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Import Libraries

```
import pandas as pd
from sklearn.tree import DecisionTreeClassifier
from sklearn.model_selection import train_test_split
from sklearn import tree
from sklearn import metrics
import matplotlib.pyplot as plt
from sklearn.metrics import confusion_matrix
from sklearn.metrics import classification_report
import seaborn as sns
```

Import Dataset

```
pd.set_option('display.max_rows', None)
d1=pd.read_csv('Diabetes.csv')
print(d1)
d1.info()
```

```
     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
5              5      116             74              0        0  25.6
6              3       78             50             32       88  31.0
7             10      115              0              0        0  35.3
8              2      197             70             45      543  30.5
9              8      125             96              0        0   0.0
10             4      110             92              0        0  37.6
11            10      168             74              0        0  38.0
12            10      139             80              0        0  27.1
13             1      189             60             23      846  30.1
14             5      166             72             19      175  25.8
15             7      100              0              0        0  30.0
16             0      118             84             47      230  45.8
17             7      107             74              0        0  29.6
18             1      103             30             38       83  43.3
19             1      115             70             30       96  34.6
```

19	1	119	70	30	96	34.6
20	3	126	88	41	235	39.3
21	8	99	84	0	0	35.4
22	7	196	90	0	0	39.8
23	9	119	80	35	0	29.0
24	11	143	94	33	146	36.6
25	10	125	70	26	115	31.1
26	7	147	76	0	0	39.4
27	1	97	66	15	140	23.2
28	13	145	82	19	110	22.2
29	5	117	92	0	0	34.1
30	5	109	75	26	0	36.0
31	3	158	76	36	245	31.6
32	3	88	58	11	54	24.8
33	6	92	92	0	0	19.9
34	10	122	78	31	0	27.6
35	4	103	60	33	192	24.0
36	11	138	76	0	0	33.2
37	9	102	76	37	0	32.9
38	2	90	68	42	0	38.2
39	4	111	72	47	207	37.1
40	3	180	64	25	70	34.0
41	7	133	84	0	0	40.2
42	7	106	92	18	0	22.7
43	9	171	110	24	240	45.4
44	7	159	64	0	0	27.4
45	0	180	66	39	0	42.0
46	1	146	56	0	0	29.7
47	2	71	70	27	0	28.0
48	7	103	66	32	0	39.1
49	7	105	0	0	0	0.0
50	1	103	80	11	82	19.4
51	1	101	50	15	36	24.2
52	5	88	66	21	23	24.4
53	8	176	90	34	300	33.7
54	7	150	66	42	342	34.7
55	1	73	50	10	0	23.0
56	7	187	68	39	304	37.7

Explore Data

```
d1.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 768 entries, 0 to 767
Data columns (total 9 columns):
 #   Column                    Non-Null Count  Dtype
---  ------                    --------------  -----
 0   Pregnancies               768 non-null    int64
 1   Glucose                   768 non-null    int64
 2   BloodPressure             768 non-null    int64
 3   SkinThickness             768 non-null    int64
 4   Insulin                   768 non-null    int64
```

```
 4   Insulin                   768 non-null    int64
 5   BMI                       768 non-null    float64
 6   DiabetesPedigreeFunction  768 non-null    float64
 7   Age                       768 non-null    int64
 8   Outcome                   768 non-null    int64
dtypes: float64(2), int64(7)
memory usage: 54.1 KB
```

```
d1.describe()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	I
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	

```
d2=d1.loc[(d1['Glucose']!=0) & (d1['BloodPressure']!=0) &
          (d1['SkinThickness']!=0) & (d1['Insulin']!=0) & (d1['BMI']!=0)]
```

```
d2.describe()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	I
count	392.000000	392.000000	392.000000	392.000000	392.000000	392.000000	
mean	3.301020	122.627551	70.663265	29.145408	156.056122	33.086224	
std	3.211424	30.860781	12.496092	10.516424	118.841690	7.027659	
min	0.000000	56.000000	24.000000	7.000000	14.000000	18.200000	
25%	1.000000	99.000000	62.000000	21.000000	76.750000	28.400000	
50%	2.000000	119.000000	70.000000	29.000000	125.500000	33.200000	
75%	5.000000	143.000000	78.000000	37.000000	190.000000	37.100000	
max	17.000000	198.000000	110.000000	63.000000	846.000000	67.100000	

```
d1['Glucose'].replace(0,d2['Glucose'].mean(),inplace=True)
d1['BloodPressure'].replace(0,d2['BloodPressure'].mean(),inplace=True)
```

```
d1['SkinThickness'].replace(0,d2['SkinThickness'].mean(),inplace=True)
d1['Insulin'].replace(0,d2['Insulin'].mean(),inplace=True)
d1['BMI'].replace(0,d2['BMI'].mean(),inplace=True)
```

```
d1.describe()
```

	Pregnancies	Glucose	BloodPressure	SkinThickness	Insulin	BMI	I
count	768.000000	768.000000	768.000000	768.000000	768.000000	768.000000	
mean	3.845052	121.692888	72.325800	29.151052	155.795560	32.466469	
std	3.369578	30.436043	12.101807	8.790943	85.021487	6.875558	
min	0.000000	44.000000	24.000000	7.000000	14.000000	18.200000	
25%	1.000000	99.750000	64.000000	25.000000	121.500000	27.500000	
50%	3.000000	117.000000	72.000000	29.145408	156.056122	32.400000	
75%	6.000000	140.250000	80.000000	32.000000	156.056122	36.600000	
max	17.000000	199.000000	122.000000	99.000000	846.000000	67.100000	

Splitting Dataset

```
x=d1.iloc[:,0:8]
y=d1.iloc[:,8]
print(x.shape)
print(y.shape)
```

```
(768, 8)
(768,)
```

```
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.2,random_state=1)
```

```
print(x_train.shape)
print(y_train.shape)
print(x_test.shape)
print(y_test.shape)
```

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

Creating Classification Model

Creating classification model

```
DC=DecisionTreeClassifier()

DC=DC.fit(x_train,y_train)
y_pred=DC.predict(x_test)
```

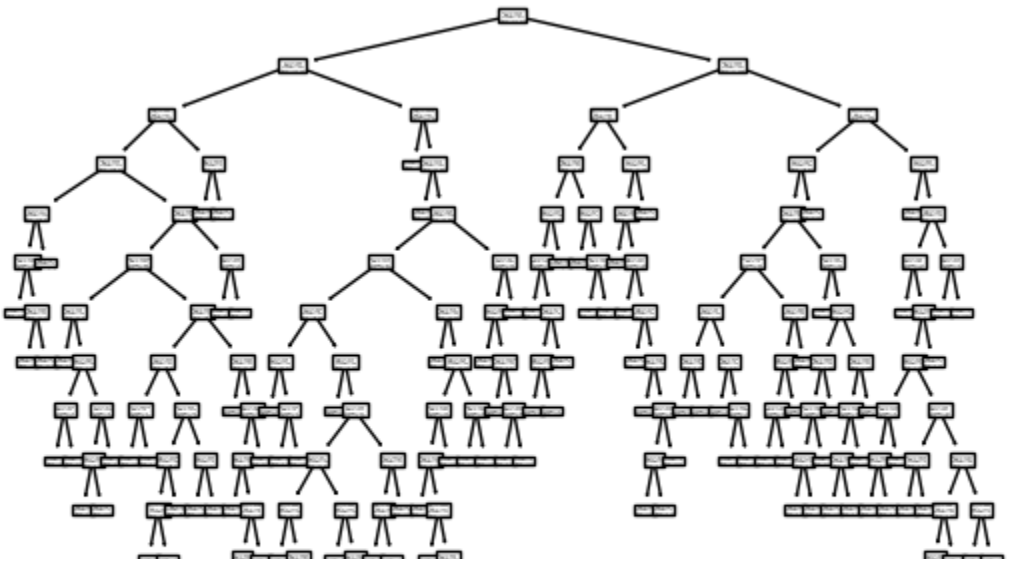
```
print(y_pred)
```

```
[0 0 0 0 0 0 0 0 0 0 1 0 0 0 0 1 0 0 0 0 1 1 1 1 0 1 1 1 0 1 0 0 0 1 1 1 0
 1 0 0 0 0 0 1 0 0 0 1 0 1 0 0 1 0 1 0 1 0 0 0 1 0 0 0 1 0 0 0 1 1 1 1 0 0
 0 1 1 1 0 1 1 0 0 1 0 0 0 0 1 1 0 0 1 0 1 1 1 0 1 0 0 1 1 0 0 0 0 1 0 0 0
 0 0 0 0 0 0 0 0 1 0 1 0 0 0 0 0 0 0 1 1 1 1 1 0 0 0 1 0 1 1 0 0 0 0 0 0 0
 0 0 0 0 1 0]
```

```
d2=pd.DataFrame({'Actual':y_test,'Predicted':y_pred})
d2.head()
```

	Actual	Predicted
285	0	0
101	0	0
581	0	0
352	0	0
726	0	0

```
tree.plot_tree(DC)
plt.savefig('o1.pdf')
```



Model Evaluation Metrics

```
print("accuracy",metrics.accuracy_score(y_test,y_pred))
```

```
accuracy 0.7012987012987013
```

```
cf=confusion_matrix(y_test,y_pred)
```

```
print(cf)
```

```
[[77 22]
 [24 31]]
```

```
print(classification_report(y_test,y_pred))
```

```
              precision    recall  f1-score   support

           0       0.76      0.78      0.77        99
           1       0.58      0.56      0.57        55

    accuracy                           0.70       154
   macro avg       0.67      0.67      0.67       154
weighted avg       0.70      0.70      0.70       154
```

Accuracy = (TP+TN) / (TP+FP+TN+FN)

Precision tells us how many of the correctly predicted cases turned out to be positive.

Precision= TP / (TP+FP)

Recall tells us how many of the actual positive cases we were able to predict correctly with our model. Recall = TP / (TP+FN)

F1 Score = 2 / ((1/Recall) + (1/Precision))

Support is the number of actual occurrences of the class in the specified dataset.

KFold Cross Validation

```
x=d1.iloc[:,0:8]
y=d1.iloc[:,8]
```

```
DC=DecisionTreeClassifier()
```

```
import sklearn.metrics as metrics
from sklearn.metrics import accuracy_score
from sklearn.model_selection import KFold
from sklearn import model_selection
kf = KFold(n_splits=5)
cv_results = model_selection.cross_val_score(DC, x, y, cv=kf)
```

```
print(cv_results)
```

```
[0.65584416 0.61038961 0.74025974 0.75163399 0.73202614]
```

```
print(cv_results.mean())
```

```
0.6980307274424922
```