

Importing dependencies

```
In [1]: 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
```

Data Collection and Analysis

```
In [5]: diabetes_data = pd.read_csv('diabetes.csv')
```

```
In [7]: #prinrting the first five row of our data
```

```
In [6]: diabetes_data.head()
```

Out[6]:

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | DiabetesPedigreeFunction | Age | O |
|---|-------------|---------|---------------|---------------|---------|------|--------------------------|-----|---|
| 0 | 6 | 148 | 72 | 35 | 0 | 33.6 | 0.627 | 50 | |
| 1 | 1 | 85 | 66 | 29 | 0 | 26.6 | 0.351 | 31 | |
| 2 | 8 | 183 | 64 | 0 | 0 | 23.3 | 0.672 | 32 | |
| 3 | 1 | 89 | 66 | 23 | 94 | 28.1 | 0.167 | 21 | |
| 4 | 0 | 137 | 40 | 35 | 168 | 43.1 | 2.288 | 33 | |

```
In [8]: #number of rows and columns in this data set
diabetes_data.shape
```

Out[8]: (768, 9)

```
In [9]: #getting the stststical measures if the data
diabetes_data.describe()
```

Out[9]:

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | DiabetesPedigreeF | |
|-------|-------------|------------|---------------|---------------|------------|------------|-------------------|----|
| count | 768.000000 | 768.000000 | 768.000000 | 768.000000 | 768.000000 | 768.000000 | | 76 |
| 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 | | |

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | DiabetesPedigree |
|-----|-------------|------------|---------------|---------------|------------|-----------|------------------|
| max | 17.000000 | 199.000000 | 122.000000 | 99.000000 | 846.000000 | 67.100000 | |



```
In [10]: diabetes_data['Outcome'].value_counts()
```

```
Out[10]: 0    500
         1    268
         Name: Outcome, dtype: int64
```

0--> Non Diabetic

1--> Diabetic

```
In [12]: diabetes_data.groupby('Outcome').mean()
```

```
Out[12]:
```

| | Pregnancies | Glucose | BloodPressure | SkinThickness | Insulin | BMI | DiabetesPedigree |
|---------|-------------|------------|---------------|---------------|------------|-----------|------------------|
| 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 | |



```
In [14]: #seperaring Data and Lables
X = diabetes_data.drop(columns = 'Outcome', axis = 1)
Y = diabetes_data['Outcome']
```

```
In [15]: 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 [16]:

```
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
```

Data Standardization

In [17]:

```
scaler = StandardScaler()
```

In [18]:

```
scaler.fit(X)
```

Out[18]:

StandardScaler()

In [19]:

```
standardized_data = scaler.transform(X)
```

In [20]:

```
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]]
```

In [21]:

```
X = standardized_data
Y = diabetes_data['Outcome']
```

In [22]:

```
print(X , Y)
```

```
[ [ 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] 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
```

Train Test Split

```
In [23]: X_train , X_test ,Y_train ,Y_test = train_test_split(X ,Y ,test_size = 0.2 ,stratify =
```

```
In [24]: print(X.shape , X_train.shape , X_test.shape)
```

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

Traning the model

```
In [27]: clasifier = svm.SVC(kernel = 'linear')
```

```
In [28]: #training the support vector machine Classifier
clasifier.fit(X_train,Y_train)
```

```
Out[28]: SVC(kernel='linear')
```

Model Evaluating

Accuray score

```
In [31]: #accuracy score on training data
X_train_prediction = clasifier.predict(X_train)
training_data_accuracy = accuracy_score(X_train_prediction , Y_train)
```

```
In [33]: print('Accuracy Score of the training data',training_data_accuracy)
```

```
Accuracy Score of the training data 0.7866449511400652
```

```
In [34]: X_test_prediction = classifier.predict(X_test)
        test_data_accuracy = accuracy_score(X_test_prediction , Y_test)
```

```
In [35]: print('Accuracy score for Test data ' , test_data_accuracy)
```

Accuracy score for Test data 0.7727272727272727

Making a predictive System

```
In [50]: input_data = (1,85,66,29,0,26.6,0.351,31)
        #changing input data to numpy array
        input_data_as_numpy = np.asarray(input_data)
        #reshape the array as we are predicting for one instance
        input_data_resaped = input_data_as_numpy.reshape(1,-1)
```

```
In [51]: #standardizing the input data
        std_data = scaler.transform(input_data_resaped)
        print(std_data)
        predition = classifier.predict(std_data)
        print(predition)
```

```
[[-0.84488505 -1.12339636 -0.16054575  0.53090156 -0.69289057 -0.68442195
  -0.36506078 -0.19067191]]
[0]
```

```
In [52]: if (predition[0] == 0):
        print('The person is not diabetic')
        else:
        print('The person is diabetic')
```

The person is not diabetic