

```
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
import matplotlib.pyplot as plt
```

```
diabetese_dataset = pd.read_csv('/content/diabetes.csv')
```

```
diabetese_dataset.head()
```

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

Next steps:

[Generate code with diabetese_dataset](#)
[View recommended plots](#)
[New interactive sheet](#)

```
diabetese_dataset.shape
```

```
(768, 9)
```

```
# Check the unique values in the last column (target column)
unique_values = diabetese_dataset['Outcome'].unique()
print("Unique target values:", unique_values)
```

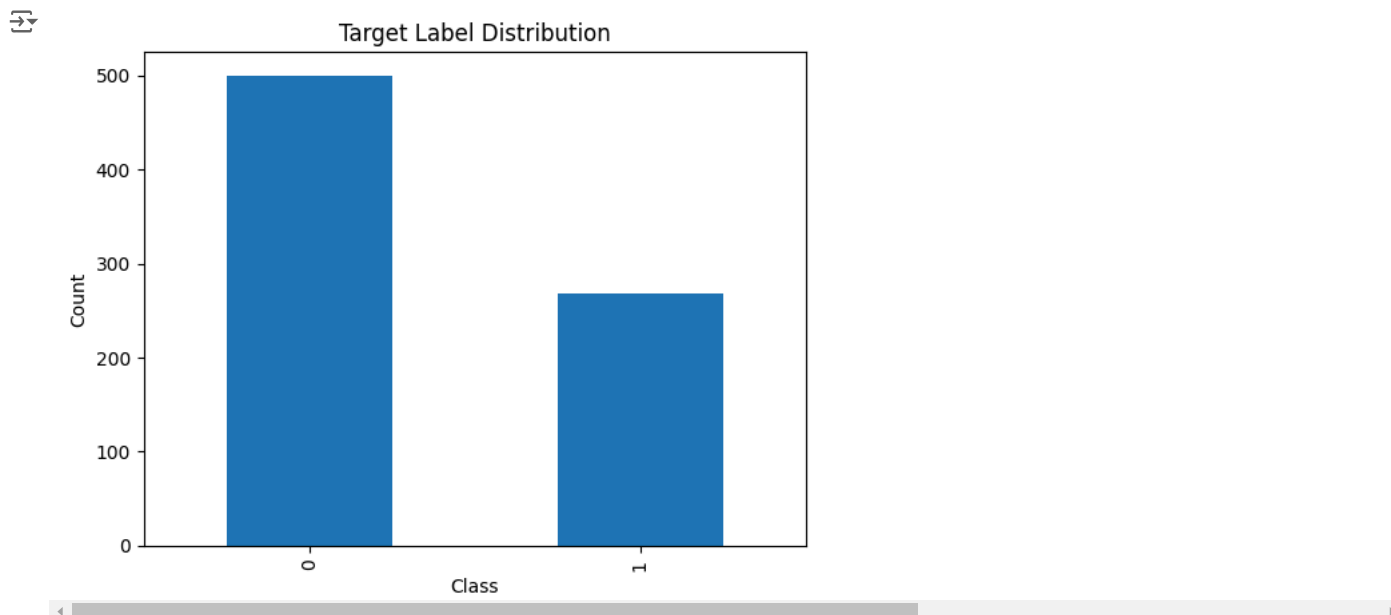
```
Unique target values: [1 0]
```

```
# Check data type and number of unique values in the target column
target = diabetese_dataset['Outcome']
print("Data type of target:", target.dtype)
print("Number of unique values:", target.nunique())
```

```
# Determine if classification or regression
if target.nunique() < 20 and target.dtype == 'object':
    print("This is a classification problem.")
else:
    print("This is a regression problem.")
```

```
Data type of target: int64
Number of unique values: 2
This is a regression problem.
```

```
# Plot the distribution of target labels
target.value_counts().plot(kind='bar')
plt.title("Target Label Distribution")
plt.xlabel("Class")
plt.ylabel("Count")
plt.show()
```



```
diabetese_dataset["Outcome"].value_counts()
```

Outcome	count
0	500
1	268

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

Outcome	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age
0	3.298000	109.980000	68.184000	19.664000	68.792000	30.304200	0.429734	31.190000
1	4.865672	141.257463	70.824627	22.164179	100.335821	35.142537	0.550500	37.067164

```
# Separating the data and variables
X = diabetese_dataset.drop(columns='Outcome', axis = 1)
Y = diabetese_dataset['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]
```

```
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
scaler = StandardScaler()

```

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

```

StandardScaler
StandardScaler()

```

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

```
display(standardized_data)
```

```

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]])

```

```
# again we separate the data after standardization
```

```

X = standardized_data
Y = diabetes_dataset['Outcome']

```

```

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
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.2, stratify=Y, random_state=2)

```

```
print("X= : ", X.shape, "\nX_train= : ", X_train.shape, "\nX_test= : ", X_test.shape)
```

```

X= : (768, 8)
X_train= : (614, 8)
X_test= : (154, 8)

```

```
# Training the model
classifier = svm.SVC(kernel='linear')
```

```
# training the SVM
classifier.fit(X_train, Y_train)
```



```
# Evaluating the model
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
```

```
X_test_prediction = classifier.predict(X_test)
tests_data_accuracy = accuracy_score(X_test_prediction, Y_test)
```

```
print("Accuracy Score of the Test data : ", tests_data_accuracy)
```

```
Accuracy Score of the Test data : 0.7727272727272727
```

```
# making the predictive system
# input_data = (4,110,92,0,0,37.6,0.191,30)
input_data = (1,189,60,23,846,30.1,0.398,59)
# convert to numpy array
input_data_as_numpy_array = np.asarray(input_data)
```

```
# reshape the array
input_data_reshaped = input_data_as_numpy_array.reshape(1, -1)
```

```
# standardize the data
std_data = scaler.transform(input_data_reshaped)
```

```
print(std_data)
```

```
prediction = classifier.predict(std_data)
print(prediction)
```

```
if (prediction[0] == 0):
    print("The Person is not diabetic")
else :
    print("The Person is diabetic")
```

```
[[[-0.84488505  2.13150675 -0.47073225  0.15453319  6.65283938 -0.24020459
    -0.2231152  2.19178518]]
```

```
[1]
```

```
The Person is diabetic
```

```
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:493: UserWarning: X does not have valid feature names, but StandardScaler was fitted
warnings.warn(
```

Start coding or [generate](#) with AI.