```
In [1]:
        import numpy as np
        import pandas as pd
        from sklearn.model_selection import train_test_split
        from sklearn.preprocessing import StandardScaler
        from sklearn.neighbors import KNeighborsClassifier
        from sklearn.metrics import confusion_matrix
        from sklearn.metrics import f1_score
        from sklearn.metrics import accuracy_score
         import seaborn as sns
         import matplotlib.pyplot as plt
        %matplotlib inline
        data = pd.read_csv('diabetes.csv')
In [2]:
In [3]: data.head()
Out[3]:
            Pregnancies Glucose BloodPressure SkinThickness Insulin BMI
                                                                            Pedigree Age O
         0
                     6
                            148
                                            72
                                                          35
                                                                   0
                                                                      33.6
                                                                               0.627
                                                                                       50
         1
                             85
                                            66
                                                          29
                                                                      26.6
                                                                               0.351
                                                                                       31
         2
                     8
                            183
                                            64
                                                           0
                                                                   0
                                                                      23.3
                                                                               0.672
                                                                                       32
         3
                             89
                                            66
                                                          23
                                                                  94
                                                                      28.1
                                                                               0.167
                                                                                       21
         4
                     0
                            137
                                            40
                                                          35
                                                                 168
                                                                      43.1
                                                                               2.288
                                                                                       33
In [5]:
        data.isnull().sum()
Out[5]: Pregnancies
                          0
         Glucose
                          0
         BloodPressure
                          0
         SkinThickness
                          0
         Insulin
                          0
         BMI
                          0
         Pedigree
                          0
         Age
         Outcome
                          0
         dtype: int64
In [6]: zero_not_accepted = ['Glucose', 'BloodPressure', 'SkinThickness', 'BMI', 'Insulin']
        for col in zero_not_accepted:
In [7]:
             data[col]= data[col].replace(0,np.NaN)
             mean = int(data[col].mean(skipna=True))
             data[col] = data[col].replace(np.NaN,mean)
In [9]: data.isnull().sum()
```

17.000000

199.000000

Out[9]: Pregnancies Glucose 0 BloodPressure 0 SkinThickness 0 Insulin 0 BMI 0 Pedigree 0 0 Age Outcome dtype: int64

In [11]: data.describe()

max

Out[11]: **BloodPressure SkinThickness BMI Pregnancies** Glucose Insulin count 768.000000 768.000000 768.000000 768.000000 768.00000 768.000000 7 mean 3.845052 121.682292 72.386719 29.108073 155.28125 32.450911 std 3.369578 30.435999 12.096642 8.791221 85.02155 6.875366 min 0.000000 44.000000 24.000000 7.000000 14.00000 18.200000 25% 1.000000 99.750000 64.000000 25.000000 121.50000 27.500000 **50**% 3.000000 117.000000 72.000000 29.000000 155.00000 32.000000 **75%** 6.000000 140.250000 80.000000 32.000000 155.00000 36.600000

122.000000

99.000000

846.00000

67.100000

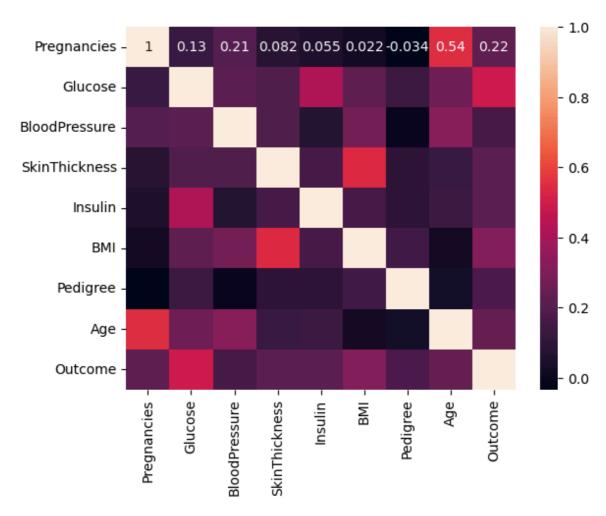
In [12]: X = data.iloc[:,0:8]

In [13]: X

					·				
Out[13]:		Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	ВМІ	Pedigree	Age
	0	6	148.0	72.0	35.0	155.0	33.6	0.627	50
	1	1	85.0	66.0	29.0	155.0	26.6	0.351	31
	2	8	183.0	64.0	29.0	155.0	23.3	0.672	32
	3	1	89.0	66.0	23.0	94.0	28.1	0.167	21
	4	0	137.0	40.0	35.0	168.0	43.1	2.288	33
	•••								
	763	10	101.0	76.0	48.0	180.0	32.9	0.171	63
	764	2	122.0	70.0	27.0	155.0	36.8	0.340	27
	765	5	121.0	72.0	23.0	112.0	26.2	0.245	30
	766	1	126.0	60.0	29.0	155.0	30.1	0.349	47
	767	1	93.0	70.0	31.0	155.0	30.4	0.315	23
	768 rows × 8 columns								
	4								<b></b>
[n [14]:	y =	data.iloc[:,	8]						

```
In [15]: y
Out[15]: 0
                 1
          1
          2
                 1
          3
                 1
          763
                0
          764
          765
          766
          767
          Name: Outcome, Length: 768, dtype: int64
In [17]: sns.heatmap(data.corr(),annot=True)
```

Out[17]: <Axes: >



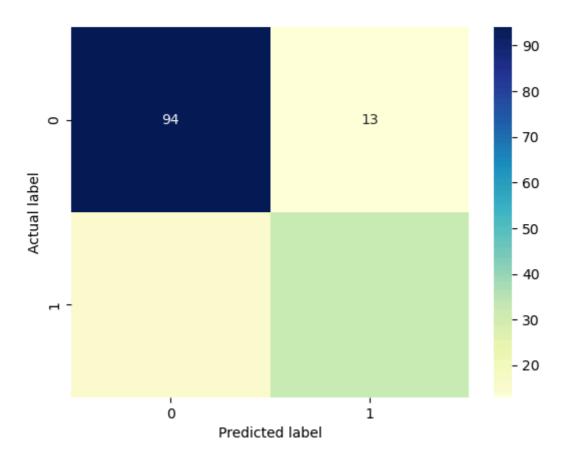
```
In [18]: X_train,X_test,y_train,y_test = train_test_split(X,y,test_size=0.2,random_state=
In [19]: scaler = StandardScaler()
         X_train = scaler.fit_transform(X_train)
         X_test = scaler.transform(X_test)
        classifier = KNeighborsClassifier(n_neighbors=11,p=2,metric='euclidean')
In [20]:
In [21]:
        classifier.fit(X_train,y_train)
Out[21]:
                            KNeighborsClassifier
         KNeighborsClassifier(metric='euclidean', n_neighbors=11)
        y_pred = classifier.predict(X_test)
In [22]:
In [23]: conf_matrix = confusion_matrix(y_test,y_pred)
         print(conf matrix)
         print(f1_score(y_test,y_pred))
        [[94 13]
         [15 32]]
        0.6956521739130435
In [31]: from sklearn.metrics import confusion_matrix
         from sklearn.metrics import accuracy_score, precision_score, recall_score, f1_sc
         y_pred = classifier.predict(X_test)
```

```
cnf_matrix = confusion_matrix(y_test, y_pred)
```

```
In [32]: p = sns.heatmap(pd.DataFrame(cnf_matrix), annot=True, cmap="YlGnBu" ,fmt='g')
    plt.title('Confusion matrix', y=1.1)
    plt.ylabel('Actual label')
    plt.xlabel('Predicted label')
```

Out[32]: Text(0.5, 23.52222222222, 'Predicted label')

## Confusion matrix



```
In [33]: accuracy_score(y_test, y_pred)
Out[33]: 0.81818181818182
In [34]: precision_score(y_test, y_pred)
Out[34]: 0.7111111111111
In [35]: recall_score(y_test, y_pred)
Out[35]: 0.6808510638297872
In [36]: f1_score(y_test, y_pred)
Out[36]: 0.6956521739130435
In []:
```