1. Aim: Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples. Read the training data from a .CSV file.

# **Program:**

```
import pandas as pd
x=pd.read_csv(r"C:\Users\SRAVANI\Downloads\Income.csv")
print(x.head())
S=[]
for i in range(len(x)):
  y=list(x.iloc[i])
  if(y[-1]==1):
    if len(s) == 0:
       s=y[:len(y)-1]
    else:
       for j in range(len(s)):
         if y[j]!=s[j]:
            s[i]="?"
print(s)
OUTPUT:
                                                          S
```

	Age	Income	Assets	Height	Qualification	Status
0	Y	High	1	2	3	1
1	Y	Med	0	3	2	1
2	Y	High	1	1	2	1
3	Y	Med	1	1	2	0
4	M	Low	0	1	1	0

['?', '?', '?', '?', '?']

2. Aim: For a given set of training data examples stored in a .CSV file, implement and demonstrate the Candidate-Elimination algorithm to output a description of the set of all hypotheses consistent with the training examples

# **Program:**

```
import numpy as np
import pandas as pd
data = pd.DataFrame(data=pd.read_csv(r'C:\Users\Downloads\ws.csv'))
```

```
concepts = np.array(data.iloc[:,0:-1])
print(concepts)
target = np.array(data.iloc[:,-1])
print(target)
def learn(concepts, target):
  specific_h = concepts[0].copy()
  print("\nInitialization of specific_h and general_h")
  print(specific_h)
  general_h = [["?" for i in range(len(specific_h))] for i in range(len(specific_h))]
  print(general_h)
  for i, h in enumerate(concepts):
     if target[i] == "Yes":
       for x in range(len(specific_h)):
          if h[x] != specific_h[x]:
             specific h[x] = '?'
             general h[x][x] = '?'
     if target[i] == "No":
       for x in range(len(specific_h)):
          if h[x] != specific_h[x]:
             general_h[x][x] = specific_h[x]
          else:
             general_h[x][x] = '?'
     print("\nSteps of Candidate Elimination Algorithm",i+1)
     print(specific_h)
     print(general h)
  indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
  for i in indices:
     general_h.remove(['?', '?', '?', '?', '?', '?'])
  return specific_h, general_h
s_final, g_final = learn(concepts, target)
print("\nFinal Specific_h:", s_final, sep="\n")
print("\nFinal General_h:", g_final, sep="\n")
```

```
OUTPUT:
```

```
[['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
['Rainy' 'Cold' 'High' 'Strong' 'Warm' 'Change']
['Sunny' 'Warm' 'High' 'Strong' 'Cool' 'Change']]
['Yes' 'No' 'Yes']
Initialization of specific_h and general_h
['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
'?'], ['?', '?', '?', '?', '?', '?']]
Steps of Candidate Elimination Algorithm 1
['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
'?'], ['?', '?', '?', '?', '?', '?']]
Steps of Candidate Elimination Algorithm 2
['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
[['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?'], ['?', '?', '?'], ['?', '?']
, '?', '?', '?', '?'], ['?', '?', '?', '?', '?', 'Same']]
Steps of Candidate Elimination Algorithm 3
['Sunny' 'Warm' 'High' 'Strong' '?' '?']
[['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?'], ['?', '?', '?'], ['?', '?']
, '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]
Final Specific h:
['Sunny' 'Warm' 'High' 'Strong' '?' '?']
Final General_h:
[['Sunny', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?', '?']]
```

**3. Aim:** Write a program to demonstrate the working of the decision tree based ID3 algorithm. Use an appropriate data set for building the decision tree and apply this knowledge to classify a new sample.

## **Program:**

import pandas as pd

from sklearn import tree

from sklearn.preprocessing import LabelEncoder

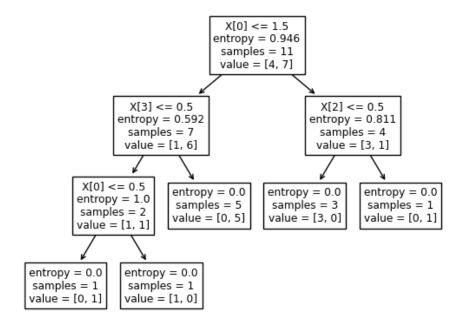
from sklearn.tree import DecisionTreeClassifier

df=pd.read csv(r"C:\Users\ADI SESHU\Downloads\play tennis.csv")

df = df.drop('day',axis=1)

```
from sklearn.preprocessing import LabelEncoder
Le = LabelEncoder()
df['outlook'] = Le.fit_transform(df['outlook'])
df['temp'] = Le.fit_transform(df['temp'])
df['humidity'] = Le.fit_transform(df['humidity'])
df['wind'] = Le.fit_transform(df['wind'])
df['play'] = Le.fit_transform(df['play'])
x = df.drop(['play'],axis=1)
y= df['play']
from sklearn.model_selection import train_test_split
X_train,X_test,y_train,y_test = train_test_split(x,y,random_state=20,test_size=0.2)
# Fitting the model
from sklearn import tree
dt = tree.DecisionTreeClassifier(criterion = 'entropy')
dt = dt.fit(X_train, y_train)
dt
dt.score(X_train,y_train)
y_pred = dt.predict(X_test)
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test,y_pred)
from sklearn.metrics import accuracy_score
print("Accuracy : ", accuracy_score(y_test,y_pred))
new_df = pd.DataFrame({'outlook': 2,
              'temp': 1,
              'humidity':0,
              'wind': 1},index=[0])
y_pred1 = dt.predict(new_df)
y_pred1
tree.plot_tree(dt)
```

# **Output:**



**4. Aim:** Exercises to solve the real-world problems using the following machine learning methods: a) Linear Regression b) Logistic Regression c) Binary Classifier

## **Program:**

```
#3.linear regression
```

import numpy as np

import matplotlib.pyplot as plt

import pandas as pd

# Importing the datasets

datasets = pd.read\_csv(r"C:\Users\ADI SESHU\Downloads\Salary\_Data (1).csv")

X = datasets.iloc[:, :-1].values

Y = datasets.iloc[:, 1].values

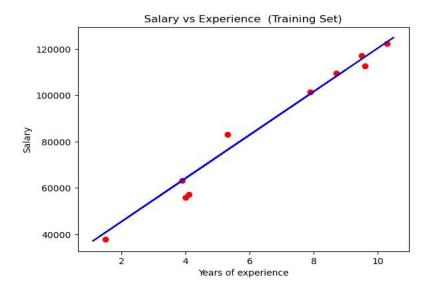
# Splitting the dataset into the Training set and Test set

from sklearn.model\_selection import train\_test\_split

X\_Train, X\_Test, Y\_Train, Y\_Test = train\_test\_split(X, Y, test\_size = 1/3, random\_state = 0)

```
# Fitting Simple Linear Regression to the training set
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_Train, Y_Train)
# Predicting the Test set result
Y_Pred = regressor.predict(X_Test)
# Visualising the Training set results
plt.scatter(X_Train, Y_Train, color = 'red')
plt.plot(X_Train, regressor.predict(X_Train), color = 'blue')
plt.title('Salary vs Experience (Training Set)')
plt.xlabel('Years of experience')
plt.ylabel('Salary')
plt.show()
# Visualising the Test set results
plt.scatter(X_Test, Y_Test, color = 'red')
plt.plot(X_Train, regressor.predict(X_Train), color = 'blue')
plt.title('Salary vs Experience (Training Set)')
plt.xlabel('Years of experience')
plt.ylabel('Salary')
plt.show()
```

# **Output:**





from sklearn.preprocessing import StandardScaler

 $sc_X = StandardScaler()$ 

 $X_{\text{Train}} = \text{sc}_X.\text{fit}_{\text{transform}}(X_{\text{Train}})$ 

 $X_Test = sc_X.transform(X_Test)$ 

from sklearn.linear\_model import LogisticRegression

 $classifier = LogisticRegression(random\_state = 0)$ 

classifier.fit(X\_Train, Y\_Train)

# Predicting the test set results

Y\_Pred = classifier.predict(X\_Test)

# Making the Confusion Matrix

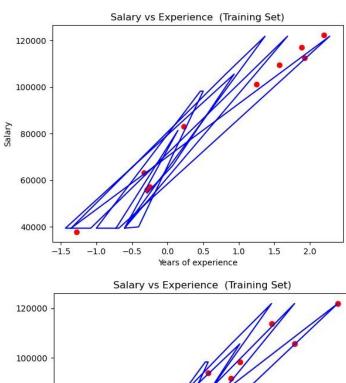
from sklearn.metrics import confusion\_matrix

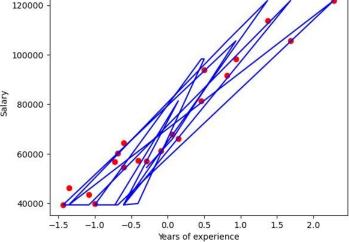
 $cm = confusion\_matrix(Y\_Test, Y\_Pred)$ 

plt.scatter(X\_Train, Y\_Train, color = 'red')

```
plt.plot(X_Train, classifier.predict(X_Train), color = 'blue')
plt.title('Salary vs Experience (Training Set)')
plt.xlabel('Years of experience')
plt.ylabel('Salary')
plt.show()
# Visualising the Test set results
plt.scatter(X_Test, Y_Test, color = 'red')
plt.plot(X_Train, classifier.predict(X_Train), color = 'blue')
plt.title('Salary vs Experience (Training Set)')
plt.xlabel('Years of experience')
plt.ylabel('Salary')
plt.show()
```

# **Output:**





# 5. Aim: Develop a program for Bias, Variance, Remove duplicates, Cross Validation

## **Program:**

```
from mlxtend.evaluate import bias variance decomp
from sklearn.model_selection import train_test_split
import pandas as pd
import matplotlib.pyplot as plt
import numpy as np
from sklearn.linear_model import LinearRegression, Lasso
import warnings
warnings.filterwarnings('ignore')
#We will load the Boston house dataset for our example
from sklearn.datasets import load_boston
from sklearn import metrics
# From the library load the necessary rows & columns
X, y = load\_boston(return\_X\_y=True)
# Split the dataset into train and test
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.33, random_state=1)
# Model definition
model_lr = LinearRegression()
# Estimation of bias and variance using bias_variance_decomp
#Note here we are using loss as 'mse' and setting default bootstrap num_rounds to 200
mse, bias, var = bias_variance_decomp(model_lr, X_train, y_train, X_test, y_test, loss='mse', num_rounds=200,
random_seed=123)
y_pred=model_lr.predict(X_test)
# summarize results
print('MSE from bias_variance lib [avg expected loss]: %.3f' % mse)
print('Avg Bias: %.3f' % bias)
print('Avg Variance: %.3f' % var)
print('Mean Square error by Sckit-learn lib: %.3f' % metrics.mean_squared_error(y_test,y_pred))
```

# output:

MSE from bias\_variance lib [avg expected loss]: 22.128

Avg Bias: 20.522 Avg Variance: 1.606

Mean Square error by Sckit-learn lib: 22.069

# **6. Aim:** write a program to implement categorical encoding, One-hot Encoding

# **Program:**

```
print("original data")
print(df1)
from sklearn.preprocessing import LabelEncoder
encoder=LabelEncoder()
df1['team']=encoder.fit_transform(df1['team'])
print("after encoding")
print(df1)
output:
original data
 team points
0 A
        25
        12
  A
2 B
        15
3 B
        14
  В
        19
  В
       23
5
  C
        25
  C
       29
after encoding
 team points
   0
        25
        12
   0
        15
   1
3
   1
        14
4
   1
        19
   1
        23
   2
        25
        29
```

# **Program:**

import pandas as pd

```
print("original data")
print(df)
from sklearn.preprocessing import OneHotEncoder
#creating instance of one-hot-encoder
encoder = OneHotEncoder(handle_unknown='ignore')
#perform one-hot encoding on 'team' column
encoder_df = pd.DataFrame(encoder.fit_transform(df[['team']]).toarray())
#merge one-hot encoded columns back with original DataFrame
final_df = df.join(encoder_df)
final_df.drop('team', axis=1, inplace=True)
#rename columns
final_df.columns = ['points', 'teamA', 'teamB', 'teamC']
#view final df
print("after one hot encoding")
print(final_df)
output:
original data
 team points
  Α
        25
        12
   Α
   В
        15
3
  В
        14
4
  В
        19
5
  В
        23
6 C
        25
7
        29
after one hot encoding
 points teamA teamB teamC
    25
        1.0 0.0 0.0
1
    12
        1.0 0.0 0.0
2
    15
        0.0 1.0 0.0
3
    14
        0.0 1.0
                  0.0
4
    19
        0.0 1.0
                   0.0
    23
5
        0.0 1.0
    25
6
        0.0 0.0
                   1.0
    29
        0.0
             0.0
                   1.0
```

**7. Aim:** Build an Artificial Neural Network by implementing the Backpropagation algorithm and test the same using appropriate data sets.

## **Program:**

```
import numpy as np
X = \text{np.array}(([2, 9], [1, 5], [3, 6]), \text{dtype=float})
y = np.array(([92], [86], [89]), dtype=float)
X = X/np.amax(X,axis=0) #maximum of X array longitudinally
y = y/100
#Sigmoid Function
def sigmoid (x):
  return 1/(1 + np.exp(-x))
#Derivative of Sigmoid Function
def derivatives_sigmoid(x):
  return x * (1 - x)
#Variable initialization
epoch=5 #Setting training iterations
lr=0.1 #Setting learning rate
inputlayer_neurons = 2 #number of features in data set
hiddenlayer_neurons = 3 #number of hidden layers neurons
output_neurons = 1 #number of neurons at output layer
#weight and bias initialization
wh=np.random.uniform(size=(inputlayer_neurons,hiddenlayer_neurons))
bh=np.random.uniform(size=(1,hiddenlayer_neurons))
wout=np.random.uniform(size=(hiddenlayer_neurons,output_neurons))
bout=np.random.uniform(size=(1,output_neurons))
#draws a random range of numbers uniformly of dim x*y
for i in range(epoch):
  #Forward Propogation
  hinp1=np.dot(X,wh)
```

```
hinp=hinp1 + bh
  hlayer_act = sigmoid(hinp)
  outinp1=np.dot(hlayer_act,wout)
  outinp= outinp1+bout
  output = sigmoid(outinp)
  #Backpropagation
  EO = y-output
  outgrad = derivatives_sigmoid(output)
  d_output = EO * outgrad
  EH = d\_output.dot(wout.T)
  hiddengrad = derivatives_sigmoid(hlayer_act)#how much hidden layer wts contributed to error
  d_hiddenlayer = EH * hiddengrad
  wout += hlayer_act.T.dot(d_output) *lr # dotproduct of nextlayererror and currentlayerop
  wh += X.T.dot(d_hiddenlayer) *lr
  print ("------Epoch-", i+1, "Starts-----")
  print("Input: \n" + str(X))
  print("Actual Output: \n" + str(y))
  print("Predicted Output: \n" ,output)
  print ("-----Epoch-", i+1, "Ends-----\n")
print("Input: \n" + str(X))
print("Actual Output: \n'' + str(y))
print("Predicted Output: \n" ,output)
Training Examples:
Training Examples:
Example Sleep Study Expected % in Exams
         2
               9
                       92
```

1

2

1

5

86

3 3 6 89

Normalize the input:

Examp	le Sleep	Study	Expected % in Exams	
1	2/3 = 0.666666667	9/9 = 1	0.92	
2	1/3 = 0.333333333	5/9 = 0.5555556	0.86	
3	3/3 = 1	6/9 = 0.66666667	0.89	

# **Output:**

Epoch- 1 Starts——-

Input:

[[0.66666667 1.]

[0.33333333 0.55555556]

[1. 0.66666667]]

Actual Output:

[[0.92]]

[0.86]

[0.89]]

Predicted Output:

[[0.81951208]

[0.8007242]

[0.82485744]]

——Epoch- 1 Ends——-

——Epoch- 2 Starts——

Input:

[[0.66666667 1.]

[0.33333333 0.55555556]

[1. 0.66666667]]

Actual Output:

[[0.92]

[0.86]
[0.89]]
Predicted Output:
[[0.82033938]
[0.80153634]
[0.82568134]]
——Epoch- 2 Ends——-
——Epoch- 3 Starts——-
Input:
[[0.66666667 1.]
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.82115226]
[0.80233463]
[0.82649072]]
——Epoch- 3 Ends——-
Epoch- 4 Starts——-
Input:
[[0.66666667 1.]
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]

Predicted Output:
[[0.82195108]
[0.80311943]
[0.82728598]]
——Epoch- 4 Ends——-
——Epoch- 5 Starts——
Input:
[[0.66666667 1. ]
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.8227362]
[0.80389106]
[0.82806747]]
——Epoch- 5 Ends——-
Input:
[[0.66666667 1.]
[0.33333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.8227362]
[0.80389106]

**8. Aim:** Write a program to implement k-Nearest Neighbor algorithm to classify the iris data set. Print both correct and wrong predictions.

# **Program:**

```
import pandas as pd
from sklearn.metrics import confusion matrix, accuracy score, classification report
from sklearn.preprocessing import LabelEncoder
ds = pd.read\_csv(r"D:\ML LAB 3-2\IRIS.csv")
encoder=LabelEncoder()
ds['species']=encoder.fit_transform(ds['species'])
x=ds.iloc[:,:-1].values
y=ds.iloc[:,-1].values
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=1)
from sklearn.neighbors import KNeighborsClassifier
clf=KNeighborsClassifier(n_neighbors=8)
clf.fit(x_train,y_train)
y_pred=clf.predict(x_test)
print("Accuracy: ", accuracy_score (y_test, y_pred))
print("Confusion Matrix :\n", confusion_matrix (y_test, y_pred))
print("Classification Report :\n", classification_report (y_test, y_pred))
```

## **Output:**

```
Accuracy: 0.97777777777777
Confusion Matrix :
[[14 0 0]
[ 0 17 1]
[0 0 13]]
Classification Report :
             precision
                        recall f1-score support
          0
                 1.00
                          1.00
                                    1.00
                                               14
                                               18
          1
                 1.00
                          0.94
                                    0.97
                 0.93
                          1.00
                                    0.96
                                               13
                                    0.98
                                               45
   accuracy
                 0.98
                          0.98
                                    0.98
                                               45
  macro avg
weighted avg
                 0.98
                           0.98
                                    0.98
                                               45
```

**9. Aim:** Implement the non-parametric Locally Weighted Regression algorithm in order to fit data points. Select appropriate data set for your experiment and draw graphs.

```
Program:-
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
% matplotlib inline
df = pd.read_csv('tips.csv')
features = np.array(df.total_bill)
labels = np.array(df.tip)
def kernel(data, point, xmat, k):
 m,n = np.shape(xmat)
 ws = np.mat(np.eye((m)))
 for j in range(m):
   diff = point - data[j]
   ws[j,j] = np.exp(diff*diff.T/(-2.0*k**2))
 return ws
def local_weight(data, point, xmat, ymat, k):
 wei = kernel(data, point, xmat, k)
 return (data.T*(wei*data)).I*(data.T*(wei*ymat.T))
def local_weight_regression(xmat, ymat, k):
 m,n = np.shape(xmat)
 ypred = np.zeros(m)
 for i in range(m):
   ypred[i] = xmat[i]*local_weight(xmat, xmat[i],xmat,ymat,k)
 return ypred
m = features.shape[0]
```

mtip = np.mat(labels)

```
data = np.hstack((np.ones((m, 1)), np.mat(features).T))

ypred = local_weight_regression(data, mtip, 0.5)

indices = data[:,1].argsort(0)

xsort = data[indices][:,0]

fig = plt.figure()

ax = fig.add_subplot(1,1,1)

ax.scatter(features, labels, color='blue')

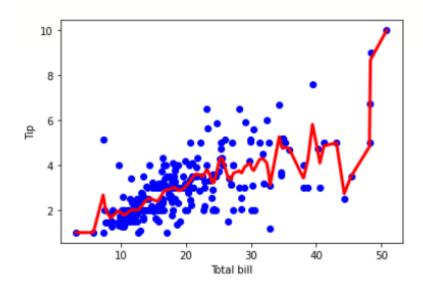
ax.plot(xsort[:,1],ypred[indices], color = 'red', linewidth=3)

plt.xlabel('Total bill')

plt.ylabel('Tip')

plt.show()
```

# Output:-



**10. Aim:** To Classify a set of documents using naive\_bayes classifier

# **Program:**

```
import numpy as np
import pandas as pd
data=pd.read_csv(r'D:\text_dataset.csv')
display(data.head())
print('The dimensions of the dataset',data.shape)
```

#### O/P:

	S.NO	Text Documents	Label
0	1	I love this sandwich	pos
1	2	This is an amazing place	pos
2	3	I feel very good about these beers	pos
3	4	This is my best work	pos
4	5	What an awesome view	pos

The dimensions of the dataset (18, 3)

data['Label']=data.Label.map({'pos':1,'neg':0})
display(data.head())

# **Output:**

	S.NO	Text Documents	Label
0	1	I love this sandwich	1
1	2	This is an amazing place	1
2	3	I feel very good about these beers	1
3	4	This is my best work	1
4	5	What an awesome view	1

x=data['Text Documents']

y=data['Label']

from sklearn.model\_selection import train\_test\_split

x\_train,x\_test,y\_train,y\_test=train\_test\_split(x,y,test\_size=0.2,random\_state=42)

print ('\n the total number of Training Data :',x\_train.shape)

print ('\n the total number of Test Data :',x\_test.shape)

## **Output:**

the total number of Training Data: (14,)

the total number of Test Data: (4,)

from sklearn.feature\_extraction.text import CountVectorizer

cv = CountVectorizer()

xtrain\_dtm = cv.fit\_transform(x\_train)

xtest\_dtm=cv.transform(x\_test)

print('\n The words or Tokens in the text documents \n')

```
print(cv.get_feature_names())
df=pd.DataFrame(xtrain_dtm.toarray(),columns=cv.get_feature_names())
```

## **Output:**

```
The words or Tokens in the text documents
['about', 'am', 'an', 'and', 'awesome', 'bad', 'beers', 'best', 'boss', 'can', 'dance', 'deal', 'do', 'enemy', 'feel', 'fun', 'good', 'great', 'have', 'holiday', 'horrible', 'house', 'is', 'juice', 'like', 'locality', 'love', 'my', 'not', 'of', 'place', 'sick', 'stay', 'stuff', 'taste', 'that', 'the', 'these', 'this', 'tired', 'to', 'today', 'tomorrow', 'very', 'view', 'we nt', 'what', 'will', 'with', 'work']
from sklearn.naive bayes import MultinomialNB
clf = MultinomialNB().fit(xtrain_dtm,y_train)
predicted = clf.predict(xtest_dtm)
from sklearn import metrics
print('\n Accuracy of the classifier is',metrics.accuracy_score(y_test,predicted))
print('\n Confusion matrix')
print(metrics.confusion_matrix(y_test,predicted))
print(\n The value of Precision', metrics.precision_score(y_test,predicted))
print('\n The value of Recall', metrics.recall_score(y_test,predicted))
```

## **Output:**

Accuracy of the classifier is 1.0

Confusion matrix [[2 0] [0 2]]

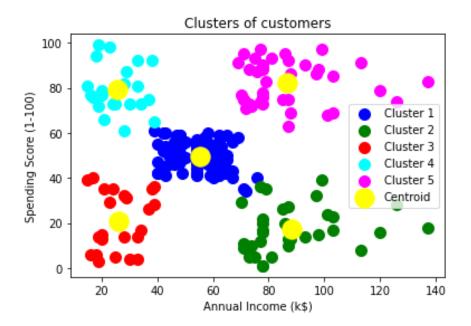
The value of Precision 1.0

The value of Recall 1.0

11. Aim: Apply EM algorithm to cluster a Heart Disease Data Set. Use the same data set for cluster ing using k-Means algorithm. Compare the results of these two algorithms and comment on the qualit y of clustering. You can add Java/Python ML library classes/API in the program. Program:

import numpy as nm import matplotlib.pyplot as plt from sklearn.cluster import KMeans

```
import pandas as pd
dataset = pd.read csv(r"C:\Users\mmuni\Downloads\archive (1)\Mall Customers.csv")
x = dataset.iloc[:, [3, 4]].values
#training the K-means model on a dataset
kmeans = KMeans(n clusters=5, init='k-means++', random state= 42)
y predict= kmeans.fit predict(x)
plt.scatter(x[y predict == 0, 0], x[y predict == 0, 1], s = 100, c = 'blue', label = 'Cluster 1') #for first
plt.scatter(x[y predict == 1, 0], x[y predict == 1, 1], s = 100, c = 'green', label = 'Cluster 2') #for second
cluster
plt.scatter(x[y_predict== 2, 0], x[y_predict == 2, 1], s = 100, c = 'red', label = 'Cluster 3') #for third
cluster
plt.scatter(x[y predict == 3, 0], x[y predict == 3, 1], s = 100, c = 'cyan', label = 'Cluster 4') #for fourth
plt.scatter(x[y_predict == 4, 0], x[y_predict == 4, 1], s = 100, c = 'magenta', label = 'Cluster 5') #for
fifth cluster
plt.scatter(kmeans.cluster centers [:, 0], kmeans.cluster centers [:, 1], s = 300, c = 'yellow', label =
'Centroid')
plt.title('Clusters of customers')
plt.xlabel('Annual Income (k$)')
plt.ylabel('Spending Score (1-100)')
plt.legend()
plt.show()
output:
```



# 12. AIM:

Exploratory Data Analysis for Classification using Pandas or Matplotlib.

# **Program:**

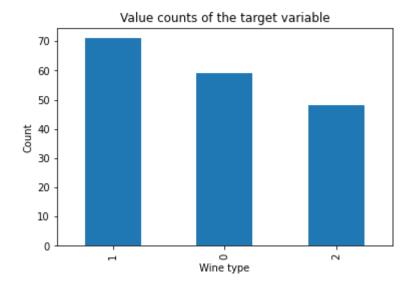
df.info()

```
# data manipulation
import pandas as pd
import numpy as np
# data viz
import matplotlib.pyplot as plt
import seaborn as sns
# use sklearn to import a dataset
from sklearn.datasets import load_wine
wine = load_wine()
df = pd.DataFrame(data=wine.data, columns=wine.feature_names)
df["target"] = wine.target
#display(df.head(2))
#display(df.tail(2))
print("shape",df.shape)
print("********")
print("information about dataset")
```

```
print("********")
print("Number of duplicate rows",df.duplicated().sum())
print("********")
print("count of target variables \n",df.target.value_counts())
df.target.value_counts().plot(kind="bar")
plt.title("Value counts of the target variable")
plt.xlabel("Wine type")
plt.ylabel("Count")
plt.show() # visualization
print("Skewness:", df['magnesium'].skew())
print("Kurtosis:",df['magnesium'].kurt())#statistical tools
OUTPUT:
shape (178, 14)
*****
information about dataset
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 178 entries, 0 to 177
Data columns (total 14 columns):
# Column
                        Non-Null Count Dtype
--- -----
                     _____
0 alcohol
                       178 non-null float64
                        178 non-null float64
1 malic_acid
2 ash
                     178 non-null float64
3 alcalinity_of_ash
                          178 non-null float64
4 magnesium
                         178 non-null float64
5 total_phenols
                         178 non-null float64
                        178 non-null float64
6 flavanoids
7 nonflavanoid_phenols
                             178 non-null float64
8 proanthocyanins
                          178 non-null float64
9 color intensity
                         178 non-null float64
10 hue
                      178 non-null float64
11 od280/od315_of_diluted_wines 178 non-null float64
                       178 non-null float64
12 proline
13 target
                      178 non-null int32
dtypes: float64(13), int32(1)
memory usage: 18.9 KB
******
Number of duplicate rows 0
******
count of target variables
   71
   59
0
```

2 48

Name: target, dtype: int64



**13.Aim:** Write a Python program to construct a Bayesian network considering medical data. Use this model to demonstrate the diagnosis of heart patients using standard Heart Disease Data Set

# **Program:**

import numpy as np

import pandas as pd

import csv

 $from\ pgmpy.estimators\ import\ Maximum Likelihood Estimator$ 

from pgmpy.models import BayesianModel

from pgmpy.inference import VariableElimination

```
heartDisease = pd.read\_csv('heart.csv')
```

heartDisease = heartDisease.replace('?',np.nan)

print('Sample instances from the dataset are given below')

print(heartDisease.head())

print('\n Attributes and datatypes')

print(heartDisease.dtypes)

```
model=
BayesianModel([('age', 'heartdisease'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'he
heartdisease','restecg'),('heartdisease','chol')])
print('\nLearning CPD using Maximum likelihood estimators')
model.fit(heartDisease,estimator=MaximumLikelihoodEstimator)
print('\n Inferencing with Bayesian Network:')
HeartDiseasetest_infer = VariableElimination(model)
print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)
print('\n 2. Probability of HeartDisease given evidence= cp ')
q2=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

# **Output:**

Learning CPD using Max Inferencing with Baye	simum likelihood estimat	tors 2.	-	rtDisease given evidenc	e= cp
3	artDisease given evidenc	ce= restecg	heartdisease	++   phi(heartdisease)   +	
heartdisease	++   phi(heartdisease)		heartdisease(0)		
heartdisease(0)	0.1012		heartdisease(1)	0.2159	
heartdisease(1)			+   heartdisease(2)	++   0.1373	
heartdisease(2)	0.2392		+	++   0.1537	
heartdisease(3)	0.2015			++	+
heartdisease(4)			heartdisease(4)	0.1321	

# 14. Aim: WRITE A PROGRAM TO IMPLEMENT SUPPORT VECTOR MACHINE

## **PROGRAM:**

from sklearn import tree from sklearn.model\_selection import train\_test\_split import pandas as pd

df=pd.read\_csv(r"C:\Users\Downloads\heart (1).csv")

df.head()

	age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	са	thal	target
0	52	1	0	125	212	0	1	168	0	1.0	2	2	3	0
1	53	1	0	140	203	1	0	155	1	3.1	0	0	3	0
2	70	1	0	145	174	0	1	125	1	2.6	0	0	3	0
3	61	1	0	148	203	0	1	161	0	0.0	2	1	3	0
4	62	0	0	138	294	1	1	106	0	1.9	1	3	2	0

y=df['target']

x=df.drop("target",axis=1)

x\_train, x\_test, y\_train, y\_test = train\_test\_split(x, y, test\_size=0.3,random\_state=1)

## **#USING KERNAL="LINEAR"**

from sklearn.svm import SVC

clf=SVC(kernel="linear")

model=clf.fit(x\_train,y\_train)

y\_pred= model.predict(x\_test)

import seaborn as sns

import matplotlib.pyplot as plt

sns.regplot(y\_pred,y\_test)

from sklearn.metrics import classification\_report,confusion\_matrix,accuracy\_score

 $result = confusion\_matrix(y\_test, y\_pred)$ 

print("Confusion Matrix:")

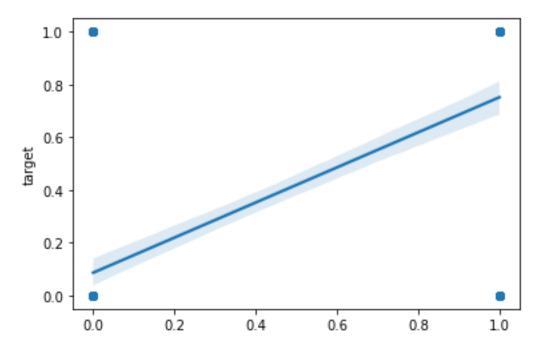
print(result)

print("Accuracy=",accuracy\_score(y\_test,y\_pred))

print("Classification Report")

print(classification\_report(y\_test,y\_pred))

## **OUTPUT:**



Confusion Matrix:

[[116 45]

[11 136]]

Accuracy= 0.81818181818182

Classification Report

precision recall f1-score support

0 0.91 0.72 0.81 161 1 0.75 0.93 0.83 147

accuracy 0.82 308 macro avg 0.83 0.82 0.82 308 weighted avg 0.84 0.82 0.82 308

## **#USING KERNAL="SIGMOID"**

from sklearn.svm import SVC  $\,$ 

clf=SVC(kernel="sigmoid")

model=clf.fit(x\_train,y\_train)

y\_pred= model.predict(x\_test)

import seaborn as sns

import matplotlib.pyplot as plt

sns.regplot(y\_pred,y\_test)

from sklearn.metrics import classification\_report,confusion\_matrix,accuracy\_score

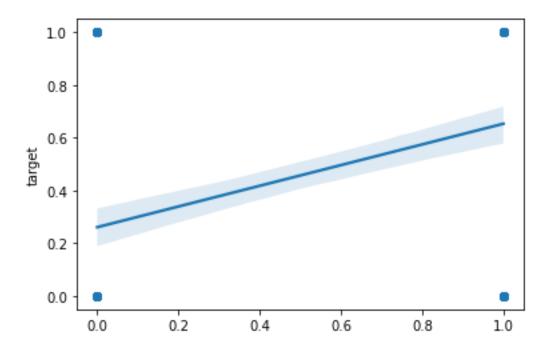
result=confusion\_matrix(y\_test,y\_pred)

print("Confusion Matrix:")

print(result)

print("Accuracy=",accuracy\_score(y\_test,y\_pred))
print("Classification Report")
print(classification\_report(y\_test,y\_pred))

## **OUTPUT:**



Confusion Matrix:

[[102 59]

[ 36 111]]

Accuracy= 0.6915584415584416

Classification Report

precision recall f1-score support

0 0.74 0.63 0.68 161 1 0.65 0.76 0.70 147

accuracy 0.69 308 macro avg 0.70 0.69 0.69 308 weighted avg 0.70 0.69 0.69 308

## **#USING KERNAL="POLY"**

from sklearn.svm import SVC

clf=SVC(kernel="poly")

model=clf.fit(x\_train,y\_train)

y\_pred= model.predict(x\_test)

import seaborn as sns

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

sns.regplot(y\_pred,y\_test)

 $from\ sklearn.metrics\ import\ classification\_report, confusion\_matrix, accuracy\_score$ 

result=confusion\_matrix(y\_test,y\_pred)

print("Confusion Matrix:")

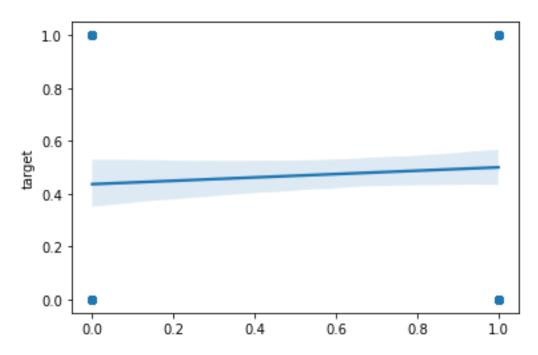
print(result)

print("Accuracy=",accuracy\_score(y\_test,y\_pred))

print("Classification Report")

print(classification\_report(y\_test,y\_pred))

## **OUTPUT:**



## Confusion Matrix:

[[62 99]

[48 99]]

Accuracy= 0.5227272727272727

Classification Report

precision recall f1-score support

0 0.56 0.39 0.46 161 1 0.50 0.67 0.57 147

accuracy 0.52 308 macro avg 0.53 0.53 0.52 308 weighted avg 0.53 0.52 0.51 308

## **#USING KERNAL="RBF"**

print("Classification Report")

print(classification\_report(y\_test,y\_pred))

```
from sklearn.svm import SVC

clf=SVC(kernel="rbf")

model=clf.fit(x_train,y_train)

y_pred= model.predict(x_test)

import seaborn as sns

import matplotlib.pyplot as plt

sns.regplot(y_pred,y_test)

from sklearn.metrics import classification_report,confusion_matrix,accuracy_score

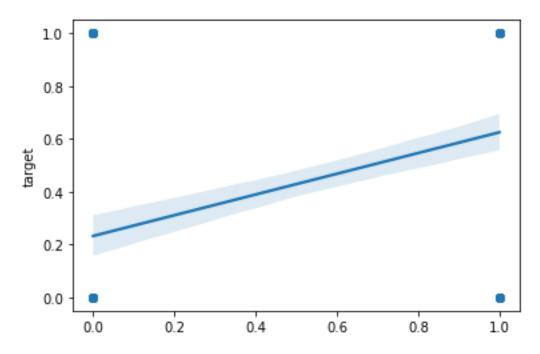
result=confusion_matrix(y_test,y_pred)

print("Confusion Matrix:")

print(result)

print("Accuracy=",accuracy_score(y_test,y_pred))
```

## **OUTPUT:**



Confusion Matrix: [[ 89 72]

[ 27 120]]

Accuracy= 0.6785714285714286

#### Classification Report

precision recall f1-score support 0.77 0 0.55 0.64 161 0.62 1 0.820.71 147 0.68 308 accuracy 0.70 0.68 308 macro avg 0.68weighted avg 0.70 0.68 0.67 308

## 15: WRITE A PROGRAM TO IMPLEMENT PRINCIPAL COMPONENT **ANALYSIS**

**15. AIM:** To implement principal component analysis

# **Program:**

```
#BEFORE PCA
from sklearn import tree
from sklearn.model_selection import train_test_split
import pandas as pd
df=pd.read_csv(r"C:\Users\Downloads\heart (1).csv")
print(df.head())
y=df['target']
x=df.drop("target",axis=1)
x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=0.3,random_state=1)
model = tree.DecisionTreeClassifier()
model = model.fit(x_train, y_train)
y_pred = model.predict(x_test)W
from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
result=confusion_matrix(y_test,y_pred)
print("Confusion Matrix:")
print(result)
print("Accuracy=",accuracy_score(y_test,y_pred))
print("Classification Report")
print(classification_report(y_test,y_pred))
```

#### **#AFTER PCA**

```
b=df['target']
a=df.drop("target",axis=1)
a_train, a_test, b_train, b_test = train_test_split(a, b, test_size=0.3,random_state=1)
from sklearn.decomposition import PCA
#USING 2 COMPONENTS
pca=PCA(n_components=2)
a_train=pca.fit_transform(a_train)
a_test=pca.transform(a_test)
model = tree.DecisionTreeClassifier()
model = model.fit(a_train, b_train)
y_pred = model.predict(a_test)
from sklearn.metrics import classification_report,confusion_matrix,accuracy_score
result=confusion_matrix(b_test,y_pred)
print("Confusion Matrix:")
print(result)
print("Accuracy=",accuracy_score(b_test,y_pred))
print("Classification Report")
print(classification_report(b_test,y_pred))
#USING 6 COMPONENTS
pca=PCA(n_components=6)
a_train=pca.fit_transform(a_train)
a_test=pca.transform(a_test)
model = tree.DecisionTreeClassifier()
model = model.fit(a_train, b_train)
y_pred = model.predict(a_test)
from sklearn.metrics import classification report, confusion matrix, accuracy score
result=confusion_matrix(b_test,y_pred)
print("Confusion Matrix:")
print(result)
print("Accuracy=",accuracy_score(b_test,y_pred))
print("Classification Report")
```

## **OUTPUT:**

#### **#BEFORE PCA**

	age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	са	thal	target
0	52	1	0	125	212	0	1	168	0	1.0	2	2	3	0
1	53	1	0	140	203	1	0	155	1	3.1	0	0	3	0
2	70	1	0	145	174	0	1	125	1	2.6	0	0	3	0
3	61	1	0	148	203	0	1	161	0	0.0	2	1	3	0
4	62	0	0	138	294	1	1	106	0	1.9	1	3	2	0

Confusion Matrix:

[[161 0] [ 3 144]]

Accuracy= 0.9902597402597403 Classification Report

precision recall f1-score support

0 0.98 1.00 0.99 161 1 1.00 0.98 0.99

0.99 

## **#AFTER PCA**

## **#USING 2 COMPONENTS**

Confusion Matrix:

[[161 0]

[ 6 141]]

Accuracy= 0.9805194805194806

Classification Report

precision recall f1-score support

0 0.96 1.00 0.98 161 1 1.00 0.96 0.98 147

0.98 308 accuracy macro avg 0.98 0.98 0.98 308 weighted avg 0.98 0.98 0.98 308

## **#USING 6 COMPONENTS**

Confusion Matrix:

[[157 4]

[ 3 144]]

Accuracy= 0.9772727272727273

Classification Report

precision recall f1-score support

0.98 0.98 0.98 161 0.97 0.98 0.98

0.98 accuracy macro avg 0.98 0.98 0.98 308 weighted avg 0.98 0.98 0.98