



CAMBRIDGE INSTITUTE OF TECHNOLOGY
KR PURAM,BANGLORE -560036 – KARNATAKA

AI AND ML APPLICATION DEVELOPMENT LABORATORY
(Effective from the academic year 2018 -2019)

SEMESTER – VII

DEPARTMENT OF ARTIFICIAL INTELLIGENCE AND MACHINE
LEARNING

LAB MANUAL

SUBJECT- AI AND ML APPLICATION DEVELOPMENT LABORATORY

CODE- 18AIL76

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Program 1

Write a program to implement k-Nearest Neighbour algorithm to classify the iris data set. Print both correct and wrong predictions.

Dataset - [link](#)

```
import pandas as pd
import numpy as np
from sklearn import datasets
from sklearn.neighbors import KNeighborsClassifier
from sklearn.model_selection import train_test_split

# Importing Dataset
dataset = datasets.load_iris()
X = pd.DataFrame(dataset.data)
y = pd.DataFrame(dataset.target)

# Splitting The Dataset Into Training Set And Testing Set
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2)

# Model Fit And Prediction
knn = KNeighborsClassifier(n_neighbors=5)
knn.fit(X_train, y_train.values.ravel())

accuracy_train = knn.score(X_train, y_train)
accuracy_test = knn.score(X_test, y_test)

print("Training Accuracy\n", accuracy_train)
print("Testing Accuracy\n", accuracy_test)

available_class = pd.DataFrame(dataset.target_names)
print("Dataset Classes\n", available_class)

example = np.array([5.7, 3, 4.2, 1.2])
example = example.reshape(1, -1)
print("Input Sample\n", example)

example_prediction = int(knn.predict(example))
print("Predicted Class\n", available_class[0][example_prediction])
```

OUTPUT -

```
Training Accuracy
0.9583333333333334
Testing Accuracy
1.0
Dataset Classes
0
0    setosa
1    versicolor
2    virginica
Input Sample
[[5.7 3. 4.2 1.2]]
Predicted Class
Versicolor
```

Program - 2

Develop a program to apply K-means algorithm to cluster a set of data stored in .CSV file. Use the same data set for clustering using EM algorithm. Compare the results of these two algorithms and comment on the quality of clustering.

Dataset- same as 1st

```
import sklearn.metrics as sm import pandas as
pd import numpy as np import
matplotlib.pyplot as plt from sklearn import
datasets from sklearn import preprocessing
from sklearn.cluster import KMeans from
sklearn.mixture import GaussianMixture

# Importing Dataset dataset
= datasets.load_iris()

# Attribute Variable(s)
X = pd.DataFrame(dataset.data)
X.columns = ['Sepal_Length', 'Sepal_Width', 'Petal_Length', 'Petal_Width'] #
Target Variable
y = pd.DataFrame(dataset.target) y.columns
= ['Targets']
```

```

plt.figure(figsize=(14, 7))
colormap = np.array(['darkorange', 'navy', 'darkgreen'])

# REAL PLOT
plt.subplot(1, 3, 1)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y.Targets], s=40)
plt.title('Real Classification') plt.xlabel('Petal Length') plt.ylabel('Petal Width')

# K-PLOT
plt.subplot(1, 3, 2) model =
KMeans(n_clusters=3)
model.fit(X)
predY = np.choose(model.labels_, [0, 1, 2]).astype(np.int64)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[predY], s=40)
plt.title('K-Means') plt.xlabel('Petal Length') plt.ylabel('Petal Width')
print('K-Means Algorithm')
print('The Accuracy Score Of K-Mean Algorithm: ', sm.accuracy_score(y, model.labels_))
print('The Confusion Matrix Of K-Mean Algorithm:\n', sm.confusion_matrix(y,
model.labels_))

# EM PLOT
scaler = preprocessing.StandardScaler()
scaler.fit(X) xsa = scaler.transform(X) xs =
pd.DataFrame(xsa, columns=X.columns)
gmm = GaussianMixture(n_components=3)
gmm.fit(xs) y_gmm =
gmm.predict(xs)
plt.subplot(1, 3, 3)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y_gmm], s=40)
plt.title('GMM Classification') plt.xlabel('Petal Length') plt.ylabel('Petal Width')
print('EM Algorithm') print('The Accuracy Score Of EM Algorithm: ',
sm.accuracy_score(y, y_gmm)) print('The Confusion Matrix Of EM Algorithm:\n',
sm.confusion_matrix(y, y_gmm)) plt.show()

```

Output –

K-Means Algorithm

The Accuracy Score Of K-Mean Algorithm: 0.24 The
Confusion Matrix Of K-Mean Algorithm:

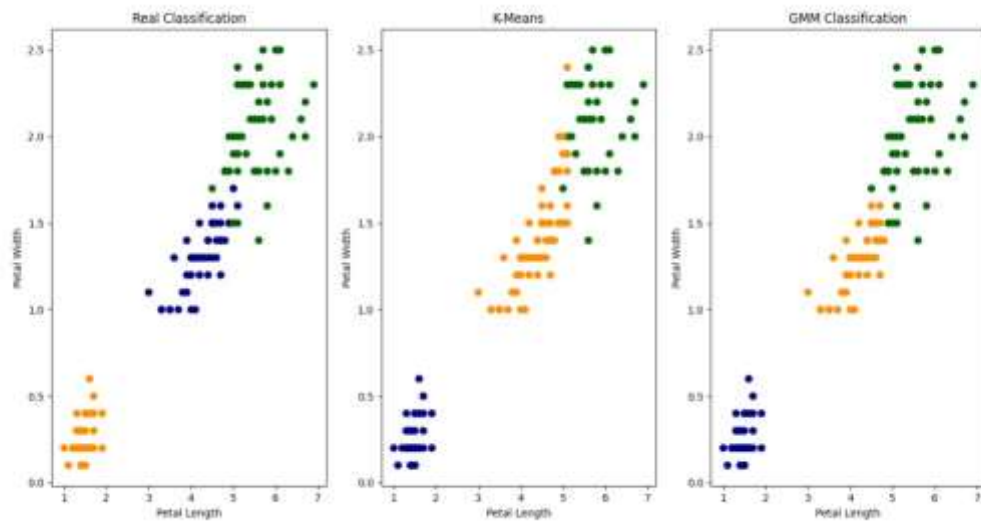
```
[[ 0 50  0]
 [48  0  2]
 [14  0 36]]
```

EM Algorithm

The Accuracy Score Of EM Algorithm: 0.3333333333333333 The
Confusion Matrix Of EM Algorithm:

```
[[ 0 50  0]
 [45  0  5]]
```

[0 0 50]]



Program – 3

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

Dataset - [link](#)

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
```

```
data_frame = pd.read_csv('data/tips.csv')
```

```
feature_array = np.array(data_frame.total_bill)
label_array = np.array(data_frame.tip)
```

```
total_cases = feature_array.shape[0]
```

```
y_matrix = np.matrix(label_array).T
```

```
x_matrix = np.hstack((np.ones((total_cases, 1)), np.matrix(feature_array).T))
```

```
y_predicted = np.zeros(total_cases)
```

```
# get weight matrix def
```

```
get_weight_matrix(current_x, x_matrix, tau):
```

```
weight_matrix = np.mat(np.eye(total_cases))
```

```
for j in range(total_cases):
```

```
    distance_vector = current_x - x_matrix[j]
```

```
    weight_matrix[j, j] = np.exp(-1 * distance_vector*distance_vector.T / (2.0 * tau ** 2))
```

```
return weight_matrix
```

```

# get theta matrix def get_theta_matrix(current_x,
x_matrix, y_matrix, tau):
    weight_matrix = get_weight_matrix(current_x, x_matrix, tau)
    theta_matrix = np.linalg.inv(x_matrix.T * (weight_matrix * x_matrix)) * x_matrix.T *
weight_matrix * y_matrix
    return theta_matrix

# get final y predicted def get_y_for_all_x(current_x, x_matrix,
y_matrix, tau):    theta_matrix = get_theta_matrix(current_x,
x_matrix, y_matrix, tau)    y_predicted_result = current_x *
theta_matrix    return y_predicted_result

# main loop for i in range(total_cases):    y_predicted[i] =
get_y_for_all_x(x_matrix[i], x_matrix, y_matrix, 1)

# Plotting the results sorted_indexes =
x_matrix[:, 1].argsort(0)
sorted_x = x_matrix[sorted_indexes][:, 0]

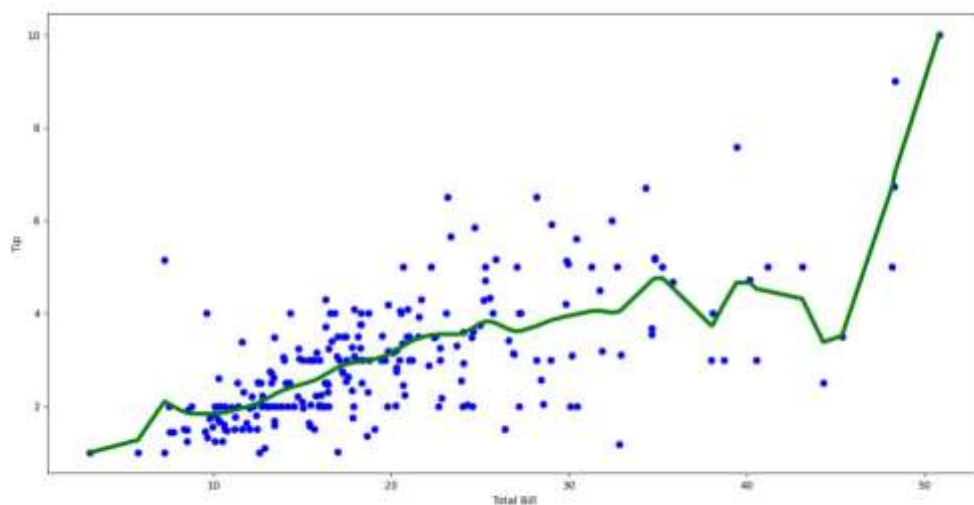
figure = plt.figure()
axes = figure.add_subplot(1, 1, 1)
axes.scatter(feature_array, label_array, color='blue')

axes.plot(sorted_x[:, 1], y_predicted[sorted_indexes], color='green', linewidth=4)

plt.xlabel('Total Bill') plt.ylabel('Tip')
plt.show()

```

Output-



Or

```

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
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

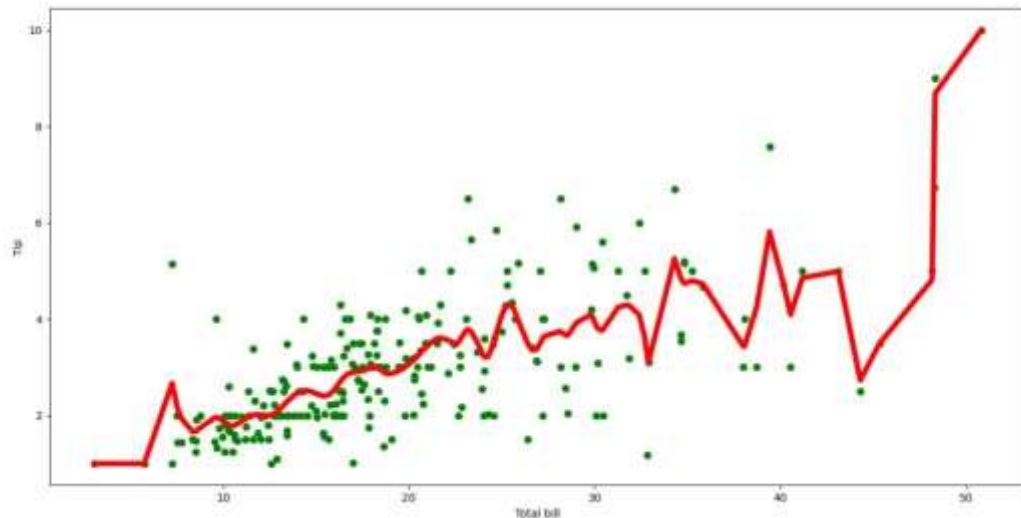
def localWeightRegression(xmat, ymat, k):    m,n =
np.shape(xmat)    ypred = np.zeros(m)    for i in
range(m):        ypred[i] =
xmat[i]*localWeight(xmat[i],xmat,ymat,k)    return ypred

# load data points data =
pd.read_csv('tips.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()

```

Output -



Program - 4

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

```
import numpy as np
X = np.array([[2, 9], [1, 5], [3, 6]], dtype=float)
y = np.array([[92], [86], [89]], dtype=float) 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=10000    lr=0.1
inputlayer_neurons = 2
hiddenlayer_neurons = 3
output_neurons  = 1
#weight initialization
wh=np.random.uniform(size=(inputlayer_neurons,hiddenlayer_neurons))
wout=np.random.uniform(size=(hiddenlayer_neurons,output_neurons)) for
i in range(epoch):
#Forward Propagation
```



```

    hinp=np.dot(X,wh)
    hlayer_act = sigmoid(hinp)
    outinp=np.dot(hlayer_act,wout)
    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
    d_hiddenlayer = EH * hiddengrad    wout +=
    hlayer_act.T.dot(d_output) *lr    wh +=
    X.T.dot(d_hiddenlayer) *lr    print("Input: \n"
+ str(X)) y=y*100 output=output*100
print("Actual Output: \n" + str(y))
print("Predicted Output: \n" ,output)

```

Output

```

Input:
[[2. 9.]
 [1. 5.]
 [3. 6.]]
Actual Output:
[[92.]
 [86.]
 [89.]]
Predicted Output:
[[89.77293616]
 [87.35314244]
 [89.77030837]]

```

Program – 5

Demonstrate Genetic algorithm by taking a suitable data for any simple application.

```

import random

# Define the knapsack problem parameters
max_weight = 50 items
= [
    {'name': 'item1', 'weight': 10, 'value': 60},
    {'name': 'item2', 'weight': 20, 'value': 100},
    {'name': 'item3', 'weight': 30, 'value': 120},
    {'name': 'item4', 'weight': 15, 'value': 50},
    {'name': 'item5', 'weight': 25, 'value': 60},
]

# Genetic algorithm parameters
population_size = 100 generations
= 50
mutation_rate = 0.1

# Initialize a population of random solutions def
initialize_population(pop_size, item_count):
    population = []    for _ in range(pop_size):        solution =
[random.randint(0, 1) for _ in range(item_count)]
    population.append(solution)    return population

# Calculate the fitness of a solution def fitness(solution):    total_weight =
sum(item['weight'] for item, bit in zip(items, solution) if bit)    total_value =
sum(item['value'] for item, bit in zip(items, solution) if bit)    return
total_value if total_weight <= max_weight else 0

# Perform single-point crossover def
crossover(parent1, parent2):
    crossover_point = random.randint(1, len(parent1) - 1)    child1
= parent1[:crossover_point] + parent2[crossover_point:]
    child2 = parent2[:crossover_point] + parent1[crossover_point:]
    return child1, child2

# Mutate a solution def mutate(solution,
mutation_rate):    mutated_solution = []
for bit in solution:        if
random.random() < mutation_rate:
    mutated_solution.append(1 - bit) # Flip
the bit with a probability of mutation_rate
else:
    mutated_solution.append(bit)
return mutated_solution

# Main genetic algorithm loop population =
initialize_population(population_size, len(items)) for generation in
range(generations):    population = sorted(population, key=lambda x:

```

```

fitness(x), reverse=True)    new_population = population[:population_size
// 2]

    for _ in range(population_size // 2):        parent1,
parent2 = random.choices(population, k=2)        child1,
child2 = crossover(parent1, parent2)        child1 =
mutate(child1, mutation_rate)        child2 =
mutate(child2, mutation_rate)
        new_population.extend([child1, child2])

population = new_population

best_solution = max(population, key=fitness) best_value = sum(item['value']
for item, bit in zip(items, best_solution) if bit) best_weight =
sum(item['weight'] for item, bit in zip(items, best_solution) if bit)

print("Best solution:", best_solution) print("Total
value:", best_value)
print("Total weight:", best_weight)

```

Output –

```

Best solution: [0, 1, 1, 0, 0]
Total value: 220
Total weight: 50

```

Program – 6

Demonstrate Q learning algorithm with suitable assumption for a problem statement.

```

import numpy as np

# Estado terminal
terminal = 5

# Possiveis acoes
actions = ['UP','DW','LF','RG']

# Recompensas rws =
np.array([-1]*6)
rws[5] = 10

```

```

# Duas trajetorias
paths = [(0, ['UP','UP','UP','RG']), (4, ['RG','RG','LF','UP'])]

# Constantes alpha
= 0.5 gamma = 0.8

def print_value(value):
    print('[' + str(value[2]) + ' ' + str(value[5]))
    print(str(value[1]) + ' ' + str(value[4]))    print(str(value[0])
    + ' ' + str(value[3]) + ']\n')

def update_value(value, state, action):
    index = actions.index(action)
    next_state = state    rw = 0

    if action == 'UP':        if
state == 2 or state == 5:
rw = -10        else:
        next_state = state + 1

    elif action == 'DW':
        if state == 0 or state == 3:
            rw = -10
    else:
        next_state = state - 1

    elif action == 'LF':        if state == 0 or
state == 1  or state == 2:
        rw = -10
    else:
        next_state = state - 3

    elif action == 'RG':        if state == 3 or
state == 4 or state == 5:        rw = -10
    else:
        next_state = state + 3

    if rw == 0:        rw =
rws[next_state]

    value[index][state] = value[index][state] + alpha * (rw + gamma *
max(value[i][next_state] for i in range(4)) - value[index][state])

    return value, next_state

```

```

def return_policy(value):
    policy = np.array([' ']*6)
    policy[5] = '+10'

    for state in range(5):
        policy[state] =
        actions[np.argmax([value[action][state] for action in range(4)])]

    print(policy[2] + ' ' + policy[5])
    print(policy[1] + ' ' + policy[4])
    print(policy[0] + ' ' + policy[3] + '\n')

def main():
    # Inicializar matriz Q com valores 0, considerando as quatro acoes
    value = [np.zeros(6),np.zeros(6),np.zeros(6),np.zeros(6)]

    for i in range(len(paths)):
        state = paths[i][0]
        actions = paths[i][1]

        for action in actions:
            value, state =
            update_value(value, state, action)

            if state == terminal:
                break

        # Acao UP
        print_value(value[0])

        # Acao DW
        print_value(value[1])    # Acao LF
        print_value(value[2])

        # Acao RG
        print_value(value[3])

        # Politica
        return_policy(value)
    if __name__ ==
    '__main__':
        main()

```

Output-

[-5.0 0.0
-0.5 0.0
-0.5 0.0]

[0.0 0.0
0.0 0.0
0.0 0.0]

[0.0 0.0
0.0 0.0
0.0 0.0]

[5.0 0.0
0.0 0.0
0.0 0.0]

RG +10 DW
UP
DW UP

[-5.0 0.0
1.25 0.0
-0.5 0.0]

[0.0 0.0
0.0 0.0
0.0 0.0]

[0.0 0.0
0.0 -0.5
0.0 0.0]

[5.0 0.0
0.0 -7.5
0.0 0.0]

RG +10
UP UP
DW UP