

## 1. Impoting Dependencies :

```
In [ ]: import numpy as np
import pandas as pd
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
from sklearn import svm
from sklearn.metrics import accuracy_score
```

## 2. Data Collcetion and Analysis:

A- Loading dataset to a panda dataframe:

```
In [ ]: data = pd.read_csv('C:/Machine_learning Python/projets/Diabet/diabetes.csv')
```

B- View the data (head)

```
In [ ]: data.head()
```

```
Out[ ]:   Pregnancies  Glucose  BloodPressure  SkinThickness  Insulin   BMI   DiabetesPedigreeF
```

0	6	148	72	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



C-Number of row & columns:

```
In [ ]: data.shape
```

```
Out[ ]: (768, 9)
```

## 2. Statisctical measures :

A. General Statistic:

```
In [ ]: data.describe()
```

Out[ ]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000
mean	3.845052	120.894531	69.105469	20.536458	79.799479	31.992578
std	3.369578	31.972618	19.355807	15.952218	115.244002	7.884160
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000
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
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000

B. Value of the Diabetic & Nom Diabetic Person

In [ ]:

```
data['Outcome'].value_counts()  
# 0 ---> Nom Diabetic  
# 1 --->Diabetic
```

Out[ ]:

Outcome  
0 500  
1 268  
Name: count, dtype: int64

C. Grouping by the Outcome

In [ ]:

```
data.groupby('Outcome').mean()
```

Out[ ]:

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI
Outcome						
0	3.298000	109.980000	68.184000	19.664000	68.792000	30.304200
1	4.865672	141.257463	70.824627	22.164179	100.335821	35.142537

D. Separating the data and labels

In [ ]:

```
X = data.drop(columns = 'Outcome', axis = 1)  
Y = data['Outcome']  
print(X)
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	\
0	6	148	72	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	
..	...	...	...	...	...	...	
763	10	101	76	48	180	32.9	
764	2	122	70	27	0	36.8	
765	5	121	72	23	112	26.2	
766	1	126	60	0	0	30.1	
767	1	93	70	31	0	30.4	

	DiabetesPedigreeFunction	Age
0	0.627	50
1	0.351	31
2	0.672	32
3	0.167	21
4	2.288	33
..	...	...
763	0.171	63
764	0.340	27
765	0.245	30
766	0.349	47
767	0.315	23

[768 rows x 8 columns]

In [ ]: `print(Y)`

```

0      1
1      0
2      1
3      0
4      1
..
763    0
764    0
765    0
766    1
767    0
Name: Outcome, Length: 768, dtype: int64

```

### 3. Data Standarization:

#### A. Creat the variable of strandarization

In [ ]: `scaler = StandardScaler()  
scaler.fit(X)`Out[ ]: `StandardScaler  
StandardScaler()`

#### B. Fiting and transforming on the new data:

In [ ]: `standardized_data = scaler.transform(X)`

## C. The new data:

```
In [ ]: print(standardized_data)
```

```
[ 0.63994726  0.84832379  0.14964075 ...  0.20401277  0.46849198
 1.4259954 ]
[-0.84488505 -1.12339636 -0.16054575 ... -0.68442195 -0.36506078
-0.19067191]
[ 1.23388019  1.94372388 -0.26394125 ... -1.10325546  0.60439732
-0.10558415]
...
[ 0.3429808  0.00330087  0.14964075 ... -0.73518964 -0.68519336
-0.27575966]
[-0.84488505  0.1597866  -0.47073225 ... -0.24020459 -0.37110101
 1.17073215]
[-0.84488505 -0.8730192  0.04624525 ... -0.20212881 -0.47378505
-0.87137393]]
```

## D. Creating the new X and Y

```
In [ ]: X = standardized_data
        Y = data['Outcome']
```

```
In [ ]: print(X)
```

```
[ 0.63994726  0.84832379  0.14964075 ...  0.20401277  0.46849198
 1.4259954 ]
[-0.84488505 -1.12339636 -0.16054575 ... -0.68442195 -0.36506078
-0.19067191]
[ 1.23388019  1.94372388 -0.26394125 ... -1.10325546  0.60439732
-0.10558415]
...
[ 0.3429808  0.00330087  0.14964075 ... -0.73518964 -0.68519336
-0.27575966]
[-0.84488505  0.1597866  -0.47073225 ... -0.24020459 -0.37110101
 1.17073215]
[-0.84488505 -0.8730192  0.04624525 ... -0.20212881 -0.47378505
-0.87137393]]
```

```
In [ ]: print(Y)
```

```
0      1
1      0
2      1
3      0
4      1
...
763    0
764    0
765    0
766    1
767    0
Name: Outcome, Length: 768, dtype: int64
```

## 4. Train test split:

## A. Variables of testing and training:

```
In [ ]: X_train,X_test, Y_train, Y_test = train_test_split(X,Y,test_size=0.2, stratify=Y
```

```
In [ ]: print(X.shape, X_train.shape, X_test.shape)
```

```
(768, 8) (614, 8) (154, 8)
```

#### 4. Training the model:

##### A. Variable of classification

```
In [ ]: classifier = svm.SVC(kernel='linear')
```

##### B.Training the support vector Machine Classifier:

```
In [ ]: classifier.fit(X_train,Y_train)
```

```
Out[ ]: SVC
SVC(kernel='linear')
```

#### 5. Evaluate the model:

##### A. Accuracy Score of training:

```
In [ ]: X_train_prediction = classifier.predict(X_train)
training_data_accuracy = accuracy_score(X_train_prediction, Y_train)
print('Accuracy score of the training data:', training_data_accuracy)
```

```
Accuracy score of the training data: 0.7866449511400652
```

##### B. Accuracy Score of testing:

```
In [ ]: X_test_prediction = classifier.predict(X_test)
training_data_accuracy = accuracy_score(X_test_prediction, Y_test)
print('Accuracy score of the training data:', training_data_accuracy)
```

```
Accuracy score of the training data: 0.7727272727272727
```

```
In [ ]: def predictionF(Pregnancies,Glucose,BloodPressure,SkinThickness,Insulin,BMI,Dia
input_data = (Pregnancies,Glucose,BloodPressure,SkinThickness,Insulin,BMI,Di
#Input the data into the numpy array:
input_dataNumpuy = np.asarray(input_data)
#Reshape the data:
input_dataReshaped = input_dataNumpuy.reshape(1,-1)
#Standariser the data:
std_data = scaler.transform(input_dataReshaped)
#Predict the model:
prediction = classifier.predict(std_data)
print(prediction[0])
if(prediction[0] == 1):
    print("The person is diabetic")
else:
    print("The person is not diabetic")

predictionF(0,131,0,0,0,43.2,0.27,26)
```

  
1

The person is diabetic

```
c:\Users\HP\AppData\Local\Programs\Python\Python312\Lib\site-packages\sklearn\base.py:465: UserWarning: X does not have valid feature names, but StandardScaler was fitted with feature names
  warnings.warn(
```