

# WEEK - 07

Assessment Bulid decision tree-based model in python for like Breast cancer Wisconsin(diagnostic) dataset from sci-kit learn Or any classification dataset from UCI, Kaggle

```
In [2]: import numpy as np  
import pandas as pd  
import matplotlib.pyplot as plt
```

```
In [13]: data=pd.read_csv(r'C:\Users\God\Downloads\data.csv')  
data
```

```
Out[13]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.30010
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.08690
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.19740
3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390	0.24140
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.19800
...	...	...	...	...	...	...	...	...	...
564	926424	M	21.56	22.39	142.00	1479.0	0.11100	0.11590	0.24390
565	926682	M	20.13	28.25	131.20	1261.0	0.09780	0.10340	0.14400
566	926954	M	16.60	28.08	108.30	858.1	0.08455	0.10230	0.09251
567	927241	M	20.60	29.33	140.10	1265.0	0.11780	0.27700	0.35140
568	92751	B	7.76	24.54	47.92	181.0	0.05263	0.04362	0.00000

569 rows × 33 columns

```
In [14]: data.head()
```

```
Out[14]:
```

	id	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	p
0	842302	M	17.99	10.38	122.80	1001.0	0.11840	0.27760	0.3001	
1	842517	M	20.57	17.77	132.90	1326.0	0.08474	0.07864	0.0869	
2	84300903	M	19.69	21.25	130.00	1203.0	0.10960	0.15990	0.1974	
3	84348301	M	11.42	20.38	77.58	386.1	0.14250	0.28390	0.2414	
4	84358402	M	20.29	14.34	135.10	1297.0	0.10030	0.13280	0.1980	

5 rows × 33 columns

```
In [15]: data.info()
```

```

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 33 columns):
 #   Column           Non-Null Count Dtype  
--- 
 0   id               569 non-null    int64  
 1   diagnosis        569 non-null    object  
 2   radius_mean      569 non-null    float64 
 3   texture_mean     569 non-null    float64 
 4   perimeter_mean   569 non-null    float64 
 5   area_mean         569 non-null    float64 
 6   smoothness_mean  569 non-null    float64 
 7   compactness_mean 569 non-null    float64 
 8   concavity_mean   569 non-null    float64 
 9   concave_points_mean 569 non-null    float64 
 10  symmetry_mean   569 non-null    float64 
 11  fractal_dimension_mean 569 non-null    float64 
 12  radius_se        569 non-null    float64 
 13  texture_se       569 non-null    float64 
 14  perimeter_se    569 non-null    float64 
 15  area_se          569 non-null    float64 
 16  smoothness_se   569 non-null    float64 
 17  compactness_se  569 non-null    float64 
 18  concavity_se    569 non-null    float64 
 19  concave_points_se 569 non-null    float64 
 20  symmetry_se    569 non-null    float64 
 21  fractal_dimension_se 569 non-null    float64 
 22  radius_worst    569 non-null    float64 
 23  texture_worst   569 non-null    float64 
 24  perimeter_worst 569 non-null    float64 
 25  area_worst       569 non-null    float64 
 26  smoothness_worst 569 non-null    float64 
 27  compactness_worst 569 non-null    float64 
 28  concavity_worst 569 non-null    float64 
 29  concave_points_worst 569 non-null    float64 
 30  symmetry_worst  569 non-null    float64 
 31  fractal_dimension_worst 569 non-null    float64 
 32  date             569 non-null    int64  
dtypes: float64(30), int64(2), object(1)
memory usage: 146.8+ KB

```

```
In [16]: data=data.drop(["id"],axis=1)
```

```
In [17]: data.head(3)
```

```
Out[17]:   diagnosis  radius_mean  texture_mean  perimeter_mean  area_mean  smoothness_mean  compactness_mean  concavity_mean  concave
              points_mean
0            M       17.99       10.38        122.8       1001.0       0.11840       0.27760       0.3001       0.14710
1            M       20.57       17.77        132.9       1326.0       0.08474       0.07864       0.0869       0.07017
2            M       19.69       21.25        130.0       1203.0       0.10960       0.15990       0.1974       0.12790
```

3 rows × 32 columns

```
In [18]: M=data[data.diagnosis=="M"]
```

```
In [20]: M.head(5)
```

```
Out[20]:   diagnosis  radius_mean  texture_mean  perimeter_mean  area_mean  smoothness_mean  compactness_mean  concavity_mean  concave
              points_mean
0            M       17.99       10.38        122.80       1001.0       0.11840       0.27760       0.3001       0.14710
1            M       20.57       17.77        132.90       1326.0       0.08474       0.07864       0.0869       0.07017
2            M       19.69       21.25        130.00       1203.0       0.10960       0.15990       0.1974       0.12790
3            M       11.42       20.38        77.58        386.1       0.14250       0.28390       0.2414       0.10520
4            M       20.29       14.34        135.10       1297.0       0.10030       0.13280       0.1980       0.10430
```

5 rows × 32 columns

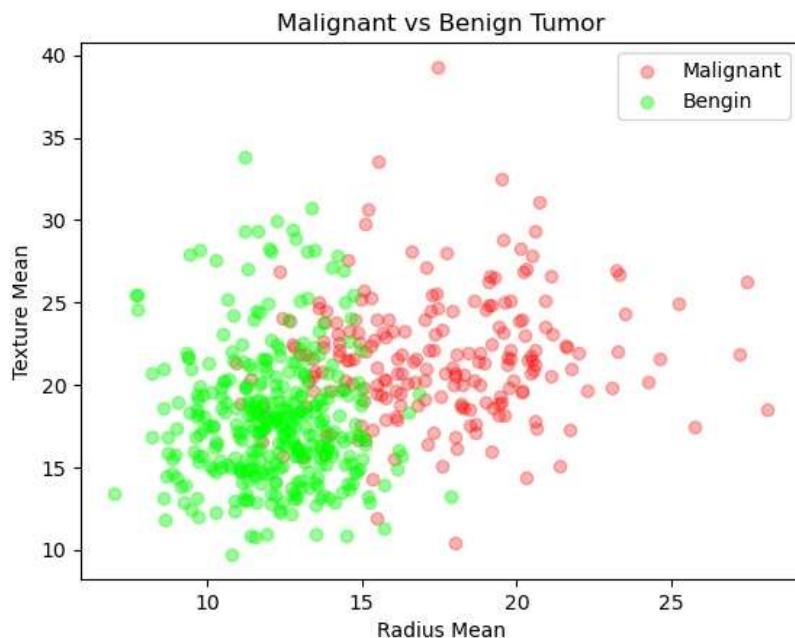
```
In [21]: B=data[data.diagnosis=="B"]
```

```
In [22]: B.head(5)
```

	diagnosis	radius_mean	texture_mean	perimeter_mean	area_mean	smoothness_mean	compactness_mean	concavity_mean	concav points_mean
19	B	13.540	14.36	87.46	566.3	0.09779	0.08129	0.06664	0.04781
20	B	13.080	15.71	85.63	520.0	0.10750	0.12700	0.04568	0.03110
21	B	9.504	12.44	60.34	273.9	0.10240	0.06492	0.02956	0.02076
37	B	13.030	18.42	82.61	523.8	0.08983	0.03766	0.02562	0.02923
46	B	8.196	16.84	51.71	201.9	0.08600	0.05943	0.01588	0.00591

5 rows × 32 columns

```
In [24]: plt.title("Malignant vs Benign Tumor")
plt.xlabel("Radius Mean")
plt.ylabel("Texture Mean")
plt.scatter(M.radius_mean,M.texture_mean,color='red',label='Malignant',alpha=0.3)
plt.scatter(B.radius_mean,B.texture_mean,color='lime',label='Bengin',alpha=0.4)
plt.legend()
plt.show()
```



```
In [25]: data.diagnosis=[1 if i == "M" else 0 for i in data.diagnosis]
```

```
In [59]: x=data.drop(["date"],axis=1)
y=data.date.values
```

```
In [61]: #Normalization
x=(x-np.min(x)) / (np.max(x)-np.min(x))
```

```
In [62]: from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test=train_test_split(x,y,test_size=0.3,random_state=42)
```

## DecisionTreeClassifier

It's type of supervised ML used to categorize or make prediction based on how a previous set of question were answered

```
In [63]: from sklearn.tree import DecisionTreeClassifier
```

```
In [64]: dt=DecisionTreeClassifier()
```

```
In [65]: dt.fit(x_train, y_train)
```

```
Out[65]: DecisionTreeClassifier()
```

```
In [66]: dt.score(x_test,y_test)
```

```
Out[66]: 1.0
```