

Apriori Algorithm --weak5

Step 1- Import required libraries

```
import csv
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from scipy.special import comb
from itertools import combinations, permutations
from apyori import apriori as apr
from mlxtend.frequent_patterns import apriori, association_rules
import scipy as sp
from mpl_toolkits.mplot3d import Axes3D
import matplotlib.pyplot as plt
import seaborn as sns
from mlxtend.preprocessing import TransactionEncoder
```

```
C:\Users\dodda\anaconda3\lib\site-packages\seaborn\rcmod.py:82:
DeprecationWarning: distutils Version classes are deprecated. Use
packaging.version instead.
```

```
    if LooseVersion(mpl.__version__) >= "3.0":
C:\Users\dodda\anaconda3\lib\site-packages\setuptools\_distutils\
version.py:351: DeprecationWarning: distutils Version classes are
deprecated. Use packaging.version instead.
    other = LooseVersion(other)
```

```
pip install apyori
```

```
Collecting apyori
```

```
  Downloading apyori-1.1.2.tar.gz (8.6 kB)
```

```
Building wheels for collected packages: apyori
```

```
  Building wheel for apyori (setup.py): started
```

```
  Building wheel for apyori (setup.py): finished with status 'done'
```

```
  Created wheel for apyori: filename=apyori-1.1.2-py3-none-any.whl
size=5974
```

```
sha256=1a8c9f8e462d6e6584ff344863721dbab45fe212311bcfd3ca8d1917e7b1933
5
```

```
  Stored in directory: c:\users\dodda\appdata\local\pip\cache\wheels\
32\2a\54\10c595515f385f3726642b10c60bf788029e8f3a1323e3913a
```

```
Successfully built apyori
```

```
Installing collected packages: apyori
```

```
Successfully installed apyori-1.1.2
```

```
Note: you may need to restart the kernel to use updated packages.
```

```
df = pd.read_csv("5_my_movies.csv")
df
```

	V1	V2	V3	V4	V5	\
0	Sixth Sense	LOTR1	Harry Potter1	Green Mile	LOTR2	
1	Gladiator	Patriot	Braveheart	NaN	NaN	
2	LOTR1	LOTR2	NaN	NaN	NaN	
3	Gladiator	Patriot	Sixth Sense	NaN	NaN	
4	Gladiator	Patriot	Sixth Sense	NaN	NaN	
5	Gladiator	Patriot	Sixth Sense	NaN	NaN	
6	Harry Potter1	Harry Potter2	NaN	NaN	NaN	
7	Gladiator	Patriot	NaN	NaN	NaN	
8	Gladiator	Patriot	Sixth Sense	NaN	NaN	
9	Sixth Sense	LOTR	Gladiator	Green Mile	NaN	

	Sixth Sense	Gladiator	LOTR1	Harry Potter1	Patriot	LOTR2	\
0	1	0	1	1	0	1	
1	0	1	0	0	1	0	
2	0	0	1	0	0	1	
3	1	1	0	0	1	0	
4	1	1	0	0	1	0	
5	1	1	0	0	1	0	
6	0	0	0	1	0	0	
7	0	1	0	0	1	0	
8	1	1	0	0	1	0	
9	1	1	0	0	0	0	

	Harry Potter2	LOTR	Braveheart	Green Mile
0	0	0	0	1
1	0	0	1	0
2	0	0	0	0
3	0	0	0	0
4	0	0	0	0
5	0	0	0	0
6	1	0	0	0
7	0	0	0	0
8	0	0	0	0
9	0	1	0	1

Step 2- Load, visualize and explore the dataset

```
df1 = df.iloc[:,5:]
df1.head()
```

	Sixth Sense	Gladiator	LOTR1	Harry Potter1	Patriot	LOTR2	\
0	1	0	1	1	0	1	
1	0	1	0	0	1	0	
2	0	0	1	0	0	1	
3	1	1	0	0	1	0	
4	1	1	0	0	1	0	

	Harry Potter2	LOTR	Braveheart	Green Mile
--	---------------	------	------------	------------

0	0	0	0	1
1	0	0	1	0
2	0	0	0	0
3	0	0	0	0
4	0	0	0	0

```
df1 = df.iloc[:,5:]
df1.head()
```

	Sixth Sense	Gladiator	LOTR1	Harry Potter1	Patriot	LOTR2	\
0	1	0	1	1	0	1	
1	0	1	0	0	1	0	
2	0	0	1	0	0	1	
3	1	1	0	0	1	0	
4	1	1	0	0	1	0	

	Harry Potter2	LOTR	Braveheart	Green Mile
0	0	0	0	1
1	0	0	1	0
2	0	0	0	0
3	0	0	0	0
4	0	0	0	0

Step 3- Clean the data set

```
df1.isnull().sum()
```

```
Sixth Sense      0
Gladiator        0
LOTR1            0
Harry Potter1    0
Patriot          0
LOTR2            0
Harry Potter2    0
LOTR             0
Braveheart       0
Green Mile       0
dtype: int64
```

```
df1
```

	Sixth Sense	Gladiator	LOTR1	Harry Potter1	Patriot	LOTR2	\
0	1	0	1	1	0	1	
1	0	1	0	0	1	0	
2	0	0	1	0	0	1	
3	1	1	0	0	1	0	
4	1	1	0	0	1	0	
5	1	1	0	0	1	0	
6	0	0	0	1	0	0	
7	0	1	0	0	1	0	

8	1	1	0	0	1	0
9	1	1	0	0	0	0

	Harry Potter2	LOTR	Braveheart	Green Mile
0	0	0	0	1
1	0	0	1	0
2	0	0	0	0
3	0	0	0	0
4	0	0	0	0
5	0	0	0	0
6	1	0	0	0
7	0	0	0	0
8	0	0	0	0
9	0	1	0	1

#Setting different thresholds

confidence = [0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.8,0.9]

Step - 4 Generating Association Rules

```
def gen_rules(df,confidence,support):
    ap = {}
    for i in confidence:
        ap_i =apriori(df1,support,True)
        rule= association_rules(ap_i,min_threshold=i)
        ap[i] = len(rule.antecedents)
    return pd.Series(ap).to_frame("Support: %s"%support)
```

```
confs = []
ap_i = gen_rules(df1,confidence=confidence,support=0.1)
confs.append(ap_i)
all_conf = pd.concat(confs,axis=1)
```

C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types result in worse computational performance and their support might be discontinued in the future.Please use a DataFrame with bool type

warnings.warn(

C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types result in worse computational performance and their support might be discontinued in the future.Please use a DataFrame with bool type

warnings.warn(

C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types result in worse computational performance and their support might be discontinued in the future.Please use a DataFrame with bool type

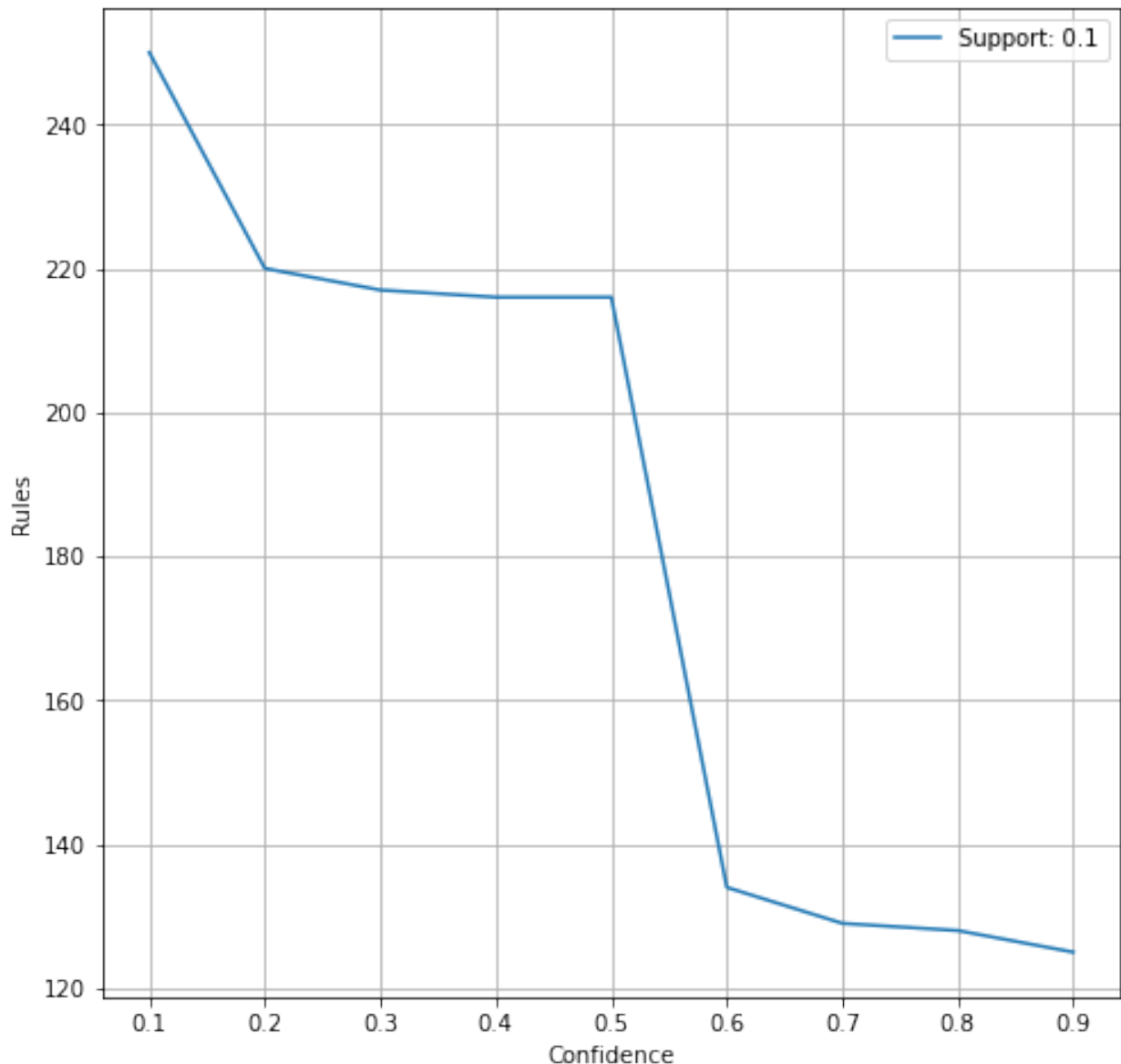
warnings.warn(

C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types

```
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discontinued in the future.Please use a DataFrame with bool type
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C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computationalperformance and their support might be
discontinued in the future.Please use a DataFrame with bool type
warnings.warn(
```

Step - 5 Visuvalizing Association Rules with different support and confidence thresholds

```
all_conf.plot(figsize=(8,8),grid=True)
plt.ylabel('Rules')
plt.xlabel('Confidence')
plt.show()
```

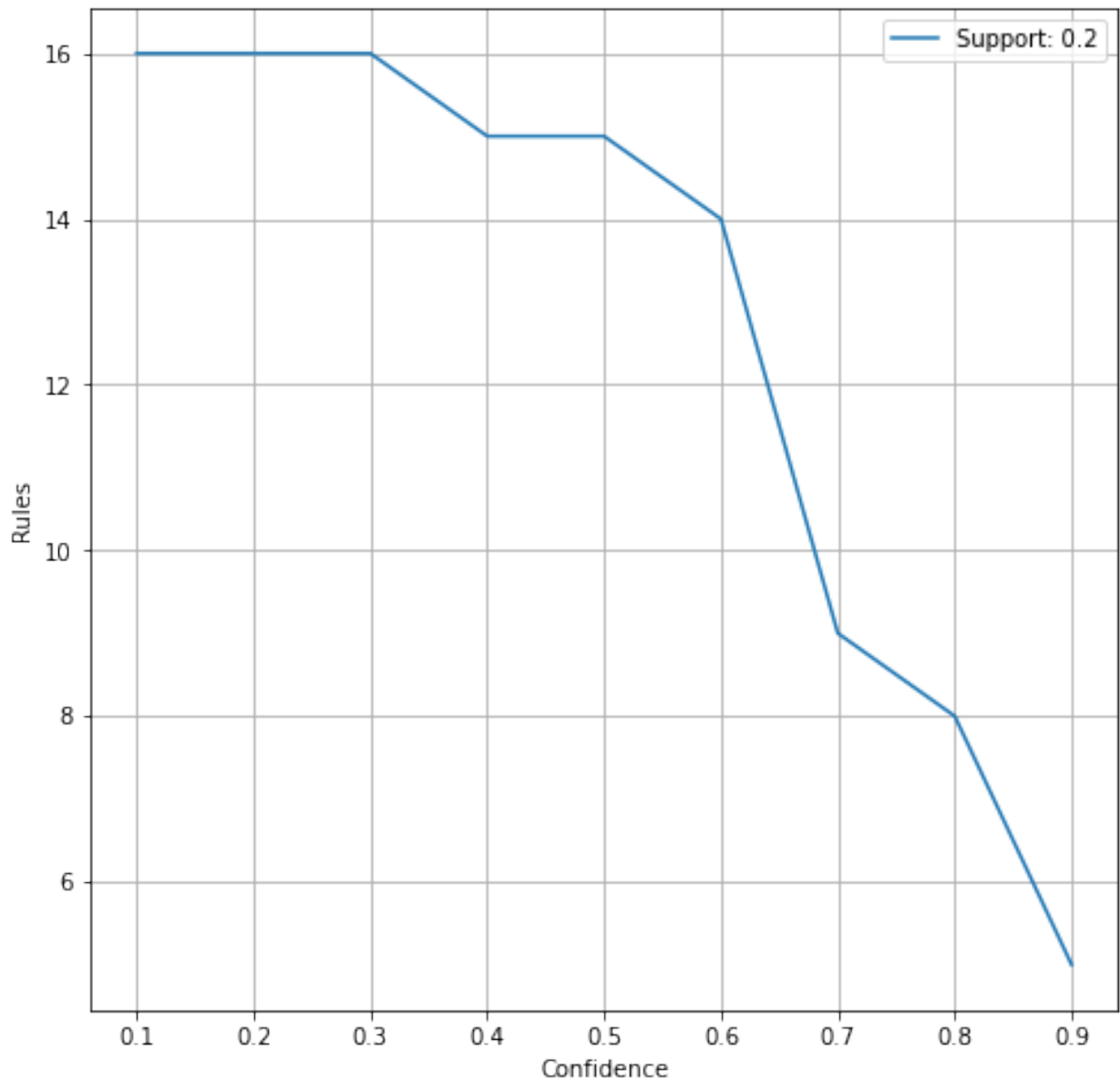


```
confs = []
ap_i = gen_rules(df1, confidence=confidence, support=0.2)
confs.append(ap_i)
all_conf = pd.concat(confs, axis=1)
```

```
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computational performance and their support might be
discontinued in the future. Please use a DataFrame with bool type
warnings.warn(
```

```
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computational performance and their support might be
discontinued in the future. Please use a DataFrame with bool type
```

```
warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computationalperformance and their support might be
discontinued in the future.Please use a DataFrame with bool type
warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
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warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computationalperformance and their support might be
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warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
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warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computationalperformance and their support might be
discontinued in the future.Please use a DataFrame with bool type
warnings.warn(
C:\Users\dodda\anaconda3\lib\site-packages\mlxtend\frequent_patterns\
fpcommon.py:111: DeprecationWarning: DataFrames with non-bool types
result in worse computationalperformance and their support might be
discontinued in the future.Please use a DataFrame with bool type
warnings.warn(
all_conf.plot(figsize=(8,8),grid=True)
plt.ylabel('Rules')
plt.xlabel('Confidence')
plt.show()
```



weak3-multiple linear regression-

```
# Importing Required files

import numpy as np
import pandas as pd
from sklearn import linear_model

# Importing the dataset
df = pd.read_csv("data.csv")

df
```


	Car	Model	Volume	Weight	CO2
0	Toyoty	Aygo	1000	790	99
1	Mitsubishi	Space Star	1200	1160	95
2	Skoda	Citigo	1000	929	95
3	Fiat	500	900	865	90
4	Mini	Cooper	1500	1140	105
5	VW	Up!	1000	929	105
6	Skoda	Fabia	1400	1109	90
7	Mercedes	A-Class	1500	1365	92
8	Ford	Fiesta	1500	1112	98
9	Audi	A1	1600	1150	99
10	Hyundai	I20	1100	980	99
11	Suzuki	Swift	1300	990	101
12	Ford	Fiesta	1000	1112	99
13	Honda	Civic	1600	1252	94
14	Hundai	I30	1600	1326	97
15	Opel	Astra	1600	1330	97
16	BMW	1	1600	1365	99
17	Mazda	3	2200	1280	104
18	Skoda	Rapid	1600	1119	104
19	Ford	Focus	2000	1328	105
20	Ford	Mondeo	1600	1584	94
21	Opel	Insignia	2000	1428	99
22	Mercedes	C-Class	2100	1365	99
23	Skoda	Octavia	1600	1415	99
24	Volvo	S60	2000	1415	99
25	Mercedes	CLA	1500	1465	102
26	Audi	A4	2000	1490	104
27	Audi	A6	2000	1725	114
28	Volvo	V70	1600	1523	109
29	BMW	5	2000	1705	114
30	Mercedes	E-Class	2100	1605	115
31	Volvo	XC70	2000	1746	117
32	Ford	B-Max	1600	1235	104
33	BMW	216	1600	1390	108
34	Opel	Zafira	1600	1405	109
35	Mercedes	SLK	2500	1395	120

selecting dependent and independent variables

```
X = df[['Weight', 'Volume']]
```

```
y = df['CO2']
```

X

	Weight	Volume
0	790	1000
1	1160	1200
2	929	1000
3	865	900
4	1140	1500

5	929	1000
6	1109	1400
7	1365	1500
8	1112	1500
9	1150	1600
10	980	1100
11	990	1300
12	1112	1000
13	1252	1600
14	1326	1600
15	1330	1600
16	1365	1600
17	1280	2200
18	1119	1600
19	1328	2000
20	1584	1600
21	1428	2000
22	1365	2100
23	1415	1600
24	1415	2000
25	1465	1500
26	1490	2000
27	1725	2000
28	1523	1600
29	1705	2000
30	1605	2100
31	1746	2000
32	1235	1600
33	1390	1600
34	1405	1600
35	1395	2500

y

0	99
1	95
2	95
3	90
4	105
5	105
6	90
7	92
8	98
9	99
10	99
11	101
12	99
13	94
14	97
15	97

```

16      99
17     104
18     104
19     105
20      94
21      99
22      99
23      99
24      99
25     102
26     104
27     114
28     109
29     114
30     115
31     117
32     104
33     108
34     109
35     120
Name: C02, dtype: int64

#fitting the model
regr = linear_model.LinearRegression()
regr.fit(X, y)

LinearRegression()

#predict the C02 emission of a car where the weight is 2300kg, and the
volume is 1300cm3:
predictedC02 = regr.predict([[2300, 1300]])

print(predictedC02)

[107.2087328]

C:\Users\dodda\anaconda3\lib\site-packages\sklearn\base.py:450:
UserWarning: X does not have valid feature names, but LinearRegression
was fitted with feature names
  warnings.warn(

```

week2-logistic regression.-

```

# Importing Required files

import numpy as np
import pandas as pd
from sklearn import linear_model

```

```

#X represents the size of a tumor in centimeters.
X = np.array([3.78, 2.44, 2.09, 0.14, 1.72, 1.65, 4.92, 4.37, 4.96,
4.52, 3.69, 5.88]).reshape(-1,1)

#Note: X has to be reshaped into a column from a row for the
LogisticRegression() function to work.
#y represents whether or not the tumor is cancerous (0 for "No", 1 for
"Yes").
y = np.array([0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1])

X
array([[3.78],
       [2.44],
       [2.09],
       [0.14],
       [1.72],
       [1.65],
       [4.92],
       [4.37],
       [4.96],
       [4.52],
       [3.69],
       [5.88]])

y
array([0, 0, 0, 0, 0, 0, 1, 1, 1, 1, 1, 1])

#Fitting the model
logr = linear_model.LogisticRegression()
logr.fit(X,y)

LogisticRegression()

#predict if tumor is cancerous where the size is 3.46mm:
predicted = logr.predict(np.array([3.46]).reshape(-1,1))
predicted
array([0])

```

We have predicted that a tumor with a size of 3.46mm will not be cancerous.

weak6-Perform clustering using k-means clustering algorithm-

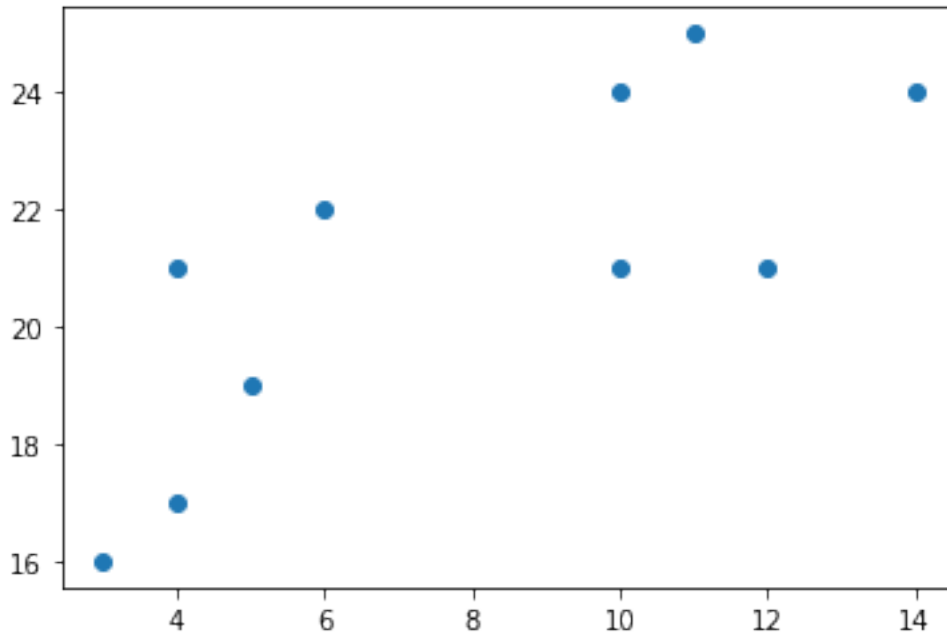
```

#1.Start by visualizing some data points:
import matplotlib.pyplot as plt

```

```
x = [4, 5, 10, 4, 3, 11, 14, 6, 10, 12]
y = [21, 19, 24, 17, 16, 25, 24, 22, 21, 21]

plt.scatter(x, y)
plt.show()
```



#2. Now we utilize the elbow method to visualize the inertia for different values of K:

```
from sklearn.cluster import KMeans

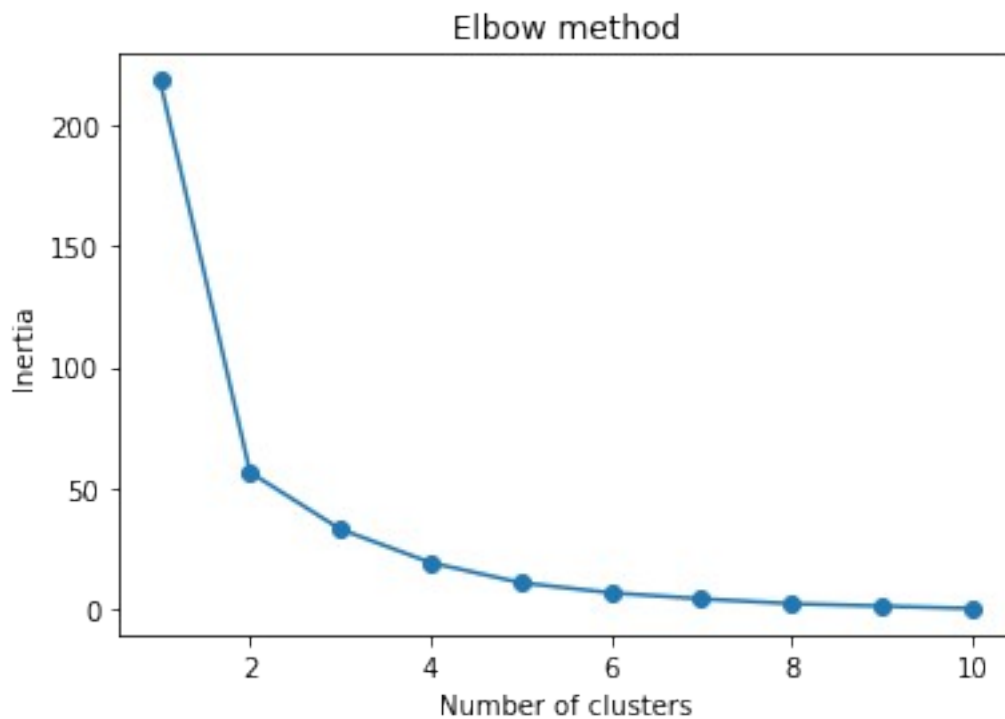
data = list(zip(x, y))
inertias = []

for i in range(1, 11):
    kmeans = KMeans(n_clusters=i)
    kmeans.fit(data)
    inertias.append(kmeans.inertia_)

plt.plot(range(1, 11), inertias, marker='o')
plt.title('Elbow method')
plt.xlabel('Number of clusters')
plt.ylabel('Inertia')
plt.show()
```

C:\Users\dodda\anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:1036: UserWarning: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks than available threads. You can avoid it by setting the environment variable

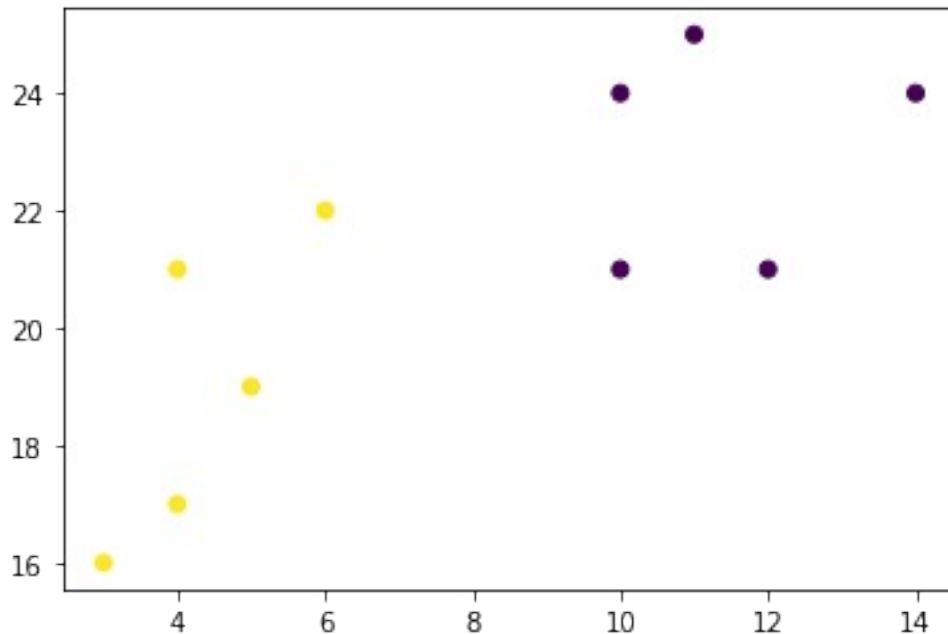
```
OMP_NUM_THREADS=1.  
warnings.warn(
```



#3. The elbow method shows that 2 is a good value for K, so we retrain and visualize the result:

```
kmeans = KMeans(n_clusters=2)  
kmeans.fit(data)
```

```
plt.scatter(x, y, c=kmeans.labels_)  
plt.show()
```



weak7-Perform Principle Component Analysis and then perform clustering-

Step 1- Import required libraries

```
import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
from sklearn.cluster import KMeans
from sklearn.preprocessing import StandardScaler
from sklearn import preprocessing
import seaborn as sns
from sklearn.decomposition import PCA
import scipy.cluster.hierarchy as sch
```

Step 2- Load, visualize and explore the dataset

```
df = pd.read_csv("7_wine.csv")
df.head()
```

	Type	Alcohol	Malic	Ash	Alcalinity	Magnesium	Phenols
0	1	14.23	1.71	2.43	15.6	127	2.80
1	1	13.20	1.78	2.14	11.2	100	2.65

2	1	13.16	2.36	2.67	18.6	101	2.80
3.24							
3	1	14.37	1.95	2.50	16.8	113	3.85
3.49							
4	1	13.24	2.59	2.87	21.0	118	2.80
2.69							
		Nonflavanoids	Proanthocyanins	Color	Hue	Dilution	Proline
0		0.28	2.29	5.64	1.04	3.92	1065
1		0.26	1.28	4.38	1.05	3.40	1050
2		0.30	2.81	5.68	1.03	3.17	1185
3		0.24	2.18	7.80	0.86	3.45	1480
4		0.39	1.82	4.32	1.04	2.93	735

```
y= df['Type']
df1 = df.iloc[:, 1:]
df1.head()
```

	Alcohol	Malic	Ash	Alcalinity	Magnesium	Phenols	Flavanoids	\
0	14.23	1.71	2.43	15.6	127	2.80	3.06	
1	13.20	1.78	2.14	11.2	100	2.65	2.76	
2	13.16	2.36	2.67	18.6	101	2.80	3.24	
3	14.37	1.95	2.50	16.8	113	3.85	3.49	
4	13.24	2.59	2.87	21.0	118	2.80	2.69	

	Nonflavanoids	Proanthocyanins	Color	Hue	Dilution	Proline
0	0.28	2.29	5.64	1.04	3.92	1065
1	0.26	1.28	4.38	1.05	3.40	1050
2	0.30	2.81	5.68	1.03	3.17	1185
3	0.24	2.18	7.80	0.86	3.45	1480
4	0.39	1.82	4.32	1.04	2.93	735

y *#actual clusters*

```
0      1
1      1
2      1
3      1
4      1
..
173    3
174    3
175    3
176    3
177    3
Name: Type, Length: 178, dtype: int64
```

Step 3- Feature Scaling


```
# normalizing the data
df_norm = StandardScaler().fit_transform(df1)

df_norm
array([[ 1.51861254, -0.5622498,  0.23205254, ...,  0.36217728,
         1.84791957,  1.01300893],
       [ 0.24628963, -0.49941338, -0.82799632, ...,  0.40605066,
         1.1134493,  0.96524152],
       [ 0.19687903,  0.02123125,  1.10933436, ...,  0.31830389,
         0.78858745,  1.39514818],
       ...,
       [ 0.33275817,  1.74474449, -0.38935541, ..., -1.61212515,
        -1.48544548,  0.28057537],
       [ 0.20923168,  0.22769377,  0.01273209, ..., -1.56825176,
        -1.40069891,  0.29649784],
       [ 1.39508604,  1.58316512,  1.36520822, ..., -1.52437837,
        -1.42894777, -0.59516041]])
```

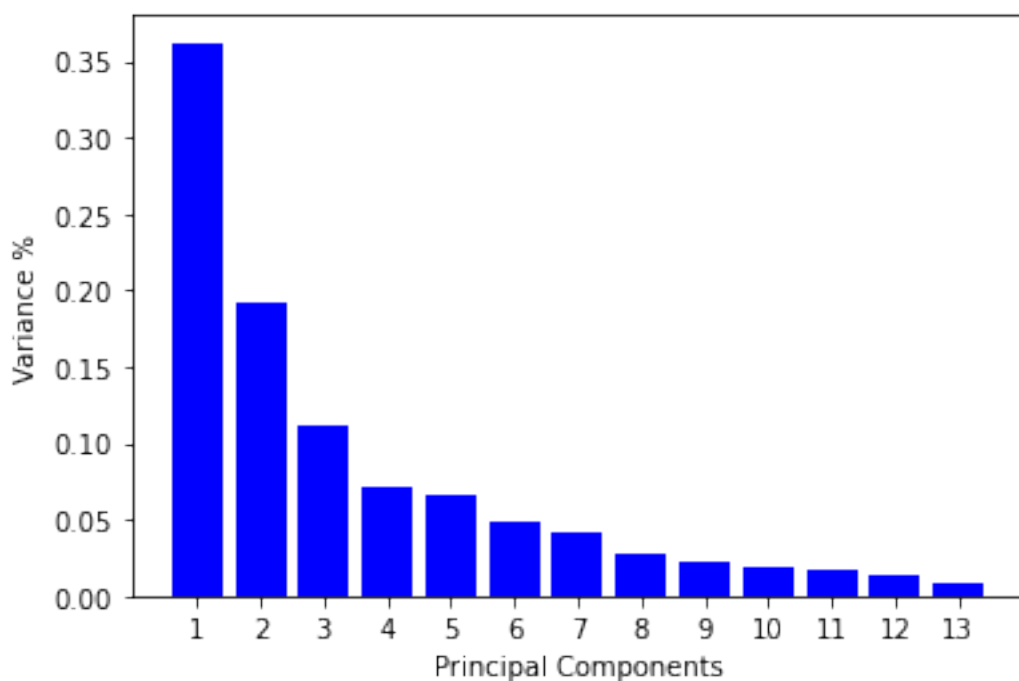
Step 4- Dimensionality Reduction with PCA

```
# Applying PCA function
pca = PCA(n_components=13)
principalComponents = pca.fit_transform(df_norm)

PC = range(1, pca.n_components_+1)
plt.bar(PC, pca.explained_variance_ratio_, color='blue')
plt.xlabel('Principal Components')
plt.ylabel('Variance %')
plt.xticks(PC)

([<matplotlib.axis.XTick at 0x21183ef3c10>,
 <matplotlib.axis.XTick at 0x21183ef3be0>,
 <matplotlib.axis.XTick at 0x21183ef3310>,
 <matplotlib.axis.XTick at 0x21183f450a0>,
 <matplotlib.axis.XTick at 0x21183f45550>,
 <matplotlib.axis.XTick at 0x21183f45be0>,
 <matplotlib.axis.XTick at 0x21183f4a370>,
 <matplotlib.axis.XTick at 0x21183f4aac0>,
 <matplotlib.axis.XTick at 0x21183f52250>,
 <matplotlib.axis.XTick at 0x21183f529a0>,
 <matplotlib.axis.XTick at 0x21183f52610>,
 <matplotlib.axis.XTick at 0x21183f4a5e0>,
 <matplotlib.axis.XTick at 0x21183f550d0>],
 [Text(0, 0, ''),
  Text(0, 0, ''),
  Text(0, 0, ''),
  Text(0, 0, ''),
  Text(0, 0, ''),
  Text(0, 0, ''),
  Text(0, 0, '')])
```

```
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, ''),
Text(0, 0, '')[0, 0, '']])
```



```
pca.explained_variance_ratio_
```

```
array([0.36198848, 0.1920749 , 0.11123631, 0.0706903 , 0.06563294,
       0.04935823, 0.04238679, 0.02680749, 0.02222153, 0.01930019,
       0.01736836, 0.01298233, 0.00795215])
```

```
PCA_components = pd.DataFrame(principalComponents)
```

```
PCA_components
```

```

      0      1      2      3      4      5
6  \
0   3.316751 -1.443463 -0.165739 -0.215631  0.693043 -0.223880
0.596427
1   2.209465  0.333393 -2.026457 -0.291358 -0.257655 -0.927120
0.053776
2   2.516740 -1.031151  0.982819  0.724902 -0.251033  0.549276
0.424205
3   3.757066 -2.756372 -0.176192  0.567983 -0.311842  0.114431 -
0.383337
```

```

4      1.008908 -0.869831  2.026688 -0.409766  0.298458 -0.406520
0.444074
...      ...      ...      ...      ...      ...      ...
...
173 -3.370524 -2.216289 -0.342570  1.058527 -0.574164 -1.108788
0.958416
174 -2.601956 -1.757229  0.207581  0.349496  0.255063 -0.026465
0.146894
175 -2.677839 -2.760899 -0.940942  0.312035  1.271355  0.273068
0.679235
176 -2.387017 -2.297347 -0.550696 -0.688285  0.813955  1.178783
0.633975
177 -3.208758 -2.768920  1.013914  0.596903 -0.895193  0.296092
0.005741

      7      8      9      10      11      12
0      0.065139  0.641443  1.020956 -0.451563  0.540810 -0.066239
1      1.024416 -0.308847  0.159701 -0.142657  0.388238  0.003637
2     -0.344216 -1.177834  0.113361 -0.286673  0.000584  0.021717
3      0.643593  0.052544  0.239413  0.759584 -0.242020 -0.369484
4      0.416700  0.326819 -0.078366 -0.525945 -0.216664 -0.079364
...      ...      ...      ...      ...      ...
173 -0.146097 -0.022498 -0.304117  0.139228  0.170786 -0.114427
174 -0.552427 -0.097969 -0.206061  0.258198 -0.279431 -0.187371
175  0.047024  0.001222 -0.247997  0.512492  0.698766  0.072078
176  0.390829  0.057448  0.491490  0.299822  0.339821 -0.021866
177 -0.292914  0.741660 -0.117969 -0.229964 -0.188788 -0.323965

[178 rows x 13 columns]

```

Step 5- We have to find the optimal K value for clustering the data. Now we are using the Elbow method to find the optimal K value.

```

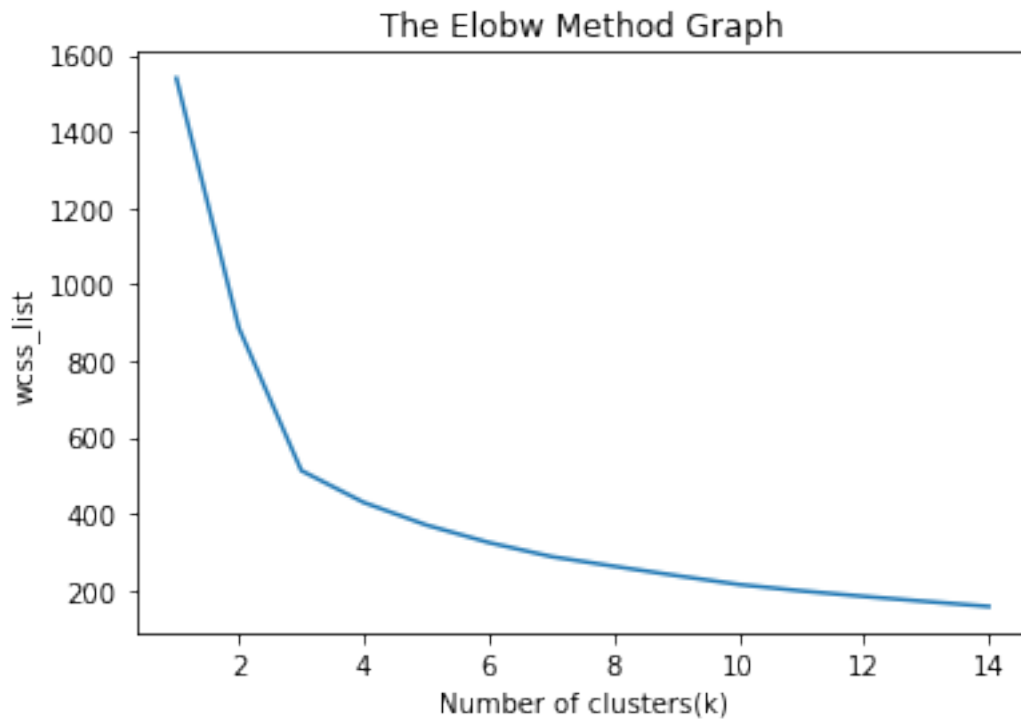
wcss = []
for i in range(1, 15):
    kmeans = KMeans(n_clusters = i, init = 'k-means++', random_state =
42)
    kmeans.fit(PCA_components.iloc[:, :3])
    wcss.append(kmeans.inertia_)

C:\Users\dodda\anaconda3\lib\site-packages\sklearn\cluster\
_kmeans.py:1036: UserWarning: KMeans is known to have a memory leak on
Windows with MKL, when there are less chunks than available threads.
You can avoid it by setting the environment variable
OMP_NUM_THREADS=1.
  warnings.warn(

plt.plot(range(1, 15), wcss)
plt.title('The Elbow Method Graph')
plt.xlabel('Number of clusters(k)')

```

```
plt.ylabel('wcss_list')
plt.show()
```

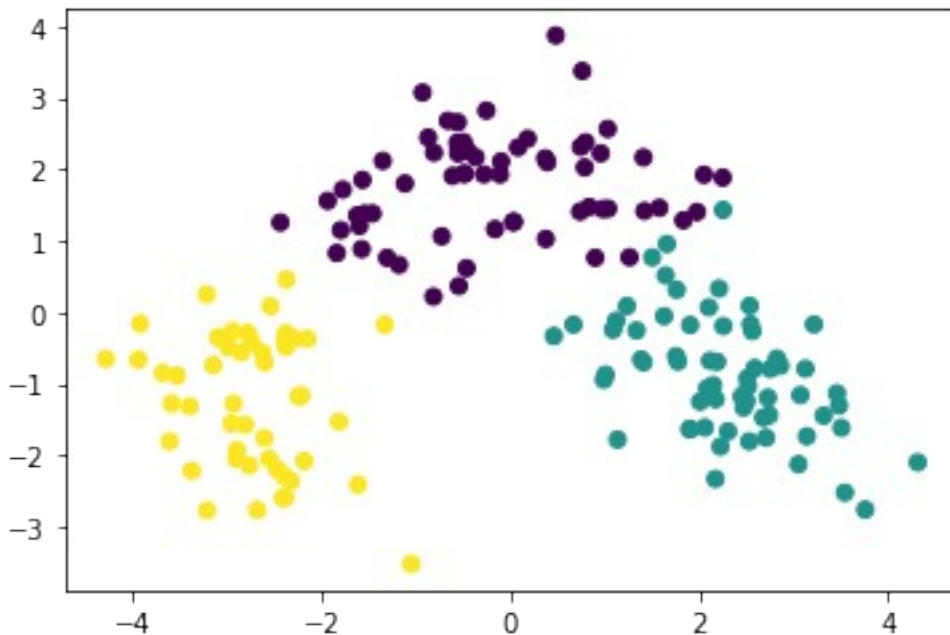


```
model = KMeans(n_clusters=3)
labels=model.fit_predict(PCA_components.iloc[:,2])

labels
array([[1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
      1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1,
1,
      1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0,
0,
      0, 0, 0, 0, 0, 1, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 2, 0, 0, 0,
0,
      0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
0,
      0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 0, 0, 0, 0, 0, 0, 0, 0, 0,
2,
      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
      2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,
2,
      2, 2])
```

Step-7: Visualizing the Clusters

```
plt.scatter(PCA_components[0], PCA_components[1], c=labels)
plt.show()
```



Step-8: Comparing actual Cluster No. with predicted Cluster No

```
df_r=pd.DataFrame({'Actual':y, 'Predicted':labels})
df_r.head()
```

	Actual	Predicted
0	1	1
1	1	1
2	1	1
3	1	1
4	1	1

weak8-Prepare a Classification model using decision tree Classifier.-

```
#Three lines to make our compiler able to draw:
import sys
import matplotlib
matplotlib.use('Agg')

import pandas as pd
from sklearn import tree
from sklearn.tree import DecisionTreeClassifier
```

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

```
df = pd.read_csv("DTree.csv")
```

```
print(df)
```

	Age	Experience	Rank	Nationality	Go
0	36	10	9	UK	NO
1	42	12	4	USA	NO
2	23	4	6	N	NO
3	52	4	4	USA	NO
4	43	21	8	USA	YES
5	44	14	5	UK	NO
6	66	3	7	N	YES
7	35	14	9	UK	YES
8	52	13	7	N	YES
9	35	5	9	N	YES
10	24	3	5	USA	NO
11	18	3	7	UK	YES
12	45	9	9	UK	YES

```
d = {'UK': 0, 'USA': 1, 'N': 2}
```

```
df['Nationality'] = df['Nationality'].map(d)
```

```
d = {'YES': 1, 'NO': 0}
```

```
df['Go'] = df['Go'].map(d)
```

```
print(df)
```

	Age	Experience	Rank	Nationality	Go
0	36	10	9	0	0
1	42	12	4	1	0
2	23	4	6	2	0
3	52	4	4	1	0
4	43	21	8	1	1
5	44	14	5	0	0
6	66	3	7	2	1
7	35	14	9	0	1
8	52	13	7	2	1
9	35	5	9	2	1
10	24	3	5	1	0
11	18	3	7	0	1
12	45	9	9	0	1

```
features = ['Age', 'Experience', 'Rank', 'Nationality']
```

```
X = df[features]
```

```
y = df['Go']
```

```
X
```

	Age	Experience	Rank	Nationality
0	36	10	9	0
1	42	12	4	1
2	23	4	6	2
3	52	4	4	1
4	43	21	8	1
5	44	14	5	0
6	66	3	7	2
7	35	14	9	0
8	52	13	7	2
9	35	5	9	2
10	24	3	5	1
11	18	3	7	0
12	45	9	9	0

y

0	0
1	0
2	0
3	0
4	1
5	0
6	1
7	1
8	1
9	1
10	0
11	1
12	1

Name: Go, dtype: int64

```
dtree = DecisionTreeClassifier()
dtree = dtree.fit(X, y)
```

```
tree.plot_tree(dtree, feature_names=features)
```

```
[Text(0.3333333333333333, 0.875, 'Rank <= 6.5\n'gini = 0.497\n'nsamples = 13\n'
nvalue = [6, 7]'),
  Text(0.16666666666666666, 0.625, 'gini = 0.0\n'nsamples = 5\n'
nvalue = [5, 0]'),
  Text(0.5, 0.625, 'Experience <= 9.5\n'gini = 0.219\n'nsamples = 8\n'
nvalue = [1, 7]'),
  Text(0.3333333333333333, 0.375, 'gini = 0.0\n'nsamples = 4\n'
nvalue = [0, 4]'),
  Text(0.6666666666666666, 0.375, 'Experience <= 11.5\n'gini = 0.375\n'
nsamples = 4\n'
nvalue = [1, 3]'),
  Text(0.5, 0.125, 'gini = 0.0\n'nsamples = 1\n'
nvalue = [1, 0]'),
  Text(0.8333333333333334, 0.125, 'gini = 0.0\n'nsamples = 3\n'
nvalue = [0, 3]')]
```

weak9-Prepare a Classification model using Navie Bayes Classifier-

Step-1:Assigning features and label variables

```
weather=['Sunny','Sunny','Overcast','Rainy','Rainy','Rainy','Overcast',
        'Sunny','Sunny',
        'Rainy','Sunny','Overcast','Overcast','Rainy']
temp=['Hot','Hot','Hot','Mild','Cool','Cool','Cool','Mild','Cool','Mild',
      'Mild','Mild','Hot','Mild']

play=['No','No','Yes','Yes','Yes','No','Yes','No','Yes','Yes','Yes','Yes',
      'Yes','Yes','No']
```

Step-2: Encoding the data

```
# Import LabelEncoder
from sklearn import preprocessing
#creating labelEncoder
le = preprocessing.LabelEncoder()
# Converting string labels into numbers.
weather_encoded=le.fit_transform(weather)
print(weather_encoded)

[2 2 0 1 1 1 0 2 2 1 2 0 0 1]

# Converting string labels into numbers
temp_encoded=le.fit_transform(temp)
label=le.fit_transform(play)
print("Temp:",temp_encoded)
print("Play:",label)

Temp: [1 1 1 2 0 0 0 2 0 2 2 2 1 2]
Play: [0 0 1 1 1 0 1 0 1 1 1 1 1 0]
```

Step-3:Combining weather and temp into single list of tuples

```
features=zip(weather_encoded,temp_encoded)
final=list(features)

final
[(2, 1),
 (2, 1),
 (0, 1),
 (1, 2),
```



```
(1, 0),  
(1, 0),  
(0, 0),  
(2, 2),  
(2, 0),  
(1, 2),  
(2, 2),  
(0, 2),  
(0, 1),  
(1, 2)]
```

Step-4: Fitting the model and predicting the classifier

```
#Import Gaussian Naive Bayes model  
from sklearn.naive_bayes import GaussianNB  
  
#Create a Gaussian Classifier  
model = GaussianNB()  
  
# Train the model using the training sets  
model.fit(final,label)  
  
#Predict Output  
predicted= model.predict([[0,0]]) # 0:Overcast, 2:Mild  
print("Predicted Value:", predicted)  
  
Predicted Value: [1]
```

#Here, 1 indicates that players can 'play'.

```
weather=['Sunny','Sunny','Overcast','Rainy','Rainy','Rainy','Overcast',  
, 'Sunny','Sunny',  
'Rainy','Sunny','Overcast','Overcast','Rainy']  
temp=['Hot','Hot','Hot','Mild','Cool','Cool','Cool','Mild','Cool','Mil  
d','Mild','Mild','Hot','Mild']  
  
play=['No','No','Yes','Yes','Yes','No','Yes','No','Yes','Yes','Yes','Y  
es','Yes','No']  
# Import LabelEncoder  
from sklearn import preprocessing  
#creating labelEncoder  
le = preprocessing.LabelEncoder()  
# Converting string labels into numbers.  
weather_encoded=le.fit_transform(weather)  
print("Weather:",weather_encoded)  
# Converting string labels into numbers  
temp_encoded=le.fit_transform(temp)  
label=le.fit_transform(play)  
print("Temp:",temp_encoded)
```

```

print("Play:",label)
features=zip(weather_encoded,temp_encoded)
final=list(features)
print("Features are:", final)
#Import Gaussian Naive Bayes model
from sklearn.naive_bayes import GaussianNB

#Create a Gaussian Classifier
model = GaussianNB()

# Train the model using the training sets
model.fit(final,label)

#Predict Output
predicted= model.predict([[0,2]]) # 0:Overcast, 2:Mild
print("Predicted Value:", predicted)

```

```

Weather: [2 2 0 1 1 1 0 2 2 1 2 0 0 1]
Temp: [1 1 1 2 0 0 0 2 0 2 2 2 1 2]
Play: [0 0 1 1 1 0 1 0 1 1 1 1 1 0]
Features are: [(2, 1), (2, 1), (0, 1), (1, 2), (1, 0), (1, 0), (0, 0),
(2, 2), (2, 0), (1, 2), (2, 2), (0, 2), (0, 1), (1, 2)]
Predicted Value: [1]

```