

Importing the Dependencies

```
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

PIMA Diabetes Dataset

```
# loading the diabetes dataset to a pandas DataFrame
diabetes_dataset = pd.read_csv('/content/diabetes.csv')
```

```
pd.read_csv?
```

```
# printing the first 5 rows of the dataset
diabetes_dataset.head()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFunction	Age	Out
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	

```
# number of rows and Columns in this dataset
diabetes_dataset.shape
```

(768, 9)

```
# getting the statistical measures of the data
diabetes_dataset.describe()
```

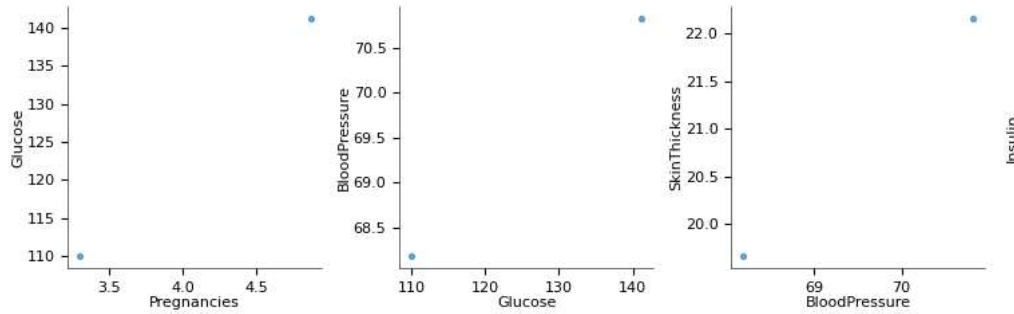
	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	DiabetesPedigreeFu	
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000		768.
mean	3.845052	120.894531	69.105469	20.536458	79.799479	31.992578		0.
std	3.369578	31.972618	19.355807	15.952218	115.244002	7.884160		0.
min	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000		0.
25%	1.000000	99.000000	62.000000	0.000000	0.000000	27.300000		0.
50%	3.000000	117.000000	72.000000	23.000000	30.500000	32.000000		0.
75%	6.000000	140.250000	80.000000	32.000000	127.250000	36.600000		0.
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000		2.

```
diabetes_dataset['Outcome'].value_counts()
```

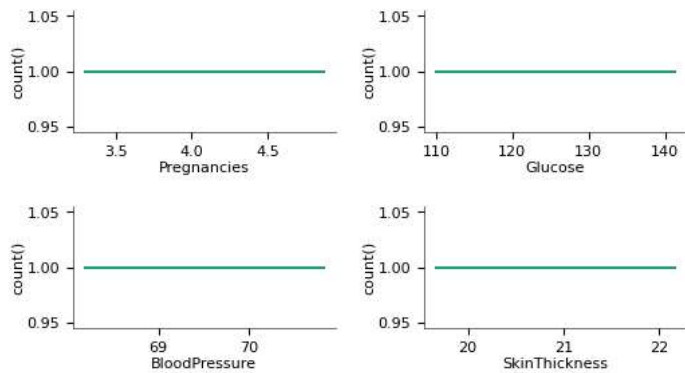
```
0    500
1    268
Name: Outcome, dtype: int64
```

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

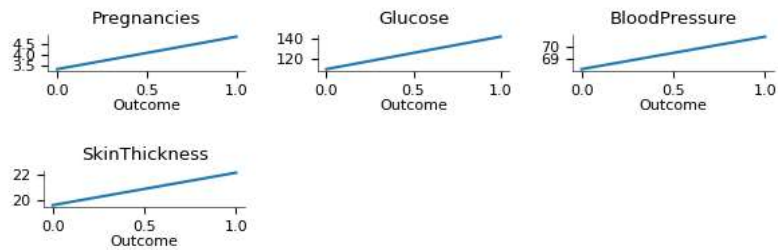
2-d distributions



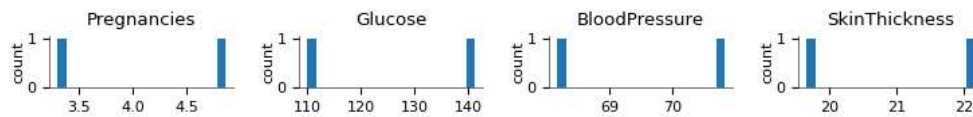
Time series



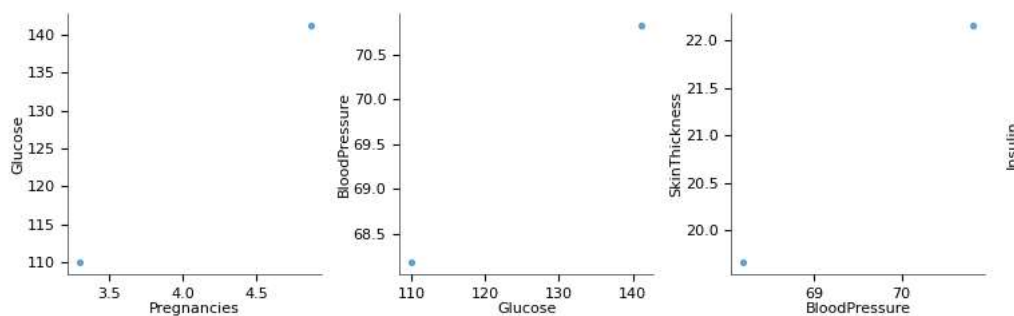
Values



Distributions



2-d distributions



Time series

```
# separating the data and labels
X = diabetes_dataset.drop(columns = 'Outcome', axis=1)
Y = diabetes_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)
```

```
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						

```

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

```

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

```
print(X.shape, X_train.shape, X_test.shape)
```

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

Training the Model

```
classifier = svm.SVC(kernel='linear')
```

```
#training the support vector Machine Classifier
classifier.fit(X_train, Y_train)
```

```

▼      SVC
SVC(kernel='linear')

```

Model Evaluation

Accuracy Score

```

# accuracy score on the training data
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

# accuracy score on the test data
X_test_prediction = classifier.predict(X_test)
test_data_accuracy = accuracy_score(X_test_prediction, Y_test)

print('Accuracy score of the test data : ', test_data_accuracy)

Accuracy score of the test data :  0.7727272727272727

```

Making a Predictive System

```
input_data = (5,166,72,19,175,25.8,0.587,51)

# changing the input_data to 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)

# standardize the input 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')

[[ 0.3429808  1.41167241  0.14964075 -0.09637905  0.82661621 -0.78595734
  0.34768723  1.51108316]]
[1]
The person is diabetic
/usr/local/lib/python3.10/dist-packages/sklearn/base.py:439: UserWarning: X does not have valid feature names, but StandardScaler was fit
warnings.warn(
```

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