

DIABETES PREDICTION USING SUPPORT VECTOR MACHINE



Importing Libraries

```
In [33]: import pandas as pd
import numpy as np
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

- PMA Diabetes Dataset

```
In [34]: # Loading diabetes dataset to a pandas dataframe
diabetes_dataset=pd.read_csv(r"C:\Users\91766\Downloads\diabetes.csv")
```

```
In [35]: # Printing the first 5 rows of the dataset
diabetes_dataset.head()
```

```
Out[35]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	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



```
In [36]: # Number of rows and column in dataframe
diabetes_dataset.shape
```

```
Out[36]: (768, 9)
```

```
In [37]: # Getting statistical Measures of data
diabetes_dataset.describe()
```

```
Out[37]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	Diabetes
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	

```
In [40]: diabetes_dataset['Outcome'].value_counts()
```

```
Out[40]: 0    500
         1    268
         Name: Outcome, dtype: int64
```

- 0 represents Non Diabetic patients
- 1 represent Diabetic patients

```
In [41]: diabetes_dataset.groupby('Outcome').mean()
```

```
Out[41]:
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	Diabetes
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	

- after grouping we can say from the result that patient having Age upto 31 year are mostly non Diabetic and 37 and above age are mostly diabetic patients
- Glucose level,Blood pressure,skineThickness,Insulin,BMI,DPF are found high in diabetic patients

Separating the data and labels

```
In [42]: x=diabetes_dataset.drop(columns='Outcome', axis=1)
y=diabetes_dataset['Outcome']
```

```
In [43]: print(x)
print(y)
```

	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]

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 [44]: scaler=StandardScaler()  
scaler.fit(x)
```

```
Out[44]: StandardScaler()
```

In a Jupyter environment, please rerun this cell to show the HTML representation or trust the notebook.

On GitHub, the HTML representation is unable to render, please try loading this page with nbviewer.org.

```
In [45]: standardized_data=scaler.transform(x)  
standardized_data
```

```
Out[45]: array([[ 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 [46]: x=standardized_data  
y=diabetes_dataset['Outcome']
```

```
In [47]: print(x)
        print(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 [48]: xtrain,xtest,ytrain,ytest=train_test_split(x,y,test_size=0.2,stratify=y,random
```

```
In [49]: print(x.shape)
        print(xtrain.shape)
        print(xtest.shape)
```

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

Training the Model

```
In [50]: classifier=svm.SVC(kernel='linear')
```

```
In [51]: # training the support vector machine classifier
classifier.fit(xtrain,ytrain)
```

Out[51]: SVC(kernel='linear')

In a Jupyter environment, please rerun this cell to show the HTML representation or trust the notebook.

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Model Evaluation

- Accuracy Score

```
In [52]: # Accuracy score on training data
xtrain_prediction=classifier.predict(xtrain)
training_data_accuracy=accuracy_score(xtrain_prediction,ytrain)
training_data_accuracy
```

Out[52]: 0.7866449511400652

```
In [53]: print('Accuracy score of the training data :',training_data_accuracy)
```

Accuracy score of the training data : 0.7866449511400652

```
In [54]: # Accuracy score on testing data
xtest_prediction=classifier.predict(xtest)
testing_data_accuracy=accuracy_score(xtest_prediction,ytest)
testing_data_accuracy
```

Out[54]: 0.7727272727272727

```
In [55]: print('Accuracy score of the testing data :',testing_data_accuracy)
```

Accuracy score of the testing data : 0.7727272727272727

Making a Diabetes Predictive System

```
In [23]: input_data = (  
    int(input('What is the pregnancy week of the patient (0 to 40): ')),  
    int(input('What is the glucose value/level of the patient in mg (80-170): ')),  
    int(input('What is the blood pressure level in mm Hg (60-80): ')),  
    int(input('What is the skin thickness of the patient in mm (2-40): ')),  
    int(input('What is the insulin level of the patient in pmol/L (0-1153): ')),  
    float(input('What is the BMI level in kg/m2 (0-100): ')),  
    float(input('What is the Diabetes Pedigree Function level (0-5): ')),  
    int(input('What is the age of the patient in years (0-100): '))  
)  
  
# changing the Input data as numpy array  
input_data_as_numpy_array=np.asarray(input_data)  
  
# Reshape the array as we are predicting for one instance  
input_data_reshaped=input_data_as_numpy_array.reshape(1,-1)  
  
# standardized the input data  
std_data=scaler.transform(input_data_reshaped)  
  
prediction=classifier.predict(std_data)  
print(prediction)  
  
if prediction==0:  
    print('patient is not Diabetic')  
else:  
    print('Patient is Diabetic')
```

```
What is the pregnancy week of the patient (0 to 40): 0  
What is the glucose value/level of the patient in mg (80-170): 85  
What is the blood pressure level in mm Hg (60-80): 70  
What is the skin thickness of the patient in mm (2-40): 3  
What is the insulin level of the patient in pmol/L (0-1153): 250  
What is the BMI level in kg/m2 (0-100): 42  
What is the Diabetes Pedigree Function level (0-5): 1.12  
What is the age of the patient in years (0-100): 30  
[0]  
patient is not Diabetic
```

```
C:\Users\91766\AppData\Local\Programs\Python\Python311\Lib\site-packages\sklearn\base.py:439: UserWarning: X does not have valid feature names, but StandardScaler was fitted with feature names  
  warnings.warn(
```




Congratulation Patient is not diabetic.....