#### **HEALTHCARE ANALYSIS**

# importing necessary libraries

```
In [44]:
           import numpy as np
           import pandas as pd
           import matplotlib.pyplot as plt
           %matplotlib inline
           plt.style.use('ggplot')
           import seaborn as sns
           sns.set(rc={'figure.figsize':(10,8)})
           sns.set_style("white")
           sns.set context({"figure.figsize": (10, 8)})
           from sklearn.model_selection import train_test_split, cross_val_score, KFold, GridSe
           from sklearn.preprocessing import StandardScaler
           from imblearn.combine import SMOTETomek
           from sklearn.metrics import classification_report, confusion_matrix
           import warnings
           warnings.filterwarnings('ignore')
 In [4]:
           print(plt.style.available)
          ['Solarize_Light2', '_classic_test_patch', 'bmh', 'classic', 'dark_background', 'fas
          t', 'fivethirtyeight', 'ggplot', 'grayscale', 'seaborn', 'seaborn-bright', 'seaborn-colorblind', 'seaborn-dark', 'seaborn-dark-palette', 'seaborn-darkgrid', 'seaborn-de
          ep', 'seaborn-muted', 'seaborn-notebook', 'seaborn-paper', 'seaborn-pastel', 'seabor
          n-poster', 'seaborn-talk', 'seaborn-ticks', 'seaborn-white', 'seaborn-whitegrid', 't
```

## read data

ableau-colorblind10']

```
In [2]:
          data=pd.read csv('health care diabetes.csv')
In [3]:
          data.head()
Out[3]:
             Pregnancies
                         Glucose BloodPressure SkinThickness Insulin BMI DiabetesPedigreeFunction
                                                                                                        Ag€
         0
                      6
                              148
                                             72
                                                                        33.6
                                                                                                 0.627
                                                                                                         50
         1
                      1
                                             66
                                                            29
                                                                        26.6
                                                                                                 0.351
                                                                                                         31
                              85
                                                                     0
         2
                      8
                              183
                                             64
                                                             0
                                                                        23.3
                                                                                                 0.672
                                                                                                         32
         3
                      1
                              89
                                             66
                                                            23
                                                                    94
                                                                        28.1
                                                                                                 0.167
                                                                                                         21
                      0
                             137
                                             40
                                                            35
                                                                   168 43.1
                                                                                                 2.288
                                                                                                         33
In [4]:
          data.isna().sum()
                                         0
         Pregnancies
Out[4]:
                                         0
         Glucose
         BloodPressure
                                         0
         SkinThickness
                                         0
```

Insulin

BMI

0

```
0
          DiabetesPedigreeFunction
                                         0
          Age
                                         0
          Outcome
          dtype: int64
 In [5]:
           lst=['Glucose','BloodPressure','SkinThickness','Insulin','BMI']
 In [ ]:
           si_median=SimpleImputer(missing_values=0, strategy='median')
 In [7]:
           data.describe()
                                                                                       BMI DiabetesPedig
 Out[7]:
                 Pregnancies
                                Glucose
                                         BloodPressure SkinThickness
                                                                          Insulin
          count
                   768.000000
                              768.000000
                                            768.000000
                                                           768.000000
                                                                      768.000000
                                                                                 768.000000
                    3.845052
                              120.894531
                                             69.105469
                                                            20.536458
                                                                       79.799479
                                                                                  31.992578
           mean
                    3.369578
                               31.972618
                                             19.355807
                                                            15.952218
                                                                      115.244002
                                                                                   7.884160
             std
                    0.000000
                                0.000000
                                              0.000000
                                                             0.000000
                                                                        0.000000
                                                                                   0.000000
            min
            25%
                    1.000000
                               99.000000
                                             62.000000
                                                             0.000000
                                                                        0.000000
                                                                                  27.300000
            50%
                    3.000000 117.000000
                                             72.000000
                                                            23.000000
                                                                       30.500000
                                                                                  32.000000
            75%
                    6.000000
                              140.250000
                                             80.000000
                                                            32.000000
                                                                      127.250000
                                                                                   36.600000
                   17.000000 199.000000
                                            122.000000
                                                                      846.000000
                                                                                  67.100000
                                                            99.000000
            max
 In [8]:
           for col in 1st:
                data[col]=data[col].replace(0, np.nan)
 In [9]:
           data.isna().sum()
          Pregnancies
                                            0
 Out[9]:
          Glucose
                                            5
          BloodPressure
                                          35
          SkinThickness
                                         227
          Insulin
                                         374
          BMI
                                          11
          DiabetesPedigreeFunction
                                            0
                                            0
          Age
                                            0
          Outcome
          dtype: int64
In [10]:
           for col in lst:
                if col=='Insulin':
                    data[col]=data[col].fillna(data[col].mean())
                else:
                    data[col]=data[col].fillna(data[col].median())
In [11]:
           data.isna().sum()
          Pregnancies
                                         0
Out[11]:
          Glucose
                                         0
          BloodPressure
                                         0
```

```
SkinThickness 0
Insulin 0
BMI 0
DiabetesPedigreeFunction 0
Age 0
Outcome 0
dtype: int64
```

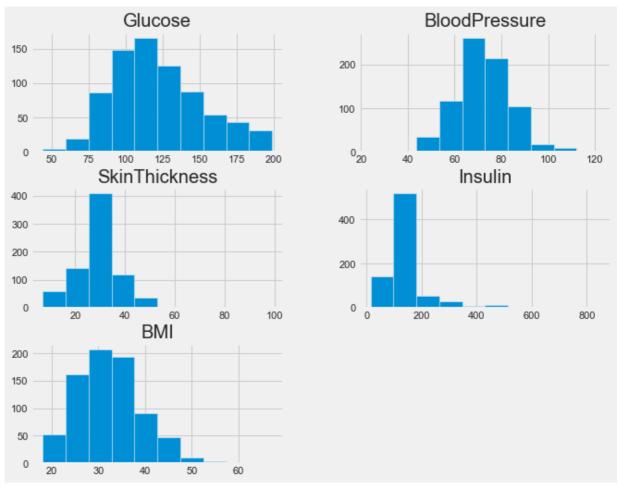
In [12]:

Out

```
data.describe()
```

t[12]:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	<b>DiabetesPedi</b> c
	count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	
	mean	3.845052	121.656250	72.386719	29.108073	155.548223	32.455208	
	std	3.369578	30.438286	12.096642	8.791221	85.021108	6.875177	
	min	0.000000	44.000000	24.000000	7.000000	14.000000	18.200000	
	25%	1.000000	99.750000	64.000000	25.000000	121.500000	27.500000	
	50%	3.000000	117.000000	72.000000	29.000000	155.548223	32.300000	
	75%	6.000000	140.250000	80.000000	32.000000	155.548223	36.600000	
	max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000	

```
In [29]: plt.style.use('fivethirtyeight')
    data[lst].hist()
```



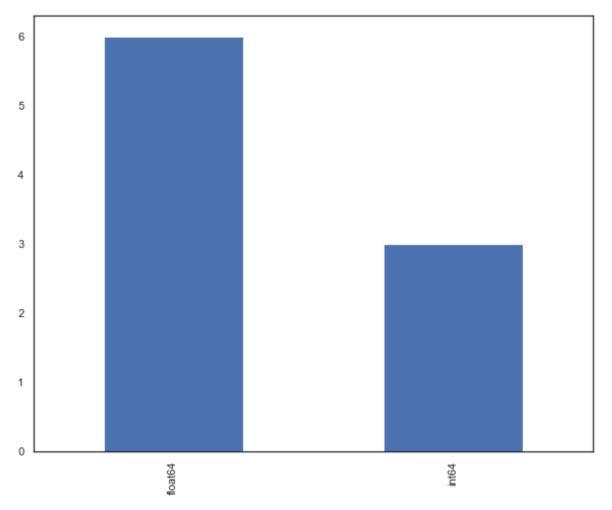
```
array([[<AxesSubplot:ylabel='Density'>, <AxesSubplot:ylabel='Density'>],
            [<AxesSubplot:ylabel='Density'>, <AxesSubplot:ylabel='Density'>], [<AxesSubplot:ylabel='Density'>, <AxesSubplot:ylabel='Density'>]],
          dtype=object)

    Glucose

                                                                                                                                 BloodPressure
  0.0125
                                                                            0.03
  0.0100
                                                                          Density
0.02
 0.0075
  0.0050
                                                                            0.01
  0.0025
  0.0000
                                                                            0.00
                 0
                         50
                                  100
                                           150
                                                    200
                                                            250
                                                                                   -25
                                                                                          0
                                                                                                 25
                                                                                                        50
                                                                                                               75
                                                                                                                      100
                                                                                                                            125
                                                                                                                                    150
                                                                                                                                           175
    0.08
                                                        SkinThickness
                                                                                                                                     - Insulin
                                                                           0.010
    0.06
                                                                           0.008
                                                                           0.006
    0.04
                                                                         0.004
    0.02
                                                                           0.002
                                                                           0.000
    0.00
               -25
                       0
                              25
                                     50
                                             75
                                                    100
                                                           125
                                                                   150
                                                                                       -250
                                                                                                        250
                                                                                                                500
                                                                                                                         750
                                                                                                                                 1000
                                                                                                                                         1250
```

```
data.dtypes.value_counts().plot.bar()
```

#### Out[14]: <AxesSubplot:>

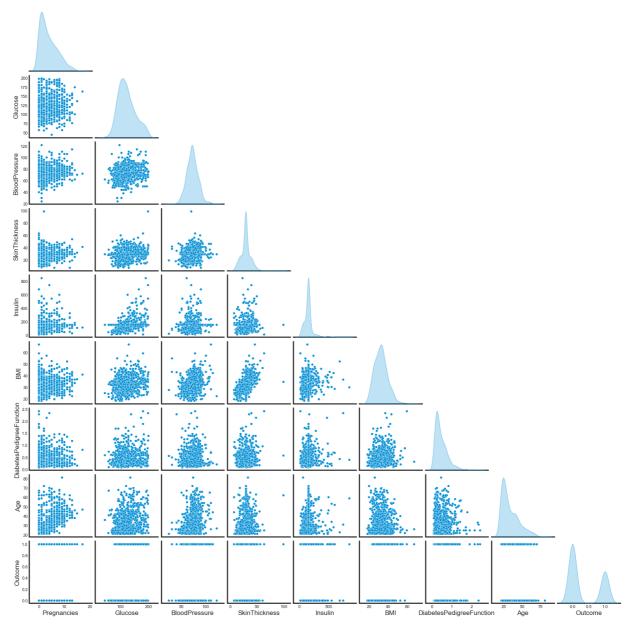


Out[16]: <AxesSubplot:>



```
plt.style.use('fivethirtyeight')
sns.set_style("white")
sns.pairplot(data=data, diag_kind='kde', corner=True)
```

Out[28]: <seaborn.axisgrid.PairGrid at 0x2b0ad88fa00>



# from the above we get some insights

- 1.BMI and SkinThickness has positive correlation
- 2.Age and Pregnancies has a relation
- 3. Insulin and Glucose has somewhat positive correlation

```
In [60]: #data.to_csv('updated_data.csv')
In [19]: X, Y=data.iloc[:,0:8], data.iloc[:, 8]
In [20]: smt=SMOTETomek(random_state=43)
    X_smt, Y_smt=smt.fit_resample(X, Y)
In [21]:
```

# as we can see we balanced our dataset using smotetek

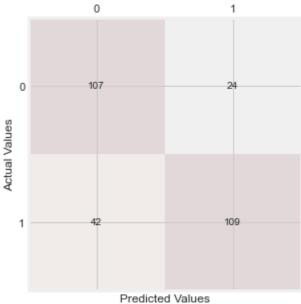
as the dependent variable is categorical(1 or 0).we go with supervised machine learning algorithms.

# we take classification algorithms from supervised learning

```
In [39]:
          from sklearn.linear_model import LogisticRegression
          from sklearn.tree import DecisionTreeClassifier
          from sklearn.neighbors import KNeighborsClassifier
          from sklearn.svm import SVC
In [40]:
          lr=LogisticRegression()
          dtc=DecisionTreeClassifier()
          knn=KNeighborsClassifier()
          svc=SVC()
In [41]:
          model_lst=[('Logistic Regression',lr),('Decision Tree',dtc),('KNearestNeighbour',knn
In [43]:
          import matplotlib.pyplot as plt
          from IPython.display import Image, display
          %matplotlib inline
          for name, model in model_lst:
              model.fit(x_train, y_train)
              y pred=model.predict(x test)
              print(name)
              #print(classification_report(y_test, y_pred))
              cm=confusion_matrix(y_test, y_pred)
              fig, ax = plt.subplots(figsize=(5, 5))
              ax.matshow(cm, cmap=plt.cm.Reds, alpha=0.1)
              for i in range(cm.shape[0]):
                   for j in range(cm.shape[1]):
```

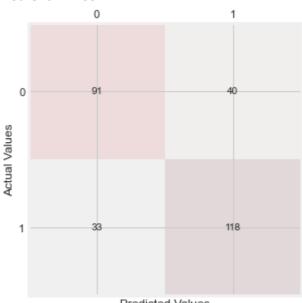
```
ax.text(x=j, y=i,
               s=cm[i, j],
                va='center', ha='center')
plt.xlabel('Predicted Values', )
plt.ylabel('Actual Values')
plt.show()
print(classification_report(y_test, y_pred ))
```

#### Logistic Regression



	precision	recall	f1-score	support
0 1	0.72 0.82	0.82 0.72	0.76 0.77	131 151
accuracy macro avg weighted avg	0.77 0.77	0.77 0.77	0.77 0.77 0.77	282 282 282

#### Decision Tree

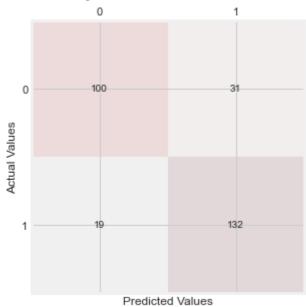


Predicted Values

	precision	recall	f1-score	support
0 1	0.73 0.75	0.69 0.78	0.71 0.76	131 151
accuracy macro avg	0.74	0.74	0.74 0.74	282 282

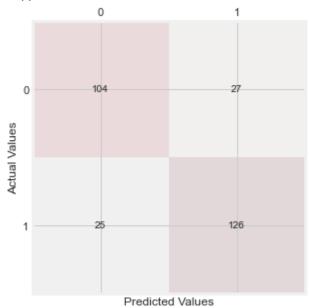
weighted avg 0.74 0.74 0.74 282

#### KNearestNeighbour



recall f1-score precision support 0 0.84 0.76 0.80 131 1 0.81 0.87 0.84 151 0.82 282 accuracy macro avg 0.83 0.82 0.82 282 weighted avg 0.82 0.82 0.82 282

#### Support Vector Machines



	precision	recall	f1-score	support
0 1	0.81 0.82	0.79 0.83	0.80 0.83	131 151
accuracy macro avg weighted avg	0.81 0.82	0.81 0.82	0.82 0.81 0.82	282 282 282

```
In [48]: X_smt=st.fit_transform(X_smt)
    for name, model in model_lst:
        print(name, cross_val_score(model, X_smt, Y_smt, scoring='accuracy', cv=10).mean
```

Logistic Regression 0.7659574468085106 Decision Tree 0.747872340425532 KNearestNeighbour 0.7946808510638298 Support Vector Machines 0.801063829787234

# **Hyper Parameter Tuning**

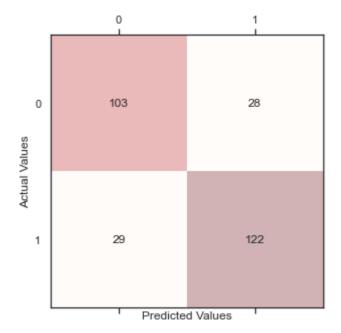
# we will do tuning for both SVC and KNN

## **SVC**

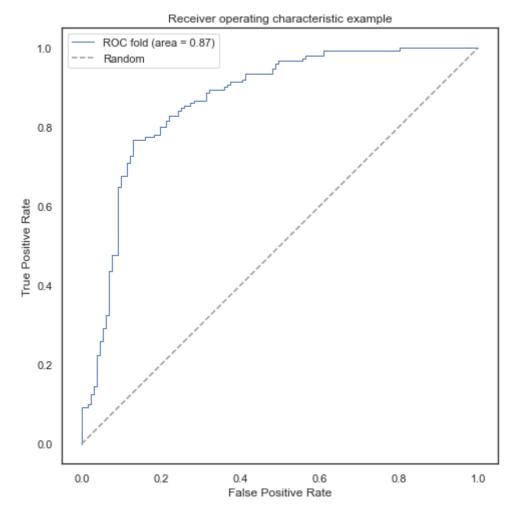
```
In [50]:
          param_grid = {'C': [0.1,1, 10, 100], 'gamma': [1,0.1,0.01,0.001], 'kernel': ['rbf',
          grid_svc=GridSearchCV(estimator=SVC(), param_grid=param_grid, cv=10, n_jobs=-1, refi
In [51]:
          grid_svc.fit(x_train, y_train)
         Fitting 10 folds for each of 48 candidates, totalling 480 fits
         ▶ GridSearchCV
Out[51]:
          ▶ estimator: SVC
                ► SVC
In [53]:
          print(grid_svc.best_estimator_)
         SVC(C=10, gamma=0.1)
In [55]:
          print("The best parameters are %s with a score of %0.2f"
                % (grid_svc.best_params_, grid_svc.best_score_))
         The best parameters are {'C': 10, 'gamma': 0.1, 'kernel': 'rbf'} with a score of 0.8
In [60]:
          grid svc.best estimator .kernel = 'rbf'
          hyp svc=grid svc.best estimator
In [61]:
          print(hyp_svc)
         SVC(C=10, gamma=0.1, probability=True)
In [63]:
          y_pred = SVC(C=10, gamma=0.1, kernel='rbf').fit(x_train, y_train).predict(x_test)
          cm = confusion_matrix(y_test, y_pred)
          #print(cm)
          print(classification_report(y_test, y_pred ))
          fig, ax = plt.subplots(figsize=(5, 5))
          ax.matshow(cm, cmap=plt.cm.Reds, alpha=0.3)
          for i in range(cm.shape[0]):
               for j in range(cm.shape[1]):
                   ax.text(x=j, y=i,
                          s=cm[i, j],
                          va='center', ha='center')
          plt.xlabel('Predicted Values', )
```

```
plt.ylabel('Actual Values')
plt.show()
```

	precision	recall	f1-score	support
0	0.78	0.79	0.78	131
1	0.81	0.81	0.81	151
accuracy			0.80	282
macro avg	0.80	0.80	0.80	282
weighted avg	0.80	0.80	0.80	282



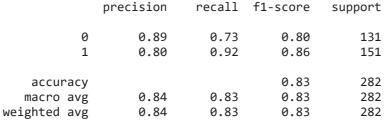
```
In [65]:
          from sklearn.metrics import roc_curve, auc
          # Plot the receiver operating characteristic curve (ROC).
          plt.figure(figsize=(10,8))
          probas_ =hyp_svc.predict_proba(x_test)
          fpr, tpr, thresholds = roc_curve(y_test, probas_[:, 1])
          roc_auc = auc(fpr, tpr)
          plt.plot(fpr, tpr, lw=1, label='ROC fold (area = %0.2f)' % (roc auc))
          plt.plot([0, 1], [0, 1], '--', color=(0.6, 0.6, 0.6), label='Random')
          plt.xlim([-0.05, 1.05])
          plt.ylim([-0.05, 1.05])
          plt.xlabel('False Positive Rate')
          plt.ylabel('True Positive Rate')
          plt.title('Receiver operating characteristic example')
          plt.legend()
          plt.axes().set_aspect(1)
```

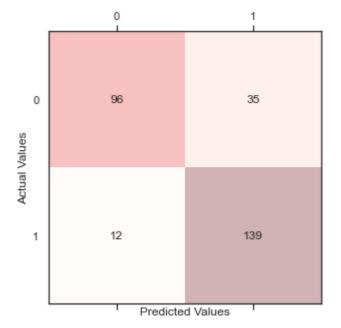


### **KNN**

```
In [66]:
          param_grid = { 'n_neighbors' : [5,7,9,11,13,15],
                         'leaf_size' : list(range(1,40)),
                          'weights' : ['uniform','distance'],
                          'metric' : ['minkowski','euclidean','manhattan']}
In [67]:
          grid_knn=GridSearchCV(estimator=KNeighborsClassifier(), param_grid=param_grid, cv=10
In [69]:
          grid_knn.fit(x_train,y_train)
         Fitting 10 folds for each of 1404 candidates, totalling 14040 fits
                      GridSearchCV
Out[69]:
          ▶ estimator: KNeighborsClassifier
                ▶ KNeighborsClassifier
In [70]:
          print(grid_knn.best_estimator_)
         KNeighborsClassifier(leaf_size=1, metric='manhattan', n_neighbors=9,
                              weights='distance')
In [72]:
          print("The best parameters are %s with a score of %0.2f"
                % (grid_knn.best_params_, grid_knn.best_score_))
```

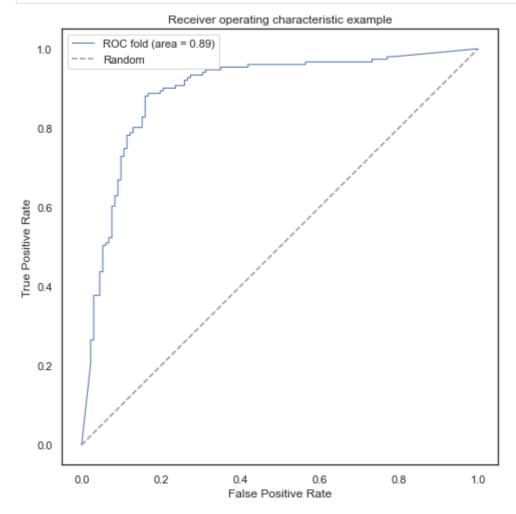
```
The best parameters are {'leaf_size': 1, 'metric': 'manhattan', 'n_neighbors': 9, 'w
         eights': 'distance'} with a score of 0.80
In [73]:
          hyp knn=grid knn.best estimator
In [74]:
          y_pred=hyp_knn.fit(x_train, y_train).predict(x_test)
          cm = confusion_matrix(y_test, y_pred)
          #print(cm)
          print(classification_report(y_test, y_pred ))
          fig, ax = plt.subplots(figsize=(5, 5))
          ax.matshow(cm, cmap=plt.cm.Reds, alpha=0.3)
          for i in range(cm.shape[0]):
               for j in range(cm.shape[1]):
                   ax.text(x=j, y=i,
                          s=cm[i, j],
                          va='center', ha='center')
          plt.xlabel('Predicted Values', )
          plt.ylabel('Actual Values')
          plt.show()
                       precision
                                    recall f1-score
                                                        support
```





```
In [75]:
    from sklearn.metrics import roc_curve, auc
    # Plot the receiver operating characteristic curve (ROC).
    plt.figure(figsize=(10,8))
    probas_ = hyp_knn.predict_proba(x_test)
    fpr, tpr, thresholds = roc_curve(y_test, probas_[:, 1])
    roc_auc = auc(fpr, tpr)
    plt.plot(fpr, tpr, lw=1, label='ROC fold (area = %0.2f)' % (roc_auc))
    plt.plot([0, 1], [0, 1], '--', color=(0.6, 0.6, 0.6), label='Random')
    plt.xlim([-0.05, 1.05])
    plt.ylim([-0.05, 1.05])
    plt.xlabel('False Positive Rate')
```

```
plt.ylabel('True Positive Rate')
plt.title('Receiver operating characteristic example')
plt.legend()
plt.axes().set_aspect(1)
```



# After Hyper Parameter Tuning its showing that KNN is slightly good than SVC

