

### 1)A\* search algorithm

**pip install heuristicsearch**

```
-----  
from heuristicsearch.a_star_search import AStar  
  
graph_nodes = { 'A': [('B', 1), ('C', 3), ('D', 7)], 'B': [('D', 5)], 'C': [('D', 12)] }  
  
heuristics = {'A':1, 'B':1, 'C':1, 'D':1}  
  
graph= AStar(graph_nodes,heuristics)  
  
graph.apply_a_star(start='A',stop='D')
```

### 2) AO\* search algorithm from heuristicsearch.ao\_star import AOSTar

```
print("Graph - 1")  
  
heuristic = { 'A': 1, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7 }  
  
graph_nodes = { 'A': [[('B', 1), ('C', 1)], [('D', 1)]], 'B': [[('G', 1)], [('H', 1)]],  
# 'C': [[('J', 1)]], 'D': [[('E', 1), ('F', 1)]],  
# 'G': [[('I', 1)]] }  
  
graph = AOSTar(graph_nodes, heuristic, 'A')  
  
graph.applyAOSTar()
```

### 3. CandidateEliminationLab3

```
import csv  
  
with open("trainingexamples.csv") as f:  
  
    csv_file = csv.reader(f)  
  
    data = list(csv_file)  
  
    specific = data[1][:-1]  
  
    general = [['?' for i in range(len(specific))] for j in range(len(specific))]
```

```

for i in data:

    if i[-1] == "Yes":

        for j in range(len(specific)):

            if i[j] != specific[j]:

                specific[j] = "?"

                general[j][j] = "?"

    elif i[-1] == "No":

        for j in range(len(specific)):

            if i[j] != specific[j]:

                general[j][j] = specific[j]

            else:

                general[j][j] = "?"

print("\nStep " + str(data.index(i)+1) + " of Candidate Elimination Algorithm")

print(specific)

print(general)

gh = [] # gh = general Hypothesis

for i in general:

    for j in i:

        if j != '?':

            gh.append(i)

```

```

        break

print("\nFinal Specific hypothesis:\n", specific)

print("\nFinal General hypothesis:\n", gh)

```

**Lab 5: 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)
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=5000          #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 Propagation
```

```
    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)
```

```
#how much hidden layer wts contributed to error
```

```
    hiddengrad = derivatives_sigmoid(hlayer_act)
    d_hiddenlayer = EH * hiddengrad
```

```
# dotproduct of nextlayererror and currentlayerop
```

```
    wout += hlayer_act.T.dot(d_output) *lr
    wh += X.T.dot(d_hiddenlayer) *lr
```

```
print("Input: \n" + str(X))
```

```
print("Actual Output: \n" + str(y))
```

```
print("Predicted Output: \n" ,output)
```

**Lab: 7 Apply EM algorithm to cluster a set of data stored in a .CSV file. Use the same data set for clustering using k-Means algorithm. Compare the results of these two algorithms and comment on the quality of clustering. You can add Java/Python ML library classes/API in the program.**

```
import matplotlib.pyplot as plt

from sklearn import datasets

from sklearn.cluster import KMeans

import pandas as pd

import numpy as np


# import some data to play with

iris = datasets.load_iris()

X = pd.DataFrame(iris.data)

X.columns = ['Sepal_Length', 'Sepal_Width', 'Petal_Length', 'Petal_Width']

y = pd.DataFrame(iris.target)

y.columns = ['Targets']


# Build the K Means Model

model = KMeans(n_clusters=3)

model.fit(X) # model.labels_ : Gives cluster no for which samples belongs to


# # Visualise the clustering results

plt.figure(figsize=(14,7))

colormap = np.array(['red', 'lime', 'black'])
```

```
# Plot the Original Classifications using Petal features
```

```
plt.subplot(1, 3, 1)
```

```
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[y.Targets], s=40)
```

```
plt.title('Real Clusters')
```

```
plt.xlabel('Petal Length')
```

```
plt.ylabel('Petal Width')
```

```
# Plot the Models Classifications
```

```
plt.subplot(1, 3, 2)
```

```
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[model.labels_], s=40)
```

```
plt.title('K-Means Clustering')
```

```
plt.xlabel('Petal Length')
```

```
plt.ylabel('Petal Width')
```

```
# General EM for GMM
```

```
from sklearn import preprocessing
```

```
# transform your data such that its distribution will have a # mean value 0 and standard  
deviation of 1.
```

```
scaler = preprocessing.StandardScaler()
```

```
scaler.fit(X)
```

```
xsa = scaler.transform(X)
```

```
xs = pd.DataFrame(xsa, columns = X.columns)
```

```
from sklearn.mixture import GaussianMixture
```

```
gmm = GaussianMixture(n_components=40)
```

```

gmm.fit(xs)

plt.subplot(1, 3, 3)

plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[0], s=40)

plt.title('GMM Clustering')

plt.xlabel('Petal Length')

plt.ylabel('Petal Width')


print('Observation: The GMM using EM algorithm based clustering matched the true labels
more closely than the Kmeans.')

```

**Lab 8: Write a program to implement K-Nearest Neighbour algorithm to classify the iris data set. Print both correct and wrong predictions. Java/Python ML library classes can be used for this problem.**

```

from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn import datasets
iris=datasets.load_iris()
print("Iris Data set loaded...")
x_train, x_test, y_train, y_test = train_test_split(iris.data,iris.target,test_size=0.1)
#random_state=0
for i in range(len(iris.target_names)):
    print("Label", i , "-",str(iris.target_names[i]))
classifier = KNeighborsClassifier(n_neighbors=2)
classifier.fit(x_train, y_train)
y_pred=classifier.predict(x_test)
print("Results of Classification using K-nn with K=1 ")
for r in range(0,len(x_test)):

```

```
print(" Sample:", str(x_test[r]), " Actual-label:", str(y_test[r]), " Predicted-label:",  
str(y_pred[r]))
```

```
print("Classification Accuracy :", classifier.score(x_test,y_test));
```

### **Lab 9: 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 numpy as np
```

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

```
def local_regression(x0, X, Y, tau):
```

```
    x0 = [1, x0]
```

```
    X = [[1, i] for i in X]
```

```
    X = np.asarray(X)
```

```
    xw = (X.T) * np.exp(np.sum((X - x0) ** 2, axis=1) / (-2 * tau))
```

```
    beta = np.linalg.pinv(xw @ X) @ xw @ Y @ x0
```

```
    return beta
```

```
def draw(tau):
```

```
    prediction = [local_regression(x0, X, Y, tau) for x0 in domain]
```

```
    plt.plot(X, Y, 'o', color='black')
```

```
    plt.plot(domain, prediction, color='red')
```

```
    plt.show()
```

```
X = np.linspace(-3, 3, num=1000)
```

```
domain = X
```

```
Y = np.log(np.abs(X ** 2 - 1) + .5)
```



`draw(10)`

`draw(0.1)`

`draw(0.01)`

`draw(0.001)`

lab 4 and 6 is hard