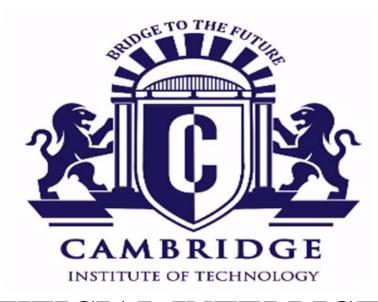
CAMBRIDGE INSTITUTE OF TECHNOLOGY KR PURAM, BENGALURU



AIRTIFICIAL INTELLIGENCE AND MACHINE LEARNING LABORATORY 18CSL76

Department of Computer Science and Engineering

Prerequisites

- 1. Programming experience in Python
- 2. Knowledge of basic Machine Learning Algorithms
- 3. Knowledge of common statistical methods and data analysis best practices.

Lab outcomes:

At the end of the course, the student will be able to;

- 1. Implement and demonstrate AI and ML algorithms.
- 2. Evaluate different algorithms

Software Requirements

- 1. Python version 3.5 and above
- 2. Machine Learning packages
 - Scikit-Learn
 - Numpy matrices and linear algebra
 - Scipy many numerical routines
 - Matplotlib- creating plots of data
 - Pandas –facilitates structured/tabular data manipulation and visualisations
 - Pomegranate –for fast and flexible probabilistic models
- 3. An Integrated Development Environment (IDE) for Python Programming

Anaconda

It contains a list of Python packages, tools like editors, Python distributions include the Python interpreter. Anaconda is one of several Python distributions. Anaconda is a new distribution of the Python and R data science package. It was formerly known as Continuum Analytics. Anaconda has more than 100 new packages. Anaconda is used for scientific computing, data science, statistical analysis, and machine learning

Operating System

Windows/Linux

Anaconda Python distribution is compatible with Linux or windows.

Python

It is an interpreted high-level programming language for general-purpose programming. Created by Guido van Rossum and first released in 1991. Python features a dynamic type system and automatic memory management. It supports multiple programming

paradigms, including object-oriented, imperative, functional and procedural, and has a large and comprehensive standard library.

Python is open source, multi-paradigm, Object-oriented and structured programming supported, Language. Python uses dynamic typing, and a combination of reference counting and a cycle-detecting garbage collector for memory management. It also features dynamic name resolution, which binds method and variable names during program execution.

Python's design offers some support for functional programming in the Lisp tradition. It has filter(), map(), and reduce() functions.

Python packages for machine learning

NumPy

NumPy (stands for Numerical Python) is one of the most famous and commonly used python package among data scientists and ML engineers. This is a part of Python's SciPy Stack, which is basically a collection of software specially designed for scientific computations. It provides several features to work with n-dimensional arrays and matrices in python. This library provides vectorization of mathematical operations on the NumPy array type which adds up to the performance of the execution.

Pandas

The Pandas library is too a well-known library in the world of Analytics and Data Sciences. This package is primarily designed to work with simple and relational data. This is one of the favorite libraries among the data scientists for easy data manipulation, visualization as well as aggregation.

If talking about the data structures, there are basically two prime data structures available in the library which are Series (one-dimensional) & Data Frames (two-dimensional) and we think these are not that significant to talk about as of now.

The basic functionalities that Pandas provides:

- We can very easily delete as well as add a columns from DataFrame
- Pandas can be used to convert the Data Structures in to DataFrame objects.
- If we have any redundancy in the dataset in the form of missing data represented as 'NaN', this is the perfect tool to remove that
- Can be used for grouping of the attributes based strictly on their functionality.

Libraries for Machine Learning

Scikit-Learn: The library is focused on modeling data. Some popular groups of models provided by scikit-learn include:

- Clustering: for grouping unlabeled data such as KMeans.
- Cross Validation: for estimating the performance of supervised models on unseen data.
- Datasets: for test datasets and for generating datasets with specific properties for investigating model behavior.
- Dimensionality Reduction: for reducing the number of attributes in data for summarization, visualization and feature selection such as Principal component analysis.
- Ensemble methods: for combining the predictions of multiple supervised models.
- Feature extraction: for defining attributes in image and text data.
- Feature selection: for identifying meaningful attributes from which to create supervised models.
- Parameter Tuning: for getting the most out of supervised models.
- Manifold Learning: For summarizing and depicting complex multi-dimensional data.
- Supervised Models: a vast array not limited to generalized linear models, discriminate analysis, naive bayes, lazy methods, neural networks, support vector machines and decision trees.

Keras

Keras is a high-level neural networks API, written in Python and capable of running on top of TensorFlow, CNTK, or Theano. It was developed with a focus on enabling fast experimentation. Being able to go from idea to result with the least possible delay is key to doing good research.

- Use Keras if you need a deep learning library that:
- Allows for easy and fast prototyping (through user friendliness, modularity, and extensibility).
- Supports both convolutional networks and recurrent networks, as well as combinations of the two.
- Runs seamlessly on CPU and GPU.

TensorFlow

TensorFlow is an open source software library for numerical computation using dataflow graphs. Nodes in the graph represents mathematical operations, while graph edges represent multi-dimensional data arrays (aka tensors) communicated between them. The flexible architecture allows you to deploy computation to one or more CPUs or GPUs in a desktop, server, or mobile device with a single API

Best Python Machine Learning IDEs

Spyder is coming to our very first focus i.e. *Spyder*. This IDE got this short name from it's name itself: "Scientific Python Development Environment". Pierre Raybaut is the author of Spyder and it got officially released on October 18, 2009 and is written solely in Python.

Features at a glance:

- Very simple and light-weight IDE with detailed documentation and quite easy to install.
- This is an open source editor and supports code completion, introspection, goto definition as well as horizontal and vertical splitting.
- This editor comes with a *Documentation Viewer* where you can see the documentation related to classes or functions you gotta use.

- Like most of the IDEs, this also supports *Variable Explorer* which is a helpful tool to explore and edit the variables that we have created during file execution.
- It supports runtime debugging i.e. the errors will be seen on the screen as soon as you type them.

Geany is primarily a Python machine learning IDE authored by Enrico Troger and got officially released on October 19, 2005. It has been written in C & C++ and is a light-weight IDE. Despite of being a small IDE it is as capable as any other IDE present out there.

Features at a glance:

- Geany's editor supports highlighting of the Syntax and line numebering.
- It comes equipped with the features like code completion, auto closing of braces, auto HTML and XML tags closing.
- It also comes with code folding.
- This IDE supports code navigation.

Rodeo: This is special we got here. This is a Python IDE that primarily focuses and built for the purpose of machine learning and data science. This particular IDE use IPython kernel (you will know this later) and was authored by Yhat.

Features at a glance

- It is mainly famous due to its ability to let users explore, compare and interact with the data frames & plots.
- Like Geany's editor this also comes with a editor that has capability of auto-completion, syntax highlighting.
- This also provides a support for IPython making the code writing fast.
- Also Rodeo comes with Python tutorials integrated within which makes it quite favourable for the users.
- This IDE is well known for the fact that for the data scientists and engineers who work on RStudio IDE can very easily adapt to it

PyCharm is the IDE which is most famous in the professional world whether it is for data science or for conventional Python programming. This IDE is built by one of the big company out there that we all might have heard about: Jetbrains, company released the official version of PyCharm in October 2010.

PyCharm comes in two different editions: Community Edition which we all can have access to essentially for free and second one is the Professional Edition for which you will need to pay some bucks.

Features at a glance

- It includes code completion, auto-indentation and code formatting.
- This also comes with runtime debugger i.e. will display the errors as soon as you type them.
- It contains PEP-8 that enables writing neat codes.

- It consist of debugger for Javascript and Python with a GUI.
- It has one of the most advanced documentation viewer along with video tutorials.
- PyCharm being accepted widely among big companies for the purpose of Machine Learning is due to its ability to provide support for important libraries like Matplotlib, NumPy and Pandas.

JuPyter Notebook or IPython Notebook

It is simple and this became a sensational IDE among the data enthusiasts as it is the descendant of IPython. Best thing about JuPyter is that there you can very easily switch between the different versions of python (or any other language) according to your preference.

Features at a glance

- It's an open source platform
- It can support up to 40 different languages to work on including languages beneficial for data sciences like R, Python, Julia, etc.
- It supports sharing live codes, and even documents with equations and visualizations.
- In JuPyter you can produce outputs in the form of images, videos and even LaTex with the help of several useful widgets.

CONTENTS:			
Expt No.	Title of the Experiments	RBT	СО
1	Implement A* Search algorithm.	L3	1,2,3,4
2	Implement AO* Search algorithm	L3	1,2,3,4
3	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.	L3	1,2,3,4
4	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.	L3	1,2,3,4
5	Build an Artificial Neural Network by implementing the Backpropagation algorithm and test the same using appropriate data sets.	L3	1,2,3,4
6	Write a program to implement the naïve Bayesian classifier for a sample training data set stored as a .CSV file. Compute the accuracy of the classifier, considering few test data sets.		1,2,3,4
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.	L3	1,2,3,4
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.	L3	1,2,3,4
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.	L3	1,2,3,4

Implement A* Search algorithm.

Objective	Finds the shortest path through a search space to goal state using heuristic
	function.
Dataset	Heuristic Values
Description	Finds the shortest path through a search space to goal state using heuristic
	function. It requires the heuristic function to evaluate the cost of path that
	passes through the particular state .

```
#!/usr/bin/env python
# coding: utf-8
# In[1]:
def aStarAlgo(start_node, stop_node):
     open_set = set(start_node)
     closed\_set = set()
     g = \{\} #store distance from starting node
     parents = {}# parents contains an adjacency map of all nodes
     #ditance of starting node from itself is zero
     g[start\_node] = 0
     #start_node is root node i.e it has no parent nodes
     #so start_node is set to its own parent node
     parents[start_node] = start_node
     while len(open\_set) > 0:
       n = None
       #node with lowest f() is found
       for v in open_set:
          if n == None \text{ or } g[v] + heuristic(v) < g[n] + heuristic(n):
             n = v
       if n == stop_node or Graph_nodes[n] == None:
          pass
       else:
          for (m, weight) in get_neighbors(n):
            #nodes 'm' not in first and last set are added to first
             #n is set its parent
             if m not in open_set and m not in closed_set:
               open_set.add(m)
```

```
parents[m] = n
         g[m] = g[n] + weight
       #for each node m,compare its distance from start i.e g(m) to the
       #from start through n node
       else:
         if g[m] > g[n] + weight:
            #update g(m)
            g[m] = g[n] + weight
            #change parent of m to n
            parents[m] = n
            #if m in closed set,remove and add to open
            if m in closed_set:
              closed_set.remove(m)
              open_set.add(m)
  if n == None:
     print('Path does not exist!')
     return None
  # if the current node is the stop_node
  # then we begin reconstructin the path from it to the start_node
  if n == stop_node:
     path = []
     while parents[n] != n:
       path.append(n)
       n = parents[n]
     path.append(start_node)
     path.reverse()
     print('Path found: { }'.format(path))
     return path
  # remove n from the open_list, and add it to closed_list
  # because all of his neighbors were inspected
  open_set.remove(n)
  closed_set.add(n)
print('Path does not exist!')
return None
```

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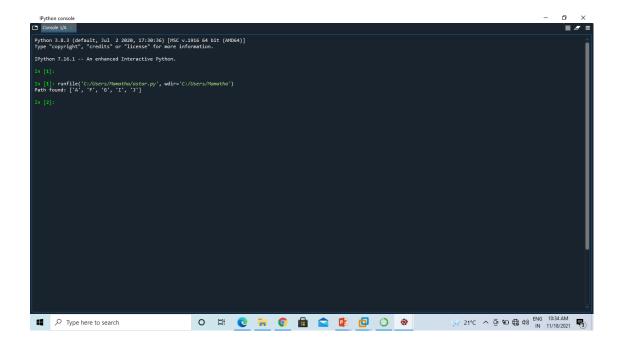
#define fuction to return neighbor and its distance

#from the passed node

```
def get_neighbors(v):
  if v in Graph_nodes:
     return Graph_nodes[v]
  else:
     return None
#for simplicity we ll consider heuristic distances given
#and this function returns heuristic distance for all nodes
def heuristic(n):
     H_dist = {
        'A': 10,
        'B': 8,
        'C': 5,
        'D': 7,
        'E': 3,
        'F': 6,
        'G': 5,
        'H': 3,
        'I': 1,
        'J': 0
      }
     return H_dist[n]
#Describe your graph here
Graph_nodes = {
  'A': [('B', 6), ('F', 3)],
  'B': [('C', 3), ('D', 2)],
  'C': [('D', 1), ('E', 5)],
  'D': [('C', 1), ('E', 8)],
  'E': [('I', 5), ('J', 5)],
  'F': [('G', 1),('H', 7)],
  'G': [('I', 3)],
  'H': [('I', 2)],
  T: [('E', 5), ('J', 3)],
aStarAlgo('A', 'J')
```

Output:

Path found: ['A', 'F', 'G', 'I', 'J']



Implement AO* Search algorithm

Objective	Finds the shortest path through a search space to goal state using heuristic
	function.
Dataset	Heuristic Values.
Description	It is an informed search and works as best first search. AO* is based on problem decomposition. It represents and AND-OR graph algorithm that is used to find more than one solution. It is an efficient methos to explore the solution path.

```
class Graph:
```

def __init__(self, graph, heuristicNodeList, startNode): #instantiate graph object with graph topology, heuristic values, start node

```
self.graph = graph
    self.H=heuristicNodeList
    self.start=startNode
    self.parent={ }
    self.status={ }
    self.solutionGraph={}
  def applyAOStar(self):
                           # starts a recursive AO* algorithm
    self.aoStar(self.start, False)
                           # gets the Neighbors of a given node
  def getNeighbors(self, v):
    return self.graph.get(v,")
  def getStatus(self,v):
                         # return the status of a given node
    return self.status.get(v,0)
  def setStatus(self,v, val): # set the status of a given node
    self.status[v]=val
  def getHeuristicNodeValue(self, n):
    return self.H.get(n,0) # always return the heuristic value of a given node
  def setHeuristicNodeValue(self, n, value):
                         # set the revised heuristic value of a given node
    self.H[n]=value
  def printSolution(self):
    print("FOR GRAPH SOLUTION, TRAVERSE THE GRAPH FROM THE START
NODE:",self.start)
    print("-----")
    print(self.solutionGraph)
    print("-----")
```

```
def computeMinimumCostChildNodes(self, v): # Computes the Minimum Cost of child
nodes of a given node v
    minimumCost=0
    costToChildNodeListDict={}
    costToChildNodeListDict[minimumCost]=[]
    flag=True
    for nodeInfoTupleList in self.getNeighbors(v): # iterate over all the set of child node/s
      cost=0
      nodeList=[]
      for c, weight in nodeInfoTupleList:
        cost=cost+self.getHeuristicNodeValue(c)+weight
        nodeList.append(c)
      if flag==True:
                               # initialize Minimum Cost with the cost of first set of
child node/s
        minimumCost=cost
        costToChildNodeListDict[minimumCost]=nodeList # set the Minimum Cost
child node/s
        flag=False
      else:
                            # checking the Minimum Cost nodes with the current
Minimum Cost
        if minimumCost>cost:
           minimumCost=cost
           costToChildNodeListDict[minimumCost]=nodeList # set the Minimum Cost
child node/s
    return minimumCost, costToChildNodeListDict[minimumCost] # return Minimum
Cost and Minimum Cost child node/s
  def aoStar(self, v, backTracking): # AO* algorithm for a start node and backTracking
status flag
    print("HEURISTIC VALUES :", self.H)
    print("SOLUTION GRAPH :", self.solutionGraph)
    print("PROCESSING NODE :", v)
    print("-----")
    if self.getStatus(v) \geq 0: # if status node v \geq 0, compute Minimum Cost nodes of v
      minimumCost, childNodeList = self.computeMinimumCostChildNodes(v)
      self.setHeuristicNodeValue(v, minimumCost)
      self.setStatus(v,len(childNodeList))
                            # check the Minimum Cost nodes of v are solved
      solved=True
      for childNode in childNodeList:
        self.parent[childNode]=v
        if self.getStatus(childNode)!=-1:
           solved=solved & False
```

```
if solved==True: # if the Minimum Cost nodes of v are solved, set the current node status as solved(-1) self.setStatus(v,-1) self.solutionGraph[v]=childNodeList # update the solution graph with the solved nodes which may be a part of solution
```

if v!=self.start: # check the current node is the start node for backtracking the current node value

self.aoStar(self.parent[v], True) # backtracking the current node value with backtracking status set to true

if backTracking==False: # check the current call is not for backtracking
for childNode in childNodeList: # for each Minimum Cost child node
self.setStatus(childNode,0) # set the status of child node to 0(needs exploration)
self.aoStar(childNode, False) # Minimum Cost child node is further explored
with backtracking status as false

```
h1 = {'A': 1, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1, 'T': 3}
graph1 = {
  'A': [[('B', 1), ('C', 1)], [('D', 1)]],
  'B': [[('G', 1)], [('H', 1)]],
  'C': [[('J', 1)]],
  'D': [[('E', 1), ('F', 1)]],
  'G': [[('I', 1)]]
G1= Graph(graph1, h1, 'A')
G1.applyAOStar()
G1.printSolution()
h2 = {'A': 1, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7} # Heuristic values of Nodes
graph2 = {
                                      # Graph of Nodes and Edges
                                         # Neighbors of Node 'A', B, C & D with repective
  'A': [[('B', 1), ('C', 1)], [('D', 1)]],
weights
                                       # Neighbors are included in a list of lists
  'B': [[('G', 1)], [('H', 1)]],
                                      # Each sublist indicate a "OR" node or "AND" nodes
  'D': [[('E', 1), ('F', 1)]]
}
                                             # Instantiate Graph object with graph, heuristic
G2 = Graph(graph2, h2, 'A')
values and start Node
G2.applyAOStar()
                                          # Run the AO* algorithm
                                         # Print the solution graph as output of the AO*
G2.printSolution()
algorithm search
```

Output:

```
HEURISTIC VALUES: {'A': 1, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1,
T': 3
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1,
'T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: B
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1,
'T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE : A
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 5, 'H': 7, 'I': 7, 'J': 1,
'T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE : G
HEURISTIC VALUES: {'A': 10, 'B': 6, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1,
'T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: B
_____
HEURISTIC VALUES: {'A': 10, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1,
'T': 3}
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 7, 'J': 1,
SOLUTION GRAPH : {}
PROCESSING NODE: I
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 8, 'H': 7, 'I': 0, 'J': 1,
'T': 3}
SOLUTION GRAPH : {'I': []}
PROCESSING NODE: G
HEURISTIC VALUES: {'A': 12, 'B': 8, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1,
SOLUTION GRAPH : {'I': [], 'G': ['I']}
PROCESSING NODE: B
HEURISTIC VALUES: {'A': 12, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1,
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
```

```
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1,
T': 3
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE : C
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1,
T': 3
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 1,
'T': 3}
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G']}
PROCESSING NODE : J
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 2, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 0,
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G'], 'J': []}
PROCESSING NODE : C
HEURISTIC VALUES: {'A': 6, 'B': 2, 'C': 1, 'D': 12, 'E': 2, 'F': 1, 'G': 1, 'H': 7, 'I': 0, 'J': 0,
SOLUTION GRAPH : {'I': [], 'G': ['I'], 'B': ['G'], 'J': [], 'C': ['J']}
PROCESSING NODE: A
FOR GRAPH SOLUTION, TRAVERSE THE GRAPH FROM THE START NODE: A
   _____
{'I': [], 'G': ['I'], 'B': ['G'], 'J': [], 'C': ['J'], 'A': ['B', 'C']}
HEURISTIC VALUES: {'A': 1, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7}
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 11, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7}
SOLUTION GRAPH : {}
PROCESSING NODE: D
HEURISTIC VALUES: {'A': 11, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7}
SOLUTION GRAPH : {}
PROCESSING NODE: A
HEURISTIC VALUES: {'A': 11, 'B': 6, 'C': 12, 'D': 10, 'E': 4, 'F': 4, 'G': 5, 'H': 7}
SOLUTION GRAPH : {}
PROCESSING NODE : E
HEURISTIC VALUES: {'A': 11, 'B': 6, 'C': 12, 'D': 10, 'E': 0, 'F': 4, 'G': 5, 'H': 7}
SOLUTION GRAPH : {'E': []}
```

PROCESSING NODE : D

HEURISTIC VALUES: {'A': 11, 'B': 6, 'C': 12, 'D': 6, 'E': 0, 'F': 4, 'G': 5, 'H': 7}

SOLUTION GRAPH : {'E': []} PROCESSING NODE : A

HEURISTIC VALUES: {'A': 7, 'B': 6, 'C': 12, 'D': 6, 'E': 0, 'F': 4, 'G': 5, 'H': 7}

SOLUTION GRAPH : {'E': []} PROCESSING NODE : F

HEURISTIC VALUES: {'A': 7, 'B': 6, 'C': 12, 'D': 6, 'E': 0, 'F': 0, 'G': 5, 'H': 7}

SOLUTION GRAPH : {'E': [], 'F': []}

PROCESSING NODE : D

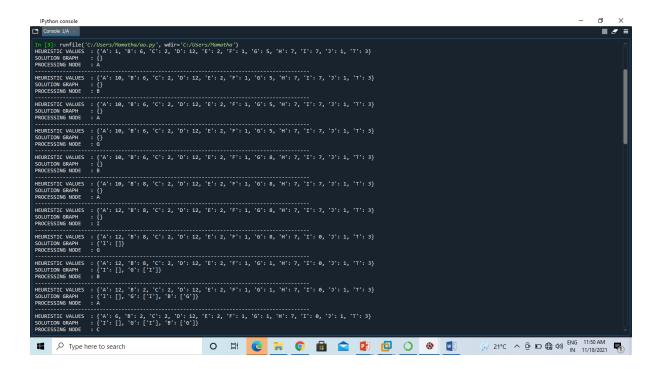
HEURISTIC VALUES: {'A': 7, 'B': 6, 'C': 12, 'D': 2, 'E': 0, 'F': 0, 'G': 5, 'H': 7}

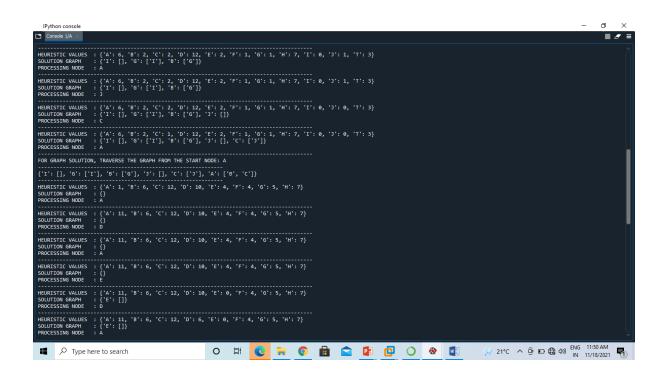
SOLUTION GRAPH : {'E': [], 'F': [], 'D': ['E', 'F']}

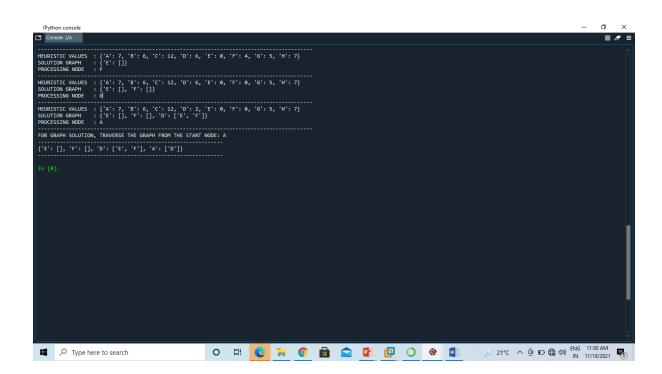
PROCESSING NODE: A

FOR GRAPH SOLUTION, TRAVERSE THE GRAPH FROM THE START NODE: A

{'E': [], 'F': [], 'D': ['E', 'F'], 'A': ['D']}







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.

Incrementally builds the version space given a hypothesis space H and a set	
E of examples.	
Tennis data set: This data set contain the set of example days on which playing	
of tennis is possible or not. Based on attribute Sky, AirTemp, Humidity,	
Wind, Water and Forecast. The dataset has 14 instances.	
Supervised Learning-Candidate elimination algorithm	
The candidate elimination algorithm incrementally builds the version space given a hypothesis space H and a set E of examples The candidate elimination algorithm does this by updating the general and specific boundary for each new example.	

```
import csv
def get_domains(examples):
d = [set() \text{ for i in examples}[0]]
for x in examples:
for i, xi in enumerate(x):
d[i].add(xi)
return [list(sorted(x)) for x in d]
def more_general(h1, h2):
more_general_parts = []
for x, y in zip(h1, h2):
mg = x == "?" \text{ or } (x != "0" \text{ and } (x == y \text{ or } y == "0"))
more_general_parts.append(mg)
return all(more general parts)
def fulfills(example, hypothesis):
# the implementation is the same as for hypotheses:
return more_general(hypothesis, example)
def min_generalizations(h, x):
h_new = list(h)
for i in range(len(h)):
if not fulfills(x[i:i+1], h[i:i+1]):
h_{new}[i] = '?' \text{ if } h[i] != '0' \text{ else } x[i]
return [tuple(h_new)]
def min specializations(h, domains, x):
results = []
for i in range(len(h)):
if h[i] == "?":
for val in domains[i]:
if x[i] != val:
h_new = h[:i] + (val_i) + h[i+1:]
results.append(h_new)
elif h[i] != "0":
h_new = h[:i] + ('0',) + h[i+1:]
```

```
results.append(h_new)
return results
def generalizeS(x, G, S):
S prev = list(S)
for s in S_prev:
if s not in S:
continue
if not fulfills(x, s):
S.remove(s)
Splus = min generalizations(s, x)
## keep only generalizations that have a counterpart in G
S.update([h for h in Splus if any([more_general(g,h)
for g in G])])
## remove hypotheses less specific than any other in S
S.difference_update([h for h in S if
any([more_general(h, h1)
for h1 in S if h != h1])])
return S
def specializeG(x, domains, G, S):
G_prev = list(G)
for g in G_prev:
if g not in G:
continue
if fulfills(x, g):
G.remove(g)
Gminus = min specializations(g, domains, x)
## keep only specializations that have a conuterpart in S
G.update([h for h in Gminus if any([more_general(h, s)
for s in S()))
## remove hypotheses less general than any other in G
G.difference_update([h for h in G if
any([more_general(g1, h)
for g1 in G if h != g1])])
return G
def candidate_elimination(examples):
domains = get domains(examples)[:-1]
n = len(domains)
G = set([("?",)*n])
S = set([("0",)*n])
print("Maximally specific hypotheses - S ")
print("Maximally general hypotheses - G")
print("\nS[0]:",str(S),"\nG[0]:",str(G))
for xcx in examples:
x, cx = xcx[:-1], xcx[-1] # Splitting data into attributes and decisions
if cx=='Y': # x is positive example
G = \{g \text{ for } g \text{ in } G \text{ if fulfills}(x, g)\}
S = generalize\_S(x, G, S)
```

```
else: # x is negative example
S = {s for s in S if not fulfills(x, s)}
G = specialize_G(x, domains, G, S)
print("\nS[{0}]:".format(i),S)
print("G[{0}]:".format(i),G)
return
with open('data22_sports.csv') as csvFile:
examples = [tuple(line) for line in csv.reader(csvFile)]
candidate_elimination(examples)
```

Output:

Data: data21_sports.csv (Sky,AirTemp,Humidity,Wind,Water,Forecast,EnjoySport)

```
sunny,warm,normal,strong,warm,same,Y sunny,warm,high,strong,warm,same,Y rainy,cold,high,strong,warm,change,N sunny,warm,high,strong,cool,change,Y
```

Maximally specific hypotheses - S

Output:

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.

Objective	To demonstrate the working of the decision tree based ID3 algorithm.
Dataset	Tennis data set: This data set contain the set of example days on which playing
	of tennis is possible or not. Based on attribute Sky, AirTemp, Humidity,
	Wind, Water and Forecast.
ML	Supervised Learning-Decision Tree algorithm
algorithm	
Description	Decision tree builds classification or regression models in the form of a tree
	structure. It breaks down a dataset into smaller and smaller subsets while at
	the same time an associated decision tree is incrementally developed. The
	final result is a tree with decision nodes and leaf nodes.

```
import math
import csv
def load_csv(filename):
lines = csv.reader(open(filename, "r"));
dataset = list(lines)
headers = dataset.pop(0)
return dataset, headers
class Node:
def __init__(self, attribute):
self.attribute = attribute
self.children = []
self.answer = "" # NULL indicates children exists.
# Not Null indicates this is a Leaf Node
def subtables(data, col, delete):
dic = \{\}
coldata = [row[col] for row in data]
attr = list(set(coldata)) # All values of attribute retrived
for k in attr:
dic[k] = []
for y in range(len(data)):
key = data[y][col]
if delete:
del data[y][col]
dic[key].append(data[y])
return attr, dic
def entropy(S):
attr = list(set(S))
if len(attr) == 1: #if all are +ve/-ve then entropy = 0
return 0
counts = [0,0] # Only two values possible 'yes' or 'no'
for i in range(2):
counts[i] = sum( [1 for x in S if attr[i] == x] ) / (len(S) * 1.0)
sums = 0
```

```
for cnt in counts:
sums += -1 * cnt * math.log(cnt, 2)
return sums
def compute gain(data, col):
attValues, dic = subtables(data, col, delete=False)
total_entropy = entropy([row[-1] for row in data])
for x in range(len(attValues)):
ratio = len(dic[attValues[x]]) / (len(data) * 1.0)
entro = entropy([row[-1] for row in dic[attValues[x]]])
total entropy -= ratio*entro
return total_entropy
def build tree(data, features):
lastcol = [row[-1] for row in data]
if (len(set(lastcol))) == 1: # If all samples have same labels return that label
node=Node("")
node.answer = lastcol[0]
return node
n = len(data[0])-1
gains = [compute_gain(data, col) for col in range(n)]
split = gains.index(max(gains)) # Find max gains and returns index
node = Node(features[split]) # 'node' stores attribute selected
#del (features[split])
fea = features[:split]+features[split+1:]
attr, dic = subtables(data, split, delete=True) # Data will be spilt in subtables
for x in range(len(attr)):
child = build tree(dic[attr[x]], fea)
node.children.append((attr[x], child))
return node
def print tree(node, level):
if node.answer != "":
print(" "*level, node.answer) # Displays leaf node yes/no
print(" "*level, node.attribute) # Displays attribute Name
for value, n in node.children:
print(" "*(level+1), value)
print tree(n, level + 2)
def classify(node,x_test,features):
if node.answer != "":
print(node.answer)
return
pos = features.index(node.attribute)
for value, n in node.children:
if x_test[pos]==value:
classify(n,x test,features)
" Main program "
dataset, features = load_csv("data3.csv") # Read Tennis data
node = build_tree(dataset, features) # Build decision tree
print("The decision tree for the dataset using ID3 algorithm is ")
print tree(node, 0)
```

```
testdata, features = load_csv("data3_test.csv")
for xtest in testdata:
print("The test instance : ",xtest)
print("The predicted label : ", end="")
classify(node,xtest,features)
Output:
Training instances: data3.csv
Outlook, Temperature, Humidity, Wind, Target
sunny,hot,high,weak,no
sunny,hot,high,strong,no
overcast,hot,high,weak,yes
rain,mild,high,weak,yes
rain,cool,normal,weak,yes
rain,cool,normal,strong,no
overcast, cool, normal, strong, yes
sunny,mild,high,weak,no
sunny,cool,normal,weak,yes
rain, mild, normal, weak, yes
sunny,mild,normal,strong,yes
overcast, mild, high, strong, yes
overcast,hot,normal,weak,yes
rain, mild, high, strong, no
Testing instances: data3_test.csv
Outlook, Temperature, Humidity, Wind
rain,cool,normal,strong
sunny,mild,normal,strong
Output:
The decision tree for the dataset using ID3 algorithm is
Outlook
overcast
yes
rain
Wind
weak
```

yes

strong

no

sunny

Humidity

normal

yes

high

no

The test instance : ['rain', 'cool', 'normal', 'strong']

The predicted label: no

The test instance : ['sunny', 'mild', 'normal', 'strong']

The predicted label: yes

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

Objective	To build an artificial neural network using the backpropagation algorithm.
Dataset	Data stored as a list having two features- number of hours slept, number of
	hours studied with the test score being the class label
ML	Supervised Learning –Backpropagation algorithm
algorithm	
Description	The neural network using back propagation will model a single hidden layer with three inputs and one output. The network will be predicting the score of an exam based on the inputs of number of hours studied and the number of
	hours slept the day before. The test score is the output.

```
import numpy as np
\# X = (hours sleeping, hours studying), y = score on test
X = \text{np.array}(([2, 9], [1, 5], [3, 6]), \text{dtype=float})
y = np.array(([92], [86], [89]), dtype=float)
# scale units
X = X/np.amax(X, axis=0) \# maximum of X array
y = y/100 \# max test score is 100
class Neural_Network(object):
 def __init__(self):
  #parameters
  self.inputSize = 2
  self.outputSize = 1
  self.hiddenSize = 3
  #weights
  self.W1 = np.random.randn(self.inputSize, self.hiddenSize) # (3x2) weight matrix from
input to hidden layer
  self.W2 = np.random.randn(self.hiddenSize, self.outputSize) # (3x1) weight matrix from
hidden to output layer
 def forward(self, X):
  #forward propagation through our network
  self.z = np.dot(X, self.W1) # dot product of X (input) and first set of 3x2 weights
  self.z2 = self.sigmoid(self.z) # activation function
  self.z3 = np.dot(self.z2, self.W2) \# dot product of hidden layer (z2) and second set of 3x1
  o = self.sigmoid(self.z3) # final activation function
  return o
 def sigmoid(self, s):
  # activation function
  return 1/(1+np.exp(-s))
```

```
def sigmoidPrime(self, s):
  #derivative of sigmoid
  return s * (1 - s)
 def backward(self, X, y, o):
  # backward propgate through the network
  self.o_error = y - o # error in output
  self.o_delta = self.o_error*self.sigmoidPrime(o) # applying derivative of sigmoid to error
  self.z2 error = self.o delta.dot(self.W2.T) # z2 error: how much our hidden layer weights
contributed to output error
  self.z2_delta = self.z2_error*self.sigmoidPrime(self.z2) # applying derivative of sigmoid
to z2 error
  self.W1 += X.T.dot(self.z2_delta) # adjusting first set (input --> hidden) weights
  self.W2 += self.z2.T.dot(self.o_delta) # adjusting second set (hidden --> output) weights
 def train (self, X, y):
  o = self.forward(X)
  self.backward(X, y, o)
NN = Neural Network()
for i in range(10): # trains the NN 1,000 times
 print(i)
 print ("Input: \n" + str(X))
 print ("Actual Output: \n" + str(y))
 print ("Predicted Output: \n" + str(NN.forward(X)))
 print ("Loss: \n" + str(np.mean(np.square(y - NN.forward(X)))) )# mean sum squared loss
 print ("\n")
 NN.train(X, y)
```

OUTPUT:

```
Input:
[[0.666666667 1. ]
[0.333333333 0.55555556]
[1. 0.66666667]]
Actual Output:
[[0.92]
[0.86]
[0.89]]
Predicted Output:
[[0.8930243 ]
[0.86917478]
[0.90947008]]
Loss:
0.0003969829742387784
```

```
Input:
[[0.66666667 1.
[0.33333333 0.55555556]
        0.66666667]]
Actual Output:
[[0.92]]
[0.86]
[0.89]]
Predicted Output:
[[0.89325922]
[0.86906258]
[0.90922387]]
Loss:
0.00038891893394468784
###output in the last epoch......
Input:
[[0.66666667 1.
                  ]
[0.33333333 0.55555556]
        0.66666667]]
[1.
Actual Output:
[[0.92]]
[0.86]
[0.89]]
Predicted Output:
[[0.89870965]
[0.86523337]
[0.90596392]]
Loss:
0.00024517132775894306
```

Write a program to implement the naïve Bayesian classifier for a sample training data set stored as a .CSV file. Compute the accuracy of the classifier, considering few test data sets.

Objective	To implement a classification model for a sample training dataset and		
	computing the accuracy of the classifier for test data.		
Dataset	Pima Indian Diabetes dataset stored as .CSV .The attributes are Number of		
	times pregnant, Plasma glucose concentration, Blood Pressure, Triceps skin		
	fold thickness, serum insulin, Body mass index, Diabetes pedigree function,		
	Age		
ML	Supervised Learning -Naïve Bayes Algorithm		
algorithm			
Description	The Naïve Bayes classifier is a probabilistic classifier that is based on Bayes		
	Theorem. The algorithm builds a model assuming that the attributes in the		
	dataset are independent of each other.		

Program

def mean(numbers):

```
import csv
import random
import math
def loadCsv(filename):
       lines = csv.reader(open(filename, "r"))
       dataset = list(lines)
       for i in range(len(dataset)):
               dataset[i] = [float(x) for x in dataset[i]]
       return dataset
def splitDataset(dataset, splitRatio):
       trainSize = int(len(dataset) * splitRatio)
       trainSet = []
       copy = list(dataset)
       while len(trainSet) < trainSize:
               index = random.randrange(len(copy))
               trainSet.append(copy.pop(index))
       return [trainSet, copy]
def separateByClass(dataset):
       separated = \{\}
       for i in range(len(dataset)):
               vector = dataset[i]
               if (vector[-1] not in separated):
                       separated[vector[-1]] = []
               separated[vector[-1]].append(vector)
       return separated
```

```
return sum(numbers)/float(len(numbers))
def stdev(numbers):
       avg = mean(numbers)
       variance = sum([pow(x-avg,2) for x in numbers])/float(len(numbers)-1)
       return math.sqrt(variance)
def summarize(dataset):
       summaries = [(mean(attribute), stdev(attribute)) for attribute in zip(*dataset)]
       del summaries[-1]
       return summaries
def summarizeByClass(dataset):
       separated = separateByClass(dataset)
       summaries = { }
       for classValue, instances in separated.items():
              summaries[classValue] = summarize(instances)
       return summaries
def calculateProbability(x, mean, stdev):
       exponent = math.exp(-(math.pow(x-mean,2)/(2*math.pow(stdev,2))))
       return (1 / (math.sqrt(2*math.pi) * stdev)) * exponent
def calculateClassProbabilities(summaries, inputVector):
       probabilities = {}
       for classValue, classSummaries in summaries.items():
              probabilities[classValue] = 1
              for i in range(len(classSummaries)):
                      mean, stdev = classSummaries[i]
                      x = inputVector[i]
                      probabilities[classValue] *= calculateProbability(x, mean, stdev)
       return probabilities
def predict(summaries, inputVector):
       probabilities = calculateClassProbabilities(summaries, inputVector)
       bestLabel, bestProb = None, -1
       for class Value, probability in probabilities.items():
              if bestLabel is None or probability > bestProb:
                      bestProb = probability
                      bestLabel = classValue
       return bestLabel
def getPredictions(summaries, testSet):
       predictions = []
       for i in range(len(testSet)):
              result = predict(summaries, testSet[i])
              predictions.append(result)
       return predictions
```

```
def getAccuracy(testSet, predictions):
       correct = 0
       for i in range(len(testSet)):
               if testSet[i][-1] == predictions[i]:
                      correct += 1
       return (correct/float(len(testSet))) * 100.0
def main():
       filename = 'naivedata.csv'
       splitRatio = 0.67
       dataset = loadCsv(filename)
       trainingSet, testSet = splitDataset(dataset, splitRatio)
       print('Split {0} rows into train={1} and test={2} rows'.format(len(dataset),
len(trainingSet), len(testSet)))
       # prepare model
       summaries = summarizeByClass(trainingSet)
       # test model
       predictions = getPredictions(summaries, testSet)
       accuracy = getAccuracy(testSet, predictions)
       print('Accuracy of the classifier is : {0}%'.format(accuracy))
main()
```

OUTPUT:

Split 768 rows into train=514 and test=254 rows Accuracy of the classifier is: 74.80314960629921%

Apply EM algorithm to cluster a set of data stored in a .csv file. Use the same data set for clustering 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.

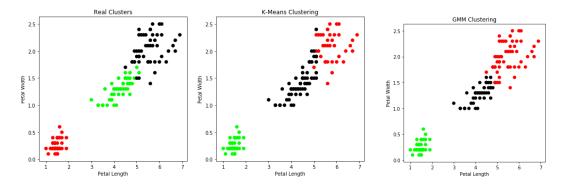
Objective	To group a set of unlabelled data into similar classes/clusters and label them			
	and to compare the quality of algorithm.			
Dataset	Delivery fleet driver dataset Data set in .csv file with features "Driver_ID",			
	"distance_feature", "speeding_feature" having more than 20 instances			
ML	EM algorithm, K means algorithm – Unsupervised clustering			
algorithm				
Packages	Scikit learn(sklearn),pandas			
Description	EM algorithm – soft clustering - can be used for variable whose value is never			
	directly observed, provided the general probability distribution governing			
	these varaiable is known. EM algorithm can be used to train Bayesian belief			
	networks as well as radial basis function network.			
	K-Means – Hard Clustering - to find groups in the data, with the number			
	of groups represented by the variable K. The algorithm works iteratively			
	to assign each data point to one of K groups based on the features that are			
	provided. Data points are clustered based on feature similarity.			

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=(14,14))
colormap = np.array(['red', 'lime', 'black'])
# Plot the Original Classifications using Petal features
plt.subplot(2, 2, 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(2, 2, 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=3)
gmm.fit(xs)
gmm_y = gmm.predict(xs)
plt.subplot(2, 2, 3)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[gmm_y], 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.')
```

Output:



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.

Objective	To implement a classification model that classifies a set of documents and
	calculates the accuracy, precision and recall for the dataset
Dataset	IRIS data set with features "petal_length", "petal_width", "sepal_length",
	"sepal_width" having more than 150 instances
ML	Supervised Learning – Lazy learning algorithm
algorithm	
Packages	Scikit learn(sklearn),pandas
Description	When we get training set/instances, machine won't learn or a model can't be
	built. Instead instances/examples will be just stored in memory. Test instance
	is given, attempt to find the closest instance/most neighboring instances in the
	instance space

```
from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, confusion_matrix
#from sklearn import metrics
#import pandas as pd
#import numpy as np
#import matplotlib.pyplot as plt
from sklearn import datasets
iris=datasets.load_iris()
iris data=iris.data
iris labels=iris.target
#print(iris_data)
#print(iris_labels)
x_train,x_test,y_train,y_test=train_test_split(iris_data,iris_labels,test_size=0.30)
classifier=KNeighborsClassifier(n_neighbors=5)
classifier.fit(x_train,y_train)
y_pred=classifier.predict(x_test)
print('Confusion matrix is as follows')
print(confusion_matrix(y_test,y_pred))
print('Accuracy Metrics')
print(classification_report(y_test,y_pred))
```

OUTPUT:

Confusion matrix is as follows

[[11 0 0]

[091]

[0 1 8]]

Accuracy Metrics

	precision		recall f1-score		support	
	0	1.00	1.00	1.00	11	
	1	0.90	0.90	0.90	10	
	2	0.89	0.89	0.89	9	
avg / to	tal	0.93	0.93	0.93	30	

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

Objective	To implement Regression algorithm to fit the given data		
Dataset	The dataset contains billing information based on the attributes		
	total_bill,tip,sex,smoker,day,time,size		
ML	Locally Weighted Regression Algorithm – Instance Based learning		
algorithm			
Description	Regression means approximating a real valued target function. Given a new		
	query instance X_q , the general approach is to construct an approximation		
	function F that fits the training example in the neighbourhood surrounding		
	X_q . This approximation is then used to estimate the target value $F(X_q)$		

Program

import matplotlib.pyplot as plt import pandas as pd import numpy as np1

```
def kernel(point,xmat, k):
  m,n = np1.shape(xmat)
  weights = np1.mat(np1.eye((m)))
  for j in range(m):
    diff = point - X[i]
    weights[j,j] = np1.exp(diff*diff.T/(-2.0*k**2))
  return weights
def localWeight(point,xmat,ymat,k):
  wei = kernel(point,xmat,k)
  B = (X.T*(wei*X)).I*(X.T*(wei*ymat.T))
  return B
def localWeightRegression(xmat,ymat,k):
  m,n = np1.shape(xmat)
  ypred = np1.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 = np1.array(data.total_bill)
tip = np1.array(data.tip)
#preparing and add 1 in bill
mbill = np1.mat(bill)
```

```
mtip = np1.mat(tip)
m= np1.shape(mbill)[1]
print(m)
one = np1.mat(np1.ones(m))
X= np1.hstack((one.T,mbill.T))
#set k here
ypred = localWeightRegression(X,mtip,0.3)
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:

