INDEX

Sr. No.	Practical	Page no.	Sign
1	 A. Design a simple machine learning model to train the training instances and test the same using Python. B. 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. 	4	
2	A. Perform Data Loading, Feature selection (Principal Component analysis) and Feature Scoring and Ranking. B. For a given set of training data examples stored in .CSV file, implement and demonstrate the CandidateElimination algorithm to output a description of the set of all hypotheses consistent with the training examples.	9	
3	Write a program to implement Decision Tree and Random Forest with Prediction, Test Score and Confusion Matrix.	15	
4	 A. For a given set of training data examples stored in a .CSV file implement Least Square Regression algorithm. (Use Univariate dataset) B. For a given set of training data examples stored in a .CSV file implement Logistic Regression algorithm. (Use Multivariate dataset) 	18	
5	Write a program to implement k-Nearest Neighbor algorithm to classify the iris data set.	26	
6	 A. Implement the different Distance methods (Euclidean, Manhattan Distance, Minkowski Distance) with Prediction, Test Score and Confusion Matrix. B. Implement the classification model using clustering for the following techniques with K means clustering with Prediction, Test Score and Confusion Matrix. 	29	
7	A. Implement the classification model using clustering for the following techniques with hierarchical clustering with Prediction, Test Score and Confusion Matrix	36	

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Rohit Kumar Prajapati (09)

8	A. Write a 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.	
	B. Implement the non-parametric Locally Weighted Regression algorithm in order to fit data points. Select appropriate data set for your experiment and draw graphs.	

Practical No - 1A

Design a simple machine learning model to train the training instances and test the same using Python.

Code:

```
# Practical 1 - Linear Regression Model
print("Practical no.1A : Linear Regression \nName: Rohit \t Roll no.:09")
print("----")
import random
from sklearn.linear model import LinearRegression
print("Read the train Data")
print("----")
feature set = []
target set = []
no of rows = 200
limit = 2000
for i in range(0, no of rows):
   x = random.randint(0, limit)
   y = random.randint(0, limit)
   z = random.randint(0, limit)
   g = 10 * x + 2 * y + 3 * z
   print("x=", x, "\ty=", y, "\tz=", z, "\tg=", g)
   feature set.append([x, y, z])
   target set.append(g)
print("Here the training of model begins. ")
model = LinearRegression()
model.fit(feature set, target set)
print("Training of model ends here!")
print("Testing started here")
test data = [[1, 1, 0]]
print("Test data:", test data)
prediction = model.predict(test data)
print("prediction:" + str(prediction) + '\t' + "Coefficient:" +
str(model.coef ))
```

Practical No - 1B

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.

Training data → dataset 1b.csv

Sky	Temp	Humidity	Wind	Water	Forecast	EnjoySport
Sunny	Warm	Normal	Strong	Warm	Same	Yes
Sunny	Warm	High	Strong	Warm	Same	Yes
Rainy	Cold	High	Strong	Warm	Change	No
Sunny	Warm	High	Strong	Cool	Change	Yes

```
import csv
  num_attributes = 6 dataset = []
print("Name: Rohit \tRoll No: 09")
print("Training data")

with open(r"data_find_s_lb.csv", "r") as csvfile:
    reader = csv.reader(csvfile)
for row in reader:
    dataset.append(row)

# print(row) # print(dataset) print("\n The
initial value of hypothesis: ") hypothesis =
['0'] * num_attributes print(hypothesis)

for j in range(0, num_attributes): hypothesis[j] =
dataset[1][j] print("\n Find S: Finding a Maximally Specific
Hypothesis \n:") for i in range(1, len(dataset)): if
```

```
Training data
Find S: Finding a Maximally Specific Hypothesis
The Maximally Specific Hypothesis for a given training examples:
['Sunny', '?', '?', '?', '?', '?']
Find S: Finding a Maximally Specific Hypothesis
  PROBLEMS (1) OUTPUT DEBUG CONSOLE TERMINAL JUPYTER
  For Training instance no:2 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', '?']
For Training instance no:3 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', '?']
For Training instance no:4 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', '?']
    The Maximally Specific Hypothesis for a given training examples:
   ['Sunny', 'Warm', '?', 'Strong', '?', '?']
    Find S: Finding a Maximally Specific Hypothesis
  For Training instance no:1 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', 'Warm', '?']
For Training instance no:2 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', 'Warm', '?']
For Training instance no:3 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', 'Warm', '?']
For Training instance no:4 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', '?']
    The Maximally Specific Hypothesis for a given training examples:
   ['Sunny', 'Warm', '?', 'Strong', '?', '?']
    Find S: Finding a Maximally Specific Hypothesis
  For Training instance no:1 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', 'Same']
For Training instance no:2 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', 'Same']
For Training instance no:3 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', 'Same']
For Training instance no:4 the hypothesis is ['Sunny', 'Warm', '?', 'Strong', '?', '?']
    The Maximally Specific Hypothesis for a given training examples:
   ['Sunny', 'Warm', '?', 'Strong', '?', '?']
   PS C:\Users\Shalu\Documents\mscit\practical\Machine Learning> data_find_s_1b.csv
```

Practical No - 2A

Perform Data Loading, Feature selection (Principal Component analysis) and Feature Scoring and Ranking.

```
import pandas as pd from sklearn.model selection
import train test split from sklearn.decomposition
import PCA from sklearn.preprocessing import
StandardScaler
print("Name: Rohit \tRoll No: 09")
# Reading dataset using pandas url =
'https://archive.ics.uci.edu/ml/machine-
learningdatabases/iris/iris.data' names = ["sepal-length", "sepal-
width", "petal-length", "petal-width",
"class"] dataset = pd.read csv(url,
names=names)
# Display dataset
print(dataset.head(10))
X = dataset.drop("class", 1)
Y = dataset["class"]
# Splitting the train and test datasets
x train, x test, y train, y test = train test split(X, Y,
test size=0.2, random state=0)
sc = StandardScaler() x train =
sc.fit transform(x train) x test =
sc.transform(x test)
# Display Training and testing data
print(f"\nDataSet before PCA :\n\nTrain
:\n{x train}\n\nTest
: \n{x test}")
```

```
# print(x_train)
# print(x_test)

# Creating PCA

pca = PCA() x_train = pca.fit_transform(x_train)
x_test = pca.transform(x_test)

# Giving a principal feature to model pca =
PCA(n_components=2) x_train =
pca.fit_transform(x_train) x_test =
pca.transform(x_test)

print(f"\nDataSet After PCA :\n\nTrain :\n{x_train}\n\nTest
:\n{x_test}")
```

```
sepal-length sepal-width petal-length petal-width class
0 5.1 3.5 1.4 0.2 Iris-setosa
1 4.9 3.0 1.4 0.2 Iris-setosa
2 4.7 3.2 1.3 0.2 Iris-setosa
3 4.6 3.1 1.5 0.2 Iris-setosa
4 5.0 3.6 1.4 0.2 Iris-setosa
4 5.0 3.6 1.4 0.2 Iris-setosa
6 4.6 3.4 1.4 0.3 Iris-setosa
6 4.6 3.4 1.4 0.3 Iris-setosa
7 5.0 3.4 1.5 0.2 Iris-setosa
8 4.4 2.9 1.4 0.2 Iris-setosa
9 4.9 3.1 1.5 0.1 Iris-setosa
9 1.7 0.0 1.4 0.2 Iris-setosa
9 1.7 0.0 1.4 0.2 Iris-setosa
9 1.7 0.0 1.4 0.2 Iris-setosa
9 1.8 0.1 Iris-setosa
9 1.9 0.1 Iris-setosa
9 1.9 0.1 Iris-setosa
1.0 0.1 Iris-setosa
9 0.1 0.1 Iris-setosa
1 0.2 Iris-setosa
1 0.2 Iris-setosa
1 0.3 Iris-set
```

Practical No - 2B

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

Data set :data_2b.csv

sky	airtemp	humidity	wind	water	forcast	enjoysport
sunny	warm	normal	strong	warm	same	yes
sunny	warm	high	strong	warm	same	yes
rainy	cold	high	strong	warm	change	no
sunny	warm	high	strong	cool	change	yes

```
import numpy as np import pandas as pd
print("Name: Rohit \tRoll No: 09")
data = pd.read csv(r'C:\Users\Shalu\Documents\mscit\practical\Machine
Learning\data find s 1b.csv') concepts = np.array(data.iloc[:, 0:-1])
print("\nInstances are:\n", concepts) target = np.array(data.iloc[:, -
1]) print("\nTarget Values are: ", target)
def learn(concepts, target):
   specific h and genearal h")
print("\nSpecific Boundary: ",
            general h = [["?" for i in range(len(specific h))] for i
specific h)
in range(len(specific h))]
print("\nGeneric Boundary: ", general h)
    for i, h in enumerate(concepts):
     "Yes":
         print("Instance is Positive ")
                                                    for x in
                          if h[x] != specific h[x]:
range(len(specific h)):
               specific_h[x] =
general h[x][x] = '?'
       if target[i] == "No":
```

```
print("Instance is Negative ")
                   range(len(specific h)):
for
              in
if h[x] != specific h[x]:
                   general h[x][x] = specific h[x]
else:
                   general h[x][x] = '?'
        print("Specific Bundary after ", i + 1, "Instance is ",
                     print("Generic Boundary after ", i + 1,
specific h)
"Instance is ", general h)
                                 print("\n")
     indices = [i for i, val in enumerate(general h) if val == ['?',
1?1,
·?·, ·?·, ·?·, ·?·]]
for i in indices:
       general h.remove(['?', '?', '?', '?', '?'])
return specific h, general h
s final, g final = learn(concepts,
target)
print("Final Specific h: ", s final, sep="\n")
print("Final General h: ", g final, sep="\n")
```

Practical No – 3

Write a program to implement Decision Tree and Random forest with Prediction, Test Score and Confusion Matrix.

```
Decision tree import pandas as pd from sklearn.model selection
import train test split from sklearn.tree import
DecisionTreeClassifier
from sklearn.metrics import confusion matrix, accuracy score,
classification report # Read Datasets def read datasets():
   print("----")
   datasets =
pd.read csv("https://archive.ics.uci.edu/ml/machinelearning-
databases/balance-scale/balance-scale.data",
sep=",", header=None) print(f"Dataset Length : {len(datasets)}")
{datasets.head()}") return datasets
# Spliting Datasets into train and test def
splitdataset(datasets):
X = datasets.values[:, 1:5]
Y = datasets.values[:, 0]
   X train, X test, y train, y test = train test split(X, Y,
test size=0.3, random state=100)
   # print(f"x train:{X train}\n x test: {X test}\n y train:
{y train}\n y test: {y test}") return X train, X test,
y_train, y_test def train_using_gini(X_train, y_train):
   clf gini = DecisionTreeClassifier(criterion='gini',
random state=100, max depth=3, min samples leaf=5)
def train using entropy(X train, y train):
```

```
clf entropy = DecisionTreeClassifier(criterion='entropy',
random state=100, max depth=3, min samples leaf=5)
prediction(X test, clf object):
   y pred = clf object.predict(X test)
print(f"Predicted values: {y pred}")
return y pred def cal accuracy(y test,
y pred):
   print(f"Confusion Metrix :{confusion matrix(y test, y pred)}")
print(f"Accuracy: {accuracy score(y test, y pred) * 100}")
print(f"Report: {classification report(y test, y pred)}") def
main():
  datasets = read datasets()
   X train, X test, y train, y test = splitdataset(datasets)
= train_using_entropy(X_train, y_train) print("Results
using Gini Index: ") y pred gini = prediction(X test,
clf_gini) cal_accuracy(y_test, y_pred_gini)
prediction(X test, clf entropy) cal accuracy(y test,
y pred entropy if name == " main ":
   main()
```

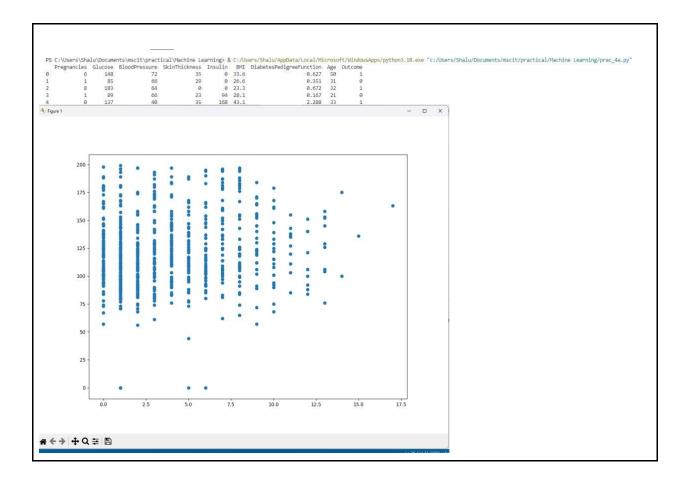
```
Direct Longit : 655
Direct Support (655, 5)
Direct Sup
```

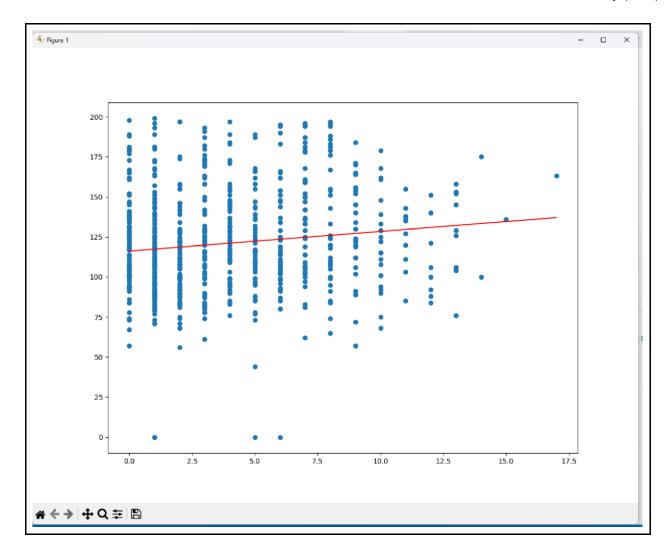
Practical No - 4A.

For a given set of training data examples stored in a .CSV file implement Least Square Regression algorithm. (Use Univariate dataset)

```
11 11 11
Practical No - 4A.
For a given set of training data examples stored in a .CSV file
implement Least Square Regression algorithm. (Use Univariate
dataset )
""" import pandas as pd import
numpy as np import
matplotlib.pyplot as plt
# from sklearn.model selection import train test split
# from sklearn.preprocessing import StandardScaler
# from sklearn.linear model import Lea
# from sklearn.metrics import confusion matrix, accuracy score
plt.rcParams['figure.figsize'] = (12.0, 9.0)
dataset =
pd.read csv("https://raw.githubusercontent.com/plotly/datasets/master/di
abetes.csv") print(dataset.head()) print("Name: Rohit, Roll no: 09") x =
dataset.iloc[:,0] y = dataset.iloc[:,1] plt.scatter(x, y) plt.show()
x mean = np.mean(x)
y mean = np.mean(y) num
= 0 den = 0 for i in
range (len(x)):
    num += (x[i] -x_mean) * (y[i] -y_mean)
   den += (x[i] -x mean) **2
m = num/den c =
y mean - m*x mean
print(m,c) y pred =
m*x +c
```

```
plt.scatter(x,y) plt.plot([min(x), max(x)], [min(y_pred),
max(y_pred)], color="red") plt.show()
```





Practical No - 4B

For a given set of training data examples stored in a .CSV file implement Logistic Regression algorithm. (Use Multivariate dataset)

```
import pandas as pd from sklearn.model selection import
train test split from sklearn.preprocessing import
StandardScaler from sklearn.linear model import
LogisticRegression from sklearn.metrics import
confusion matrix, accuracy score
 print("Name: Rohit \tRoll No: 09")
dataset =
pd.read csv("https://raw.githubusercontent.com/plotly/da
tasets/master/di abetes.csv") print(dataset.head()) x =
dataset.iloc[:, [0, 1, 2, 3, 4, 5, 6, 7]].values y =
dataset.iloc[:, [-1]].values print(x) print(y)
x train, x test, y train, y test = train test split(x,
y, test size=0.2, random state=0) sc = StandardScaler()
x_train = sc.fit_transform(x_train) x_test =
sc.transform(x test)
print(x train[0:15, :])
classifier = LogisticRegression()
classifier.fit(x train, y train)
y pred = classifier.predict(x test)
cm = confusion matrix(y test, y pred)
print("Confusion Matrix :\n ", cm)
print("Accuracy :", accuracy_score(y_test, y_pred))
```

Practical No - 5

Write a program to implement k-Nearest Neighbor algorithm to classify the iris data set.

```
""" Write a program to implement k-Nearest
Neighbour algorithm to classify the iris data set.
 """ from sklearn.neighbors import KNeighborsClassifier from
sklearn.preprocessing import StandardScaler from sklearn.pipeline
import make pipeline from sklearn import datasets from
sklearn.model selection import train test split, GridSearchCV
iris=datasets.load iris()
X=iris.data
Y=iris.target
## Create train and test split X train, X test, Y train, Y test =
train test split(X,Y,test size=0.3,random state=42,stratify=Y)
#feature Scaling using StandardScalar
sc= StandardScaler() sc.fit(X train)
X train std= sc.transform(X train)
X test std= sc.transform(X test)
#Fit the model
knn=KNeighborsClassifier(n neighbors=5,p=2,weights='uniform',algorithm='
auto') knn.fit(X train std, Y train)
#Evaluate the training and test score print("Name: Rohit \t Roll no:
09") print("Traing Accuracy score: %.3f "
%knn.score(X train std,Y train)) print("Test Accuracy score: %.3f"
%knn.score(X test std,Y test))
iris=datasets.load iris()
X=iris.data
Y=iris.target
```

```
## Create train and test split X train, X test, Y train, Y test =
train test split(X,Y,test size=0.3,random state=42,stratify=Y)
#create a pipeline
pipeline=make pipeline(StandardScaler(), KNeighborsClassifier())
#Create a parameter
param grid=[{
    'kneighborsclassifier__n_neighbors':[2,3,4,5,6,7,8,9,10],
    'kneighborsclassifier p':[1,2],
    'kneighborsclassifier weights':['uniform','distance'],
'kneighborsclassifier algorithm':['auto','ball tree','kd tree','brute'
} ]
#Create a grid search instance
gs=GridSearchCV(pipeline, param grid=param grid,
scoring="accuracy",
                                    refit=True,
cv=10,
                      verbose=1,
n jobs=2)
gs.fit(X train, Y train)
#print best model parameter and score
print("Best Score: %.3f " %gs.best score ," \nBest
Parameters:", gs.best params )
```

```
Traing Accuracy score: 0.901
Test Accuracy score: 0.901
Fitting 10 folds for each of 144 candidates, totalling 1440 fits
Best Score: 0.972
Best Parameters: {'kneighborsclassifier_algorithm': 'auto', 'kneighborsclassifier_n_neighbors': 5, 'kneighborsclassifier_p': 1, 'kneighborsclassifier_weights': 'uniform'}
PS C:\Users\Shalu\Documents\mscit\practical\Wachine Learning> []
```

Practical No - 6A.

Implement the different Distance methods (Euclidean, Manhattan Distance, Minkowski Distance) with Prediction, Test Score and Confusion Matrix.

```
# Q. 6A. Implement the different Distance methods (Euclidean, Manhattan
Distance, Minkowski Distance)
# with Prediction, Test Score and Confusion Matrix. from math import
sqrt from sklearn.metrics import confusion matrix,
classification report
print("Practical no.3 : Decision tree \nName: Rohit \t Roll no.:09")
print("----")
euclidian distance(a, b):
   return sqrt(sum((e1 - e2) ** 2 for e1, e2 in zip(a, b))) def
manhattan distance(a, b):
   return sum(abs(e1 - e2) for e1, e2 in zip(a, b))
def minkowski_distance(a, b, p):
   return sum(abs(e1 - e2) ** p for e1, e2 in zip(a, b)) ** (1 / p)
actual = [1, 0, 0, 1, 0, 0, 1, 0, 0, 1] predicted = [1, 0, 0, 1, 0,
0, 0, 1, 0, 0] dist1 = euclidian distance(actual, predicted) dist2 =
manhattan distance(actual, predicted) dist3 =
minkowski distance (actual, predicted, 1)
```

```
print("Confusion_matrix: \n", matrix)

tp, fn, fp, tn = confusion_matrix(actual, predicted, labels=[1,
0]).reshape(-1)

print("Outcome values: \n", tp, fn, fp, tn)

matrix = classification_report(actual, predicted, labels=[1, 0])

print("Classification_report: \n", matrix)
```

```
print(f"Euclidian_dist: {dist1}\nManhattan_dist: {dist2}\nMinkowski_dist
with value 1: {dist3}") dist4 = minkowski_distance(actual, predicted, 2)
print(f"Minkowski_dist with value 2: {dist4}\n")
matrix = confusion_matrix(actual, predicted, labels=[1, 0])
```

```
Euclidian_dist: 1.7320508075688772
Manhattan dist: 3
Minkowski_dist with value 1: 3.0
Minkowski_dist with value 2: 1.7320508075688772
Confusion_matrix:
 [[2 2]
 [1 5]]
Outcome values:
 2 2 1 5
Classification report:
           precision recall f1-score support
        1
            0.67 0.50 0.57
                                       4
            0.71 0.83 0.77
                                       6
                               0.70
                                       18
   accuracy
  macro avg 0.69 0.67
                               0.67
weighted avg 0.70 0.70
                             0.69
PS C:\Users\Shalu\Documents\mscit\practical\Machine Learning>
```

Practical No – 6B

Implement the classification model using clustering for the following techniques with K means clustering with Prediction, Test Score and Confusion Matrix.

```
# Common imports
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.preprocessing import StandardScaler
from sklearn.model selection import train test split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification report
from sklearn.metrics import confusion matrix
# Import the data set raw data
= pd.read csv(
    'https://raw.githubusercontent.com/imhardikj/A-Model-a-
Day/master/KNearest%20Neighbors/Classified%20Data.csv', index col=0)
# Standardize the data set
scaler = StandardScaler()
scaler.fit(raw data.drop('TARGET CLASS', axis=1))
scaled features = scaler.transform(raw data.drop('TARGET CLASS',
axis=1))
scaled data = pd.DataFrame(scaled features,
columns=raw data.drop('TARGET CLASS', axis=1).columns)
# Split the data set into training data and test data
```

```
x = scaled data
y = raw data['TARGET CLASS']
x training data, x test data, y training data, y test data =
train test split(x, y, test size=0.3)
# Train the model and make predictions model =
KNeighborsClassifier(n neighbors=1)
model.fit(x_training_data, y_training_data) predictions
= model.predict(x test data)
# Performance measurement
print(classification report(y test data, predictions))
print(confusion matrix(y test data, predictions))
# Selecting an optimal K value error rates = []
# for i in np.arange(1, 101):
new model = KNeighborsClassifier(n neighbors=1)
new_model.fit(x_training_data, y_training_data) new_predictions =
new_model.predict(x_test_data)
error rates.append(np.mean(new predictions != y test data))
plt.figure(figsize=(16, 12)) plt.plot(error rates) plt.show()
```

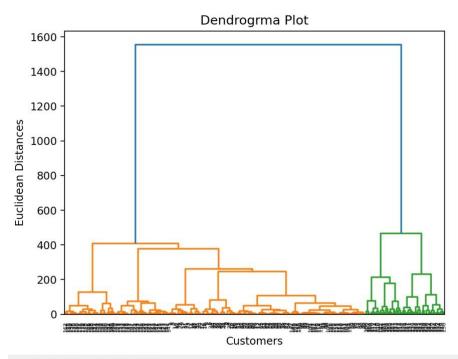
	precision	recall	f1-score	support	
0	0.92	0.91	0.92	143	
1	0.92	0.93	0.92	157	
accuracy			0.92	300	
macro avg	0.92	0.92	0.92	300	
weighted avg	0.92	0.92	0.92	300	
[[130 13] [11 146]]					

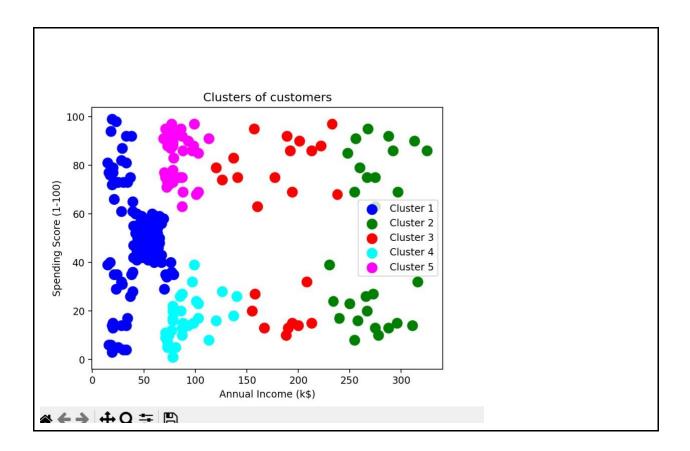
Practical No – 7A

A. Implement the classification model using clustering for the following techniques with hierarchical clustering with Prediction, Test Score and Confusion Matrix

```
Implement the classification model using clustering for the following
techniques with hierarchical clustering with Prediction,
Test Score and Confusion Matrix
1 1 1
# Importing the libraries import
numpy as nm
import matplotlib.pyplot as mtp import
pandas as pd
# Importing the dataset
dataset = pd.read csv('7 Mall Customers.csv') x
= dataset.iloc[:, [3, 4]].values
# Finding the optimal number of clusters using the dendrogram import
scipy.cluster.hierarchy as shc
dendro = shc.dendrogram(shc.linkage(x, method="ward"))
mtp.title("Dendrogrma Plot") mtp.ylabel("Euclidean
Distances") mtp.xlabel("Customers") mtp.show()
# training the hierarchical model on dataset from
sklearn.cluster import AgglomerativeClustering
hc = AgglomerativeClustering(n clusters=5, affinity='euclidean',
linkage='ward')
y pred = hc.fit predict(x)
# visulaizing the clusters
mtp.scatter(x[y\_pred == 0, 0], x[y pred == 0, 1], s=100, c='blue',
label='Cluster 1')
mtp.scatter(x[y\_pred == 1, 0], x[y pred == 1, 1], s=100, c='green',
label='Cluster \overline{2}')
mtp.scatter(x[y\_pred == 2, 0], x[y\_pred == 2, 1], s=100, c='red',
label='Cluster 3')
mtp.scatter(x[y\_pred == 3, 0], x[y pred == 3, 1], s=100, c='cyan',
label='Cluster \overline{4}')
mtp.scatter(x[y\_pred == 4, 0], x[y pred == 4, 1], s=100, c='magenta',
label='Cluster 5')
```

```
mtp.title('Clusters of customers')
mtp.xlabel('Annual Income (k$)')
mtp.ylabel('Spending Score (1-100)')
mtp.legend()
mtp.show()
```





Practical No - 8A

Write a 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.

```
A. Write a program to construct a Bayesian network considering medical
Use this model to demonstrate the diagnosis of heart patients using
standard Heart Disease Data Set.
import numpy as np
import pandas as pd
import csv
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianNetwork from
pgmpy.inference import VariableElimination
heartDisease = pd.read csv('8a heart data.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=
BayesianNetwork([('age','heartdisease'),('gender','heartdisease'),('exan
g', 'heartdisease'), ('cp', 'heartdisease'), ('heartdisease', 'restecg'), ('he
artdisease','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={'re
stecg':1}) print(q1)
print('\n 2. Probability of HeartDisease given evidence= cp ')
q2=HeartDiseasetest infer.query(variables=['heartdisease'],evidence={'cp
':2})
```

```
print(q2)
```

```
Sample instances from the dataset are given below
  age gender cp trestbps chol ... oldpeak slope ca thal heartdisease
        1 1
               145 233 ... 2.3 3 0 6
160 286 ... 1.5 2 3 3
                 160 286 ...
        1 4
                                              3
                                                        2

    2
    67
    1
    4
    100
    286
    ...
    1.5
    2
    3
    3

    2
    67
    1
    4
    120
    229
    ...
    2.6
    2
    2
    7

    3
    37
    1
    3
    130
    250
    ...
    3.5
    3
    0
    3

    4
    41
    0
    2
    130
    204
    ...
    1.4
    1
    0
    3

[5 rows x 14 columns]
Attributes and datatypes
             int64
age
gender
             int64
             int64
trestbps
             int64
chol
             int64
fbs
             int64
restecq
             int64
thalach
            int64
exang
            int64
oldpeak
           float64
slope
             int64
            object
thal
            object
heartdisease
dtype: object
 Inferencing with Bayesian Network:
 1. Probability of HeartDisease given evidence= restecg
+------
| heartdisease | phi(heartdisease) |
| heartdisease(0) |
+-----+
| heartdisease(1) |
+-----
| heartdisease(2) |
+-----+
| heartdisease(3) |
                               0.2015 |
+----+
| heartdisease(4) |
+----+
 2. Probability of HeartDisease given evidence= cp
+----+
 | heartdisease | phi(heartdisease) |
+=========+
| heartdisease(0) |
                               0.3610
+------
| heartdisease(1) |
| heartdisease(2) |
                               0.1373 |
| heartdisease(3) |
                                0.1537 I
+------
| heartdisease(4) |
                                0.1321 |
```

Practical No – 8B

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

```
""" NON Parametric model
 Implement the non-parametric Locally Weighted Regression algorithm in order
to fit data points.
 Select appropriate data set for your experiment and draw graphs.
import matplotlib.pyplot as plt
import pandas as pd import numpy
as np
def kernel(point, xmat, k):
m, n = np.shape(xmat) weights
= np.mat(np.eye((m))) for j in
= np.mat(np.eye((m)))
                         for j in
range (m):
        diff = point - X[j]
        weights[j, j] = np.exp(diff * diff.T / (-2.0 * k ** 2))
return weights
 def localWeight(point, xmat, ymat, k):
    wei = kernel(point, xmat, k)
    W = (X.T * (wei * X)).I * (X.T * (wei * ymat.T))
return W
 def localWeightRegression(xmat, ymat, k):
    m, n = np.shape(xmat)
ypred = np.zeros(m)
i in range(m):
        ypred[i] = xmat[i] * localWeight(xmat[i], xmat, ymat, k)
return ypred
# load data points
data = pd.read csv('8b resturant data.csv')
bill = np.array(data.total_bill) tip =
np.array(data.tip)
# preparing and add 1 in bill
mbill = np.mat(bill) mtip =
np.mat(tip)
m = np.shape(mbill)[1] one =
np.mat(np.ones(m)) X =
np.hstack((one.T, mbill.T))
# set k here
ypred = localWeightRegression(X, mtip, 0.5)
SortIndex = X[:, 1].argsort(0) xsort
= X[SortIndex][:, 0]
```

```
fig = plt.figure()
ax = fig.add_subplot(1, 1, 1) ax.scatter(bill,
tip, color='green')
ax.plot(xsort[:, 1], ypred[SortIndex], color='red', linewidth=5) plt.xlabel('Total bill') plt.ylabel('Tip')
plt.show()
```

