

## Practical No.7

### **Aim:-** Logistic Regression and Decision Tree

(A)

- Build a Logistic Regression Model to predict a binary outcome.
- Evaluate the model's performance using classification metrics.

Logistic Regression

Import the Libraries

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

Importing the dataset

```
[6]: dataset = pd.read_csv('Social_Network_Ads.csv')
X = dataset.iloc[:, :-1].values
Y = dataset.iloc[:, -1].values
```

Splitting the Dataset

```
[8]: from sklearn.model_selection import train_test_split
X_train, X_test, Y_train, Y_test = train_test_split(X, Y, test_size=0.25, random_state=0)
```

```
[9]: print(X_train)
print(Y_train)
```

```
[[ 44 39000]
 [ 32 120000]
 [ 38 50000]
 [ 32 135000]
```

Feature Scaling

```
[10]: from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)
```

Training the Logistic Regression Model

```
[11]: from sklearn.linear_model import LogisticRegression
classifier = LogisticRegression(random_state=0)
classifier.fit(X_train, Y_train)
```

```
[11]: * LogisticRegression 1 2
LogisticRegression(random_state=0)
```

Predicting a New Result

```
[12]: # Predicting result for certain age and salary, person will purchase or not purchase  
print(classifier.predict(sc.transform([[30, 87000]])))
```

[0]

Predicting the test results

```
[15]: y_pred = classifier.predict(X_test)  
print(np.concatenate(  
      (y_pred.reshape(len(y_pred), 1), Y_test.reshape(len(Y_test), 1)), 1))
```

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

Making the Confusion Matrix

```
[16]: from sklearn.metrics import confusion_matrix, accuracy_score  
cm = confusion_matrix(Y_test, y_pred)  
print(cm)  
accuracy = accuracy_score(Y_test, y_pred)  
print(accuracy)
```

```
[[65  3]  
 [ 8 24]]  
0.89
```

[B]

**Aim :-** Construct a Decision Tree model and interpret rules for classification.



```
In [12]: #Accuracy of the model
print("Accuracy:", accuracy_score(y_test, Y_pred))

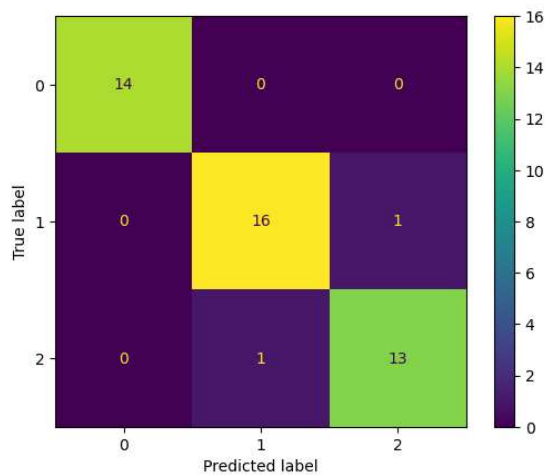
Accuracy: 0.9555555555555556
```

```
In [15]: from sklearn.metrics import confusion_matrix
cm = np.array(confusion_matrix(y_test, Y_pred))
cm
```

```
Out[15]: array([[14,  0,  0],
               [ 0, 16,  1],
               [ 0,  1, 13]], dtype=int64)
```

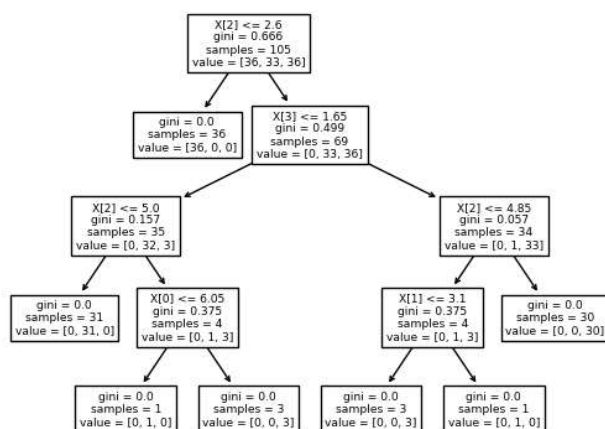
```
In [16]: from sklearn.metrics import plot_confusion_matrix
plot_confusion_matrix(clf, X_test, y_test)
plt.show()
```

C:\ProgramData\Anaconda3\lib\site-packages\sklearn\utils\deprecation.py:87: FutureWarning: Function plot\_confusion\_matrix is deprecated; Function 'plot\_confusion\_matrix' is deprecated in 1.0 and will be removed in 1.2. Use one of the class methods: ConfusionMatrixDisplay.from\_predictions or ConfusionMatrixDisplay.from\_estimator.  
warnings.warn(msg, category=FutureWarning)



```
In [17]: tree.plot_tree(clf)
```

```
Out[17]: [Text(0.4, 0.9, 'X[2] <= 2.6\ngini = 0.666\nsamples = 105\nvalue = [36, 33, 36]'),
Text(0.3, 0.7, 'gini = 0.0\nsamples = 36\nvalue = [36, 0, 0]'),
Text(0.5, 0.7, 'X[3] <= 1.65\ngini = 0.499\nsamples = 69\nvalue = [0, 33, 36]'),
Text(0.2, 0.5, 'X[2] <= 5.0\ngini = 0.157\nsamples = 35\nvalue = [0, 32, 3]'),
Text(0.1, 0.3, 'gini = 0.0\nsamples = 31\nvalue = [0, 31, 0]'),
Text(0.3, 0.3, 'X[0] <= 6.05\ngini = 0.375\nsamples = 4\nvalue = [0, 1, 3]'),
Text(0.2, 0.1, 'gini = 0.0\nsamples = 1\nvalue = [0, 1, 0]'),
Text(0.4, 0.1, 'gini = 0.0\nsamples = 3\nvalue = [0, 0, 3]'),
Text(0.8, 0.5, 'X[2] <= 4.85\ngini = 0.057\nsamples = 34\nvalue = [0, 1, 33]'),
Text(0.7, 0.3, 'X[1] <= 3.1\ngini = 0.375\nsamples = 4\nvalue = [0, 1, 3]'),
Text(0.6, 0.1, 'gini = 0.0\nsamples = 3\nvalue = [0, 0, 3]'),
Text(0.8, 0.1, 'gini = 0.0\nsamples = 1\nvalue = [0, 1, 0]'),
Text(0.9, 0.3, 'gini = 0.0\nsamples = 30\nvalue = [0, 0, 30]')]
```



## Practical No.8

### **Aim:-** K-Means Clustering

- Apply the K-Means Algorithm to group similar data points into clusters.
- Determine optimal number of clusters using elbow method or silhouette analysis.
- Visualize the clustering results and analyse the cluster characteristics.

```
In [7]: #Importing Libraries
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import sklearn.cluster as cluster
from sklearn.cluster import KMeans
import seaborn as sns
import sklearn.metrics as metrics
```

```
In [8]: dataset = pd.read_csv('Iris.csv')
x = dataset.iloc[:,[0,1,2,3]].values
```

```
In [9]: print(x)
```

```
[[5.1 3.5 1.4 0.2]
 [4.9 3.  1.4 0.2]
 [4.7 3.2 1.3 0.2]
 [4.6 3.1 1.5 0.2]
 [5.  3.6 1.4 0.2]
 [5.4 3.9 1.7 0.4]
 [4.6 3.4 1.4 0.3]
 [5.  3.4 1.5 0.2]
 [4.4 2.9 1.4 0.2]
 [4.9 3.1 1.5 0.1]]
```

```
In [10]: K = range(1,10)
# within-cluster-sum-of-square
wss = []
for k in K:
    kmeans=cluster.KMeans(n_clusters=k,init="k-means++")
    kmeans=kmeans.fit(x)
    wss_iter = kmeans.inertia_
    wss.append(wss_iter)
```

```
C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cluster\_kmeans.py
on Windows with MKL, when there are less chunks than available thread
P_NUM_THREADS=1.
warnings.warn(
```