

VISVESVARAYA TECHNOLOGICAL UNIVERSITY

“JnanaSangama”, Belgaum -590014, Karnataka.



LAB REPORT
on

MACHINE LEARNING **(20CS6PCMAL)**

Submitted by

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in partial fulfillment for the award of the degree of
BACHELOR OF ENGINEERING
in
COMPUTER SCIENCE AND ENGINEERING



B.M.S. COLLEGE OF ENGINEERING

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CERTIFICATE

This is to certify that the Lab work entitled “**MACHINE LEARNING**” was carried out by **SAMPREETH P (1BM19CS142)**, who is a bona fide student of **B. M. S. College of Engineering**. It is in partial fulfillment for the award of **Bachelor of Engineering in Computer Science and Engineering** of the Visvesvaraya Technological University, Belgaum during the year 2022. The Lab report has been approved as it satisfies the academic requirements in respect of the course **MACHINE LEARNING(20CS6PCMAL)** work prescribed for the said degree.

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Course Outcome:-

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

| | |
|-----|---|
| CO1 | Ability to apply the different learning algorithms. |
| CO2 | Ability to analyze the learning techniques for given dataset. |
| CO3 | Ability to design a model using machine learning to solve a problem. |
| CO4 | Ability to conduct practical experiments to solve problems using appropriate machine learning techniques. |

Lab Program -1 :-

Implement and demonstrate the FIND-S algorithm for finding the most specific hypothesis based on a given set of training data samples.

Source code and output :-

```
+*In[1]:*+
[source, ipython3]
----
import csv
hypo = ['%', '%', '%', '%', '%', '%'];

with open(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\lab 1\finds.csv') as csv_file:
    readcsv = csv.reader(csv_file, delimiter=',')
    print(readcsv)

    data = []
    print("\nThe given training examples are:")
    for row in readcsv:
        print(row)
        if row[len(row)-1].upper() == "YES":
            data.append(row)
    ----

+*Out[1]:*+
----
<_csv.reader object at 0x0000013B7E4DFD60>
```

The given training examples are:

```
['sky', 'air temp', 'humidity', 'wind', 'water', 'forecast', 'enjoy sport']
```

```
['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']
```

```
+*In[2]:*+
```

```
[source, ipython3]
```

```
print("\nThe positive examples are:");
```

```
for x in data:
```

```
    print(x);
```

```
print("\n");
```

```
+*Out[2]:*+
```

The positive examples are:

```
['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes']
```

```
['sunny', 'warm', 'high', 'strong', 'warm', 'same', 'yes']
```

```
['sunny', 'warm', 'high', 'strong', 'cool', 'change', 'yes']
```

+*ln[3]:*+

[source, ipython3]

TotalExamples = len(data);

i=0;

j=0;

k=0;

print("The steps of the Find-s algorithm are :\n",hypo);

list = [];

p=0;

d=len(data[p])-1;

for j in range(d):

list.append(data[i][j]);

hypo=list;

i=1;

for i in range>TotalExamples):

for k in range(d):

if hypo[k]!=data[i][k]:

hypo[k]='?';

k=k+1;

else:

hypo[k];

print(hypo);

i=i+1;

```
+*Out[3]:*+
```

```
----
```

The steps of the Find-s algorithm are :

```
['%', '%', '%', '%', '%', '%']
```

```
['sunny', 'warm', 'normal', 'strong', 'warm', 'same']
```

```
['sunny', 'warm', '?', 'strong', 'warm', 'same']
```

```
['sunny', 'warm', '?', 'strong', '?', '?']
```

```
----
```

```
+*In[4]:*+
```

```
[source, ipython3]
```

```
----
```

```
print("\nThe maximally specific Find-s hypothesis for the given training examples is :");
```

```
list=[];
```

```
for i in range(d):
```

```
    list.append(hypo[i]);
```

```
print(list);
```

```
----
```

```
+*Out[4]:*+
```

```
----
```

The maximally specific Find-s hypothesis for the given training examples is :

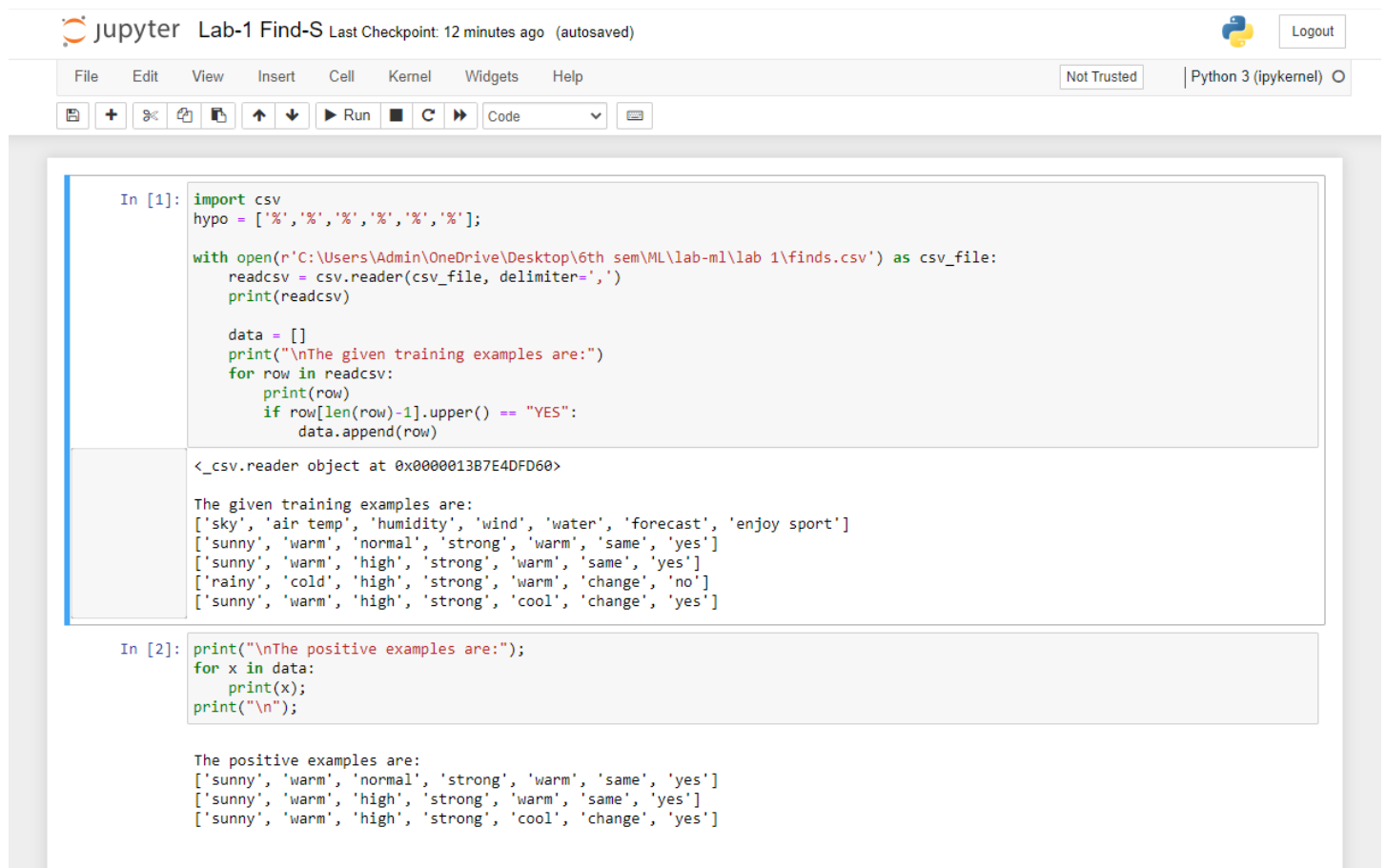
```
['sunny', 'warm', '?', 'strong', '?', '?']
```

```
----
```


+*In[]:*

[source, ipython3]

Output screenshots :-



The screenshot displays a JupyterLab environment with the following components:

- Header:** "jupyter Lab-1 Find-S Last Checkpoint: 12 minutes ago (autosaved)" and a "Logout" button.
- Menu Bar:** File, Edit, View, Insert, Cell, Kernel, Widgets, Help.
- Toolbar:** Includes icons for file operations, a "Run" button, and a "Code" dropdown menu.
- Code Cell 1 (In [1]):**

```
import csv
hypo = ['%', '%', '%', '%', '%', '%'];

with open(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\lab 1\finds.csv') as csv_file:
    readcsv = csv.reader(csv_file, delimiter=',')
    print(readcsv)

    data = []
    print("\nThe given training examples are:")
    for row in readcsv:
        print(row)
        if row[len(row)-1].upper() == "YES":
            data.append(row)
```

The output of this cell is:

```
<_csv.reader object at 0x0000013B7E4DFD60>

The given training examples are:
['sky', 'air temp', 'humidity', 'wind', 'water', 'forecast', 'enjoy sport']
['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']
```
- Code Cell 2 (In [2]):**

```
print("\nThe positive examples are:");
for x in data:
    print(x);
print("\n");
```

The output of this cell is:

```
The positive examples are:
['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes']
['sunny', 'warm', 'high', 'strong', 'warm', 'same', 'yes']
['sunny', 'warm', 'high', 'strong', 'cool', 'change', 'yes']
```



[sunny , warm , high , strong , cool , change , yes]

```
In [3]: TotalExamples = len(data);
i=0;
j=0;
k=0;
print("The steps of the Find-s algorithm are :\n",hypo);
list = [];
p=0;
d=len(data[p])-1;
for j in range(d):
    list.append(data[i][j]);
hypo=list;
i=1;
for i in range(TotalExamples):
    for k in range(d):
        if hypo[k]!=data[i][k]:
            hypo[k]='?';
            k=k+1;
        else:
            hypo[k];
    print(hypo);
i=i+1;
```

The steps of the Find-s algorithm are :
 ['%', '%', '%', '%', '%', '%']
 ['sunny', 'warm', 'normal', 'strong', 'warm', 'same']
 ['sunny', 'warm', '?', 'strong', 'warm', 'same']
 ['sunny', 'warm', '?', 'strong', '?', '?']

```
In [4]: print("\nThe maximally specific Find-s hypothesis for the given training examples is :");
list=[];
for i in range(d):
    list.append(hypo[i]);
print(list);
```

The maximally specific Find-s hypothesis for the given training examples is :
 ['sunny', 'warm', '?', 'strong', '?', '?']

In []:

| | A | B | C | D | E | F | G | H |
|---|-------|----------|----------|--------|-------|----------|-------------|---|
| 1 | sky | air temp | humidity | wind | water | forecast | enjoy sport | |
| 2 | sunny | warm | normal | strong | warm | same | yes | |
| 3 | sunny | warm | high | strong | warm | same | yes | |
| 4 | rainy | cold | high | strong | warm | change | no | |
| 5 | sunny | warm | high | strong | cool | change | yes | |
| 6 | | | | | | | | |
| 7 | | | | | | | | |
| 8 | | | | | | | | |

Lab Program -2 :-

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.

Source code and output :-

```
+*In[7]:*+
```

```
[source, ipython3]
```

```
----
```

```
import numpy as np
```

```
import pandas as pd
```

```
----
```

```
+*In[10]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Loading Data from a CSV File
```

```
data = pd.DataFrame(data=pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th  
sem\ML\lab-ml\lab 2\trainingdata.csv'))
```

```
print(data)
```

```
----
```

```
+*Out[10]:*+
```

sky airtemp humidity wind water forecast enjoySport

0 Sunny Warm Normal Strong Warm Same Yes

1 Sunny Warm High Strong Warm Same Yes

2 Rainy Cold High Strong Warm Change No

3 Sunny Warm High Strong Cool Change Yes

+*In[11]:*+

[source, ipython3]

Separating concept features from Target

concepts = np.array(data.iloc[:,0:-1])

print(concepts)

+*Out[11]:*+

['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']

['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']

['Rainy' 'Cold' 'High' 'Strong' 'Warm' 'Change']

['Sunny' 'Warm' 'High' 'Strong' 'Cool' 'Change']

```
+*In[12]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Isolating target into a separate DataFrame
```

```
# copying last column to target array
```

```
target = np.array(data.iloc[:,-1])
```

```
print(target)
```

```
----
```

```
+*Out[12]:*+
```

```
----
```

```
['Yes' 'Yes' 'No' 'Yes']
```

```
----
```

```
+*In[13]:*+
```

```
[source, ipython3]
```

```
----
```

```
def learn(concepts, target):
```

```
'''
```

```
learn() function implements the learning method of the Candidate elimination algorithm.
```

```
Arguments:
```

```
    concepts - a data frame with all the features
```

target - a data frame with corresponding output values

'''

Initialise S0 with the first instance from concepts

.copy() makes sure a new list is created instead of just pointing to the same memory location

specific_h = concepts[0].copy()

print("\nInitialization of specific_h and general_h")

print(specific_h)

#h=["#" for i in range(0,5)]

#print(h)

general_h = [["?" for i in range(len(specific_h))] for i in range(len(specific_h))]

print(general_h)

The learning iterations

for i, h in enumerate(concepts):

Checking if the hypothesis has a positive target

if target[i] == "Yes":

for x in range(len(specific_h)):

Change values in S & G only if values change

if h[x] != specific_h[x]:

specific_h[x] = '?'

general_h[x][x] = '?'

Checking if the hypothesis has a positive target

```

if target[i] == "No":
    for x in range(len(specific_h)):
        # For negative hypothesis change values only in G
        if h[x] != specific_h[x]:
            general_h[x][x] = specific_h[x]
        else:
            general_h[x][x] = '?'

```

```

print("\nSteps of Candidate Elimination Algorithm",i+1)
print(specific_h)
print(general_h)

```

```

# find indices where we have empty rows, meaning those that are unchanged
indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
for i in indices:
    # remove those rows from general_h
    general_h.remove(['?', '?', '?', '?', '?', '?'])
# Return final values
return specific_h, general_h

```

```

+*In[14]:*+

```

```

[source, ipython3]

```

```

s_final, g_final = learn(concepts, target)

```

```
print("\nFinal Specific_h:", s_final, sep="\n")
```

```
print("\nFinal General_h:", g_final, sep="\n")
```

```
+*Out[14]:*+
```

Initialization of specific_h and general_h

```
['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
```

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

Steps of Candidate Elimination Algorithm 1

```
['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
```

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

Steps of Candidate Elimination Algorithm 2

```
['Sunny' 'Warm' '?' 'Strong' 'Warm' 'Same']
```

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

Steps of Candidate Elimination Algorithm 3

```
['Sunny' 'Warm' '?' 'Strong' 'Warm' 'Same']
```

```
[[ 'Sunny', '?', '?', '?', '?', '?' ], [ '?', 'Warm', '?', '?', '?', '?' ], [ '?', '?', '?', '?', '?', '?' ], [ '?', '?', '?', '?', '?',  
'?' ], [ '?', '?', '?', '?', '?', '?' ], [ '?', '?', '?', '?', '?', 'Same' ]]
```


Steps of Candidate Elimination Algorithm 4

['Sunny' 'Warm' '?' 'Strong' '?' '?']

[['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?'], ['?', '?', '?', '?', '?', '?']]

Final Specific_h:

['Sunny' 'Warm' '?' 'Strong' '?' '?']

Final General_h:

[['Sunny', '?', '?', '?', '?', '?'], ['?', 'Warm', '?', '?', '?', '?']]

+*In[]:*+

[source, ipython3]

Output screenshots :-

```
In [7]: import numpy as np
import pandas as pd
```

```
In [10]: # Loading Data from a CSV File
data = pd.DataFrame(data=pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\lab 2\trainingdata.csv'))
print(data)

   sky airtemp humidity   wind water forecast enjoySport
0  Sunny    Warm   Normal  Strong   Warm    Same       Yes
1  Sunny    Warm    High  Strong   Warm    Same       Yes
2  Rainy    Cold    High  Strong   Warm  Change       No
3  Sunny    Warm    High  Strong   Cool  Change       Yes
```

```
In [11]: # Separating concept features from Target
concepts = np.array(data.iloc[:,0:-1])
print(concepts)

[['Sunny' 'Warm' 'Normal' 'Strong' 'Warm' 'Same']
 ['Sunny' 'Warm' 'High' 'Strong' 'Warm' 'Same']
 ['Rainy' 'Cold' 'High' 'Strong' 'Warm' 'Change']
 ['Sunny' 'Warm' 'High' 'Strong' 'Cool' 'Change']]
```

```
In [12]: # Isolating target into a separate DataFrame
# copying last column to target array
target = np.array(data.iloc[:, -1])
print(target)

['Yes' 'Yes' 'No' 'Yes']
```

```
In [13]: def learn(concepts, target):

    ...
    learn() function implements the learning method of the Candidate elimination algorithm.
    Arguments:
        concepts - a data frame with all the features
        target - a data frame with corresponding output values
    ...

    # Initialise S0 with the first instance from concepts
    # .copy() makes sure a new List is created instead of just pointing to the same memory location
    specific_h = concepts[0].copy()
    print("\nInitialization of specific_h and general_h")
    print(specific_h)
    #h=["#" for i in range(0,5)]
    #print(h)

    general_h = [["?" for i in range(len(specific_h))] for i in range(len(specific_h))]
    print(general_h)
    # The Learning iterations
    for i, h in enumerate(concepts):

        # Checking if the hypothesis has a positive target
        if target[i] == "Yes":
            for x in range(len(specific_h)):

                # Change values in S & G only if values change
                if h[x] != specific_h[x]:
                    specific_h[x] = '?'
                    general_h[x][x] = '?'

        # Checking if the hypothesis has a positive target
        if target[i] == "No":
            for x in range(len(specific_h)):
                # For negative hypothesis change values only in G
                if h[x] != specific_h[x]:
                    general_h[x][x] = specific_h[x]
                else:
                    general_h[x][x] = '?'

    print("\nSteps of Candidate Elimination Algorithm",i+1)
```


Lab Program -3 :-

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.

Source code and output :-

```
+#In[1]:*+
```

```
[source, ipython3]
```

```
----
```

```
import numpy as np
```

```
import math
```

```
import csv
```

```
----
```

```
+#In[2]:*+
```

```
[source, ipython3]
```

```
----
```

```
def read_data(filename):
```

```
    with open(filename, 'r') as csvfile:
```

```
        datareader = csv.reader(csvfile, delimiter=',')
```

```
        headers = next(datareader)
```

```
        metadata = []
```

```
        traindata = []
```

```
        for name in headers:
```

```
            metadata.append(name)
```

```
for row in datareader:
    traindata.append(row)

return (metadata, traindata)
```

```
+*In[5]:*+
```

```
[source, ipython3]
```

```
class Node:
```

```
    def __init__(self, attribute):
```

```
        self.attribute = attribute
```

```
        self.children = []
```

```
        self.answer = ""
```

```
    def __str__(self):
```

```
        return self.attribute
```

```
+*In[6]:*+
```

```
[source, ipython3]
```

```
def subtables(data, col, delete):
```

```
    dict = {}
```

```
items = np.unique(data[:, col])
count = np.zeros((items.shape[0], 1), dtype=np.int32)
```

```
for x in range(items.shape[0]):
    for y in range(data.shape[0]):
        if data[y, col] == items[x]:
            count[x] += 1
```

```
for x in range(items.shape[0]):
    dict[items[x]] = np.empty((int(count[x]), data.shape[1]), dtype="|S32")
    pos = 0
    for y in range(data.shape[0]):
        if data[y, col] == items[x]:
            dict[items[x]][pos] = data[y]
            pos += 1
    if delete:
        dict[items[x]] = np.delete(dict[items[x]], col, 1)
```

```
return items, dict
```

```
----
```

```
+#ln[7]:*+
```

```
[source, ipython3]
```

```
----
```

```
def entropy(S):
```

```
items = np.unique(S)
```

```
if items.size == 1:
```

```
    return 0
```

```
counts = np.zeros((items.shape[0], 1))
```

```
sums = 0
```

```
for x in range(items.shape[0]):
```

```
    counts[x] = sum(S == items[x]) / (S.size * 1.0)
```

```
for count in counts:
```

```
    sums += -1 * count * math.log(count, 2)
```

```
return sums
```

```
----
```

```
+#ln[8]:#+
```

```
[source, ipython3]
```

```
----
```

```
def gain_ratio(data, col):
```

```
    items, dict = subtables(data, col, delete=False)
```

```
    total_size = data.shape[0]
```

```
    entropies = np.zeros((items.shape[0], 1))
```

```
    intrinsic = np.zeros((items.shape[0], 1))
```

```

for x in range(items.shape[0]):
    ratio = dict[items[x]].shape[0]/(total_size * 1.0)
    entropies[x] = ratio * entropy(dict[items[x]][:, -1])
    intrinsic[x] = ratio * math.log(ratio, 2)

```

```

total_entropy = entropy(data[:, -1])

```

```

iv = -1 * sum(intrinsic)

```

```

for x in range(entropies.shape[0]):

```

```

    total_entropy -= entropies[x]

```

```

return total_entropy / iv

```

```

----

```

```

+*ln[9]:*+

```

```

[source, ipython3]

```

```

----

```

```

def create_node(data, metadata):

```

```

    if (np.unique(data[:, -1])).shape[0] == 1:

```

```

        node = Node("")

```

```

        node.answer = np.unique(data[:, -1])[0]

```

```

        return node

```

```

gains = np.zeros((data.shape[1] - 1, 1))

```



```
for col in range(data.shape[1] - 1):  
    gains[col] = gain_ratio(data, col)
```

```
split = np.argmax(gains)
```

```
node = Node(metadata[split])  
metadata = np.delete(metadata, split, 0)
```

```
items, dict = subtables(data, split, delete=True)
```

```
for x in range(items.shape[0]):  
    child = create_node(dict[items[x]], metadata)  
    node.children.append((items[x], child))
```

```
return node
```

```
----
```

```
+*ln[10]:*+
```

```
[source, ipython3]
```

```
----
```

```
def empty(size):
```

```
    s = ""
```

```
    for x in range(size):
```

```
        s += " "
```

```
return s
```

```
def print_tree(node, level):  
    if node.answer != "":  
        print(empty(level), node.answer)  
        return  
    print(empty(level), node.attribute)  
    for value, n in node.children:  
        print(empty(level + 1), value)  
        print_tree(n, level + 2)
```

```
----
```

```
+*In[11]:*+
```

```
[source, ipython3]
```

```
----
```

```
metadata, traindata = read_data(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-  
ml\Lab 3\id3 training dataset.csv")  
data = np.array(traindata)  
node = create_node(data, metadata)  
print_tree(node, 0)
```

```
----
```

```
+*Out[11]:*+
```

```
----
```

Outlook

overcast

b'yes'

rain

Wind

b'strong'

b'no'

b'weak'

b'yes'

sunny

Humidity

b'high'

b'no'

b'normal'

b'yes'

+*ln[]:*+

[source, ipython3]

Output screenshots :-



```
In [1]: import numpy as np
import math
import csv
```

```
In [2]: def read_data(filename):
    with open(filename, 'r') as csvfile:
        datareader = csv.reader(csvfile, delimiter=',')
        headers = next(datareader)
        metadata = []
        traindata = []
        for name in headers:
            metadata.append(name)
        for row in datareader:
            traindata.append(row)

    return (metadata, traindata)
```

```
In [5]: class Node:
    def __init__(self, attribute):
        self.attribute = attribute
        self.children = []
        self.answer = ""

    def __str__(self):
        return self.attribute
```

```
In [6]: def subtables(data, col, delete):
    dict = {}
    items = np.unique(data[:, col])
    count = np.zeros((items.shape[0], 1), dtype=np.int32)

    for x in range(items.shape[0]):
        for y in range(data.shape[0]):
            if data[y, col] == items[x]:
                count[x] += 1

    for x in range(items.shape[0]):
        dict[items[x]] = np.empty((int(count[x]), data.shape[1]), dtype="<S32")
        pos = 0
        for y in range(data.shape[0]):
            if data[y, col] == items[x]:
                dict[items[x]][pos] = data[y]
                pos += 1
    if delete:
```

```
dict[items[x]] = np.delete(dict[items[x]], col, 1)

return items, dict
```

```
In [7]: def entropy(S):
        items = np.unique(S)

        if items.size == 1:
            return 0

        counts = np.zeros((items.shape[0], 1))
        sums = 0

        for x in range(items.shape[0]):
            counts[x] = sum(S == items[x]) / (S.size * 1.0)

        for count in counts:
            sums += -1 * count * math.log(count, 2)
        return sums
```

```
In [8]: def gain_ratio(data, col):
        items, dict = subtables(data, col, delete=False)

        total_size = data.shape[0]
        entropies = np.zeros((items.shape[0], 1))
        intrinsic = np.zeros((items.shape[0], 1))

        for x in range(items.shape[0]):
            ratio = dict[items[x]].shape[0]/(total_size * 1.0)
            entropies[x] = ratio * entropy(dict[items[x]][:-1])
            intrinsic[x] = ratio * math.log(ratio, 2)

        total_entropy = entropy(data[:, -1])
        iv = -1 * sum(intrinsic)

        for x in range(entropies.shape[0]):
            total_entropy -= entropies[x]

        return total_entropy / iv
```

```

In [9]: def create_node(data, metadata):
        if (np.unique(data[:, -1])).shape[0] == 1:
            node = Node("")
            node.answer = np.unique(data[:, -1])[0]
            return node

        gains = np.zeros((data.shape[1] - 1, 1))

        for col in range(data.shape[1] - 1):
            gains[col] = gain_ratio(data, col)

        split = np.argmax(gains)

        node = Node(metadata[split])
        metadata = np.delete(metadata, split, 0)

        items, dict = subtables(data, split, delete=True)

        for x in range(items.shape[0]):
            child = create_node(dict[items[x]], metadata)
            node.children.append((items[x], child))

        return node

```

jupyter Lab-3 ID3 1BM19CS159 Last Checkpoint: 6 minutes ago (autosaved)



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Trusted

Python 3 (ipykernel)

Code

```

In [10]: def empty(size):
        s = ""
        for x in range(size):
            s += " "
        return s

        def print_tree(node, level):
            if node.answer != "":
                print(empty(level), node.answer)
                return
            print(empty(level), node.attribute)
            for value, n in node.children:
                print(empty(level + 1), value)
                print_tree(n, level + 2)

```

```

In [11]: metadata, traindata = read_data(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 3\id3 training dataset.csv")
        data = np.array(traindata)
        node = create_node(data, metadata)
        print_tree(node, 0)

```

```

Outlook
  overcast
    b'yes'
  rain
    Wind
      b'strong'
      b'no'
      b'weak'
      b'yes'
    sunny
      Humidity
        b'high'
        b'no'
        b'normal'
        b'yes'

```

In []:

| | | | | | | | | |
|---------|----------|----------|----------|--------|--------|---|---|--|
| A1 | | | | | | | | |
| Outlook | | | | | | | | |
| | A | B | C | D | E | F | G | |
| 1 | Outlook | Temperat | Humidity | Wind | Answer | | | |
| 2 | sunny | hot | high | weak | no | | | |
| 3 | sunny | hot | high | strong | no | | | |
| 4 | overcast | hot | high | weak | yes | | | |
| 5 | rain | mild | high | weak | yes | | | |
| 6 | rain | cool | normal | weak | yes | | | |
| 7 | rain | cool | normal | strong | no | | | |
| 8 | overcast | cool | normal | strong | yes | | | |
| 9 | sunny | mild | high | weak | no | | | |
| 10 | sunny | cool | normal | weak | yes | | | |
| 11 | rain | mild | normal | weak | yes | | | |
| 12 | sunny | mild | normal | strong | yes | | | |
| 13 | overcast | mild | high | strong | yes | | | |
| 14 | overcast | hot | normal | weak | yes | | | |
| 15 | rain | mild | high | strong | no | | | |
| 16 | | | | | | | | |
| 17 | | | | | | | | |
| 18 | | | | | | | | |

Lab Program -4.a.-

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

Source code and output :-

```
.*In[1]:.*+
```

```
[source, ipython3]
```

```
----
```

```
# import necessary libarities
```

```
import pandas as pd
```

```
from sklearn import tree
```

```
from sklearn.preprocessing import LabelEncoder
```

```
from sklearn.naive_bayes import GaussianNB
```

```
# load data from CSV
```

```
data = pd.read_csv(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 4\Naive  
Bayesian classifier training dataset.csv")
```

```
print("The first 5 values of data is :\n",data.head())
```

```
----
```

```
.*Out[1]:.*+
```

```
----
```

```
The first 5 values of data is :
```

```
Outlook Temperature Humidity Windy PlayTennis
```


| | | | | | |
|---|----------|------|--------|-------|-----|
| 0 | Sunny | Hot | High | False | No |
| 1 | Sunny | Hot | High | True | No |
| 2 | Overcast | Hot | High | False | Yes |
| 3 | Rainy | Mild | High | False | Yes |
| 4 | Rainy | Cool | Normal | False | Yes |

+*In[2]:*+

[source, ipython3]

obtain Train data and Train output

X = data.iloc[:, :-1]

print("\nThe First 5 values of train data is\n", X.head())

+*Out[2]:*+

The First 5 values of train data is

Outlook Temperature Humidity Windy

| | | | | |
|---|----------|------|------|-------|
| 0 | Sunny | Hot | High | False |
| 1 | Sunny | Hot | High | True |
| 2 | Overcast | Hot | High | False |
| 3 | Rainy | Mild | High | False |

```
4    Rainy    Cool    Normal    False
```

```
----
```

```
+*In[3]:*+
```

```
[source, ipython3]
```

```
----
```

```
y = data.iloc[:, -1]
```

```
print("\nThe first 5 values of Train output is\n", y.head())
```

```
----
```

```
+*Out[3]:*+
```

```
----
```

```
The first 5 values of Train output is
```

```
0    No
```

```
1    No
```

```
2    Yes
```

```
3    Yes
```

```
4    Yes
```

```
Name: PlayTennis, dtype: object
```

```
----
```

```
+*In[4]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Convert then in numbers
```

```
le_outlook = LabelEncoder()
```

```
X.Outlook = le_outlook.fit_transform(X.Outlook)
```

```
le_Temperature = LabelEncoder()
```

```
X.Temperature = le_Temperature.fit_transform(X.Temperature)
```

```
le_Humidity = LabelEncoder()
```

```
X.Humidity = le_Humidity.fit_transform(X.Humidity)
```

```
le_Windy = LabelEncoder()
```

```
X.Windy = le_Windy.fit_transform(X.Windy)
```

```
print("\nNow the Train data is :\n",X.head())
```

```
----
```

```
+*Out[4]:*+
```

```
----
```

```
Now the Train data is :
```

```
Outlook Temperature Humidity Windy
```

```
0    2         1    0    0
```

```
1    2         1    0    1
```

```
2    0    1    0    0
3    1    2    0    0
4    1    0    1    0
```

```
+*In[5]:*+
```

```
[source, ipython3]
```

```
le_PlayTennis = LabelEncoder()
```

```
y = le_PlayTennis.fit_transform(y)
```

```
print("\nNow the Train output is\n",y)
```

```
+*Out[5]:*+
```

```
Now the Train output is
```

```
[0 0 1 1 1 0 1 0 1 1 1 1 1 0]
```

```
+*In[6]:*+
```

```
[source, ipython3]
```

```
from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X,y, test_size=0.20)
```

```
classifier = GaussianNB()
classifier.fit(X_train,y_train)
```

```
from sklearn.metrics import accuracy_score
print("Accuracy is:",accuracy_score(classifier.predict(X_test),y_test))
```

```
+*Out[6]:*+
```

```
Accuracy is: 0.3333333333333333
```

```
+*In[ ]:*+
```

```
[source, ipython3]
```

Output screenshots :-



```
In [1]: # import necessary libraries
import pandas as pd
from sklearn import tree
from sklearn.preprocessing import LabelEncoder
from sklearn.naive_bayes import GaussianNB

# Load data from CSV
data = pd.read_csv(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 4\Naive Bayesian classifier training dataset.csv")
print("The first 5 values of data is :\n",data.head())
```

The first 5 values of data is :

| | Outlook | Temperature | Humidity | Windy | PlayTennis |
|---|----------|-------------|----------|-------|------------|
| 0 | Sunny | Hot | High | False | No |
| 1 | Sunny | Hot | High | True | No |
| 2 | Overcast | Hot | High | False | Yes |
| 3 | Rainy | Mild | High | False | Yes |
| 4 | Rainy | Cool | Normal | False | Yes |

```
In [2]: # obtain Train data and Train output
X = data.iloc[:, :-1]
print("\nThe First 5 values of train data is\n",X.head())
```

The First 5 values of train data is

| | Outlook | Temperature | Humidity | Windy |
|---|----------|-------------|----------|-------|
| 0 | Sunny | Hot | High | False |
| 1 | Sunny | Hot | High | True |
| 2 | Overcast | Hot | High | False |
| 3 | Rainy | Mild | High | False |
| 4 | Rainy | Cool | Normal | False |

```
In [3]: y = data.iloc[:, -1]
print("\nThe first 5 values of Train output is\n",y.head())
```

The first 5 values of Train output is

| | |
|---|-----|
| 0 | No |
| 1 | No |
| 2 | Yes |
| 3 | Yes |
| 4 | Yes |

Name: PlayTennis, dtype: object



```
In [4]: # Convert then in numbers
le_outlook = LabelEncoder()
X.Outlook = le_outlook.fit_transform(X.Outlook)

le_Temperature = LabelEncoder()
X.Temperature = le_Temperature.fit_transform(X.Temperature)

le_Humidity = LabelEncoder()
X.Humidity = le_Humidity.fit_transform(X.Humidity)

le_Windy = LabelEncoder()
X.Windy = le_Windy.fit_transform(X.Windy)

print("\nNow the Train data is :\n",X.head())
```

```
Now the Train data is :
   Outlook  Temperature  Humidity  Windy
0         2             1         0      0
1         2             1         0      1
2         0             1         0      0
3         1             2         0      0
4         1             0         1      0
```

```
In [5]: le_PlayTennis = LabelEncoder()
y = le_PlayTennis.fit_transform(y)
print("\nNow the Train output is\n",y)
```

```
Now the Train output is
[0 0 1 1 1 0 1 0 1 1 1 1 0]
```

```
In [6]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X,y, test_size=0.20)

classifier = GaussianNB()
classifier.fit(X_train,y_train)

from sklearn.metrics import accuracy_score
print("Accuracy is:",accuracy_score(classifier.predict(X_test),y_test))
```

```
Accuracy is: 0.3333333333333333
```

In []:

| | | | | | | |
|----|----------|----------|----------|-------|------------|---------|
| A1 | | | | | | Outlook |
| | A | B | C | D | E | F |
| 1 | Outlook | Temperat | Humidity | Windy | PlayTennis | |
| 2 | Sunny | Hot | High | FALSE | No | |
| 3 | Sunny | Hot | High | TRUE | No | |
| 4 | Overcast | Hot | High | FALSE | Yes | |
| 5 | Rainy | Mild | High | FALSE | Yes | |
| 6 | Rainy | Cool | Normal | FALSE | Yes | |
| 7 | Rainy | Cool | Normal | TRUE | No | |
| 8 | Overcast | Cool | Normal | TRUE | Yes | |
| 9 | Sunny | Mild | High | FALSE | No | |
| 10 | Sunny | Cool | Normal | FALSE | Yes | |
| 11 | Rainy | Mild | Normal | FALSE | Yes | |
| 12 | Sunny | Mild | Normal | TRUE | Yes | |
| 13 | Overcast | Mild | High | TRUE | Yes | |
| 14 | Overcast | Hot | Normal | FALSE | Yes | |
| 15 | Rainy | Mild | High | TRUE | No | |
| 16 | | | | | | |
| 17 | | | | | | |
| 18 | | | | | | |
| 19 | | | | | | |

Lab Program -4.b.-

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 (without packages).

Source code and output :-

```
+*In[1]:*+
```

```
[source, ipython3]
```

```
----
```

```
import math
```

```
import csv
```

```
import random
```

```
----
```

```
+*In[2]:*+
```

```
[source, ipython3]
```

```
----
```

This make sures that the dataset is in an ordered format. If we have some arbitrary names in that column it difficult to deal with that.

```
def encode_class(dataset):
```

```
    classes=[]
```

```
    for i in range(len(dataset)):
```

```
        if dataset[i][-1] not in classes:
```

```
            classes.append(dataset[i][-1])
```

```
# Looping across the classes which we have derived above.This will make sure that we have definitive classes (numeric) and not arbitrary
```

```
for i in range(len(classes)):
```

```
    # Looping across all rows of dataset
```

```
    for j in range(len(dataset)):
```

```
        if dataset[j][-1] == classes[i]:
```

```
            dataset[j][-1]=i
```

```
return dataset
```

```
----
```

```
+*In[3]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Splitting the data between training set and testing set. Normally its a general understanding the training:testing=7:3
```

```
def train_test_split(dataset,ratio):
```

```
    test_num=int(ratio*len(dataset))
```

```
    train=list(dataset)
```

```
    test=[]
```

```
    for i in range(test_num):
```

```
        rand=random.randrange(len(train))
```

```
        test.append(train.pop(rand))
```

```
    return train,test
```

```
----
```

```
+*In[4]:*+
```

```
[source, ipython3]
```

```
----
```

Now depending on resultant value (last column values), we need to group the rows. It will be usefult for calculating mean and std_dev

```
def groupUnderClass(train):
```

```
    dict={}
```

```
    for row in train:
```

```
        if row[-1] not in dict:
```

```
            dict[row[-1]]=[]
```

```
            dict[row[-1]].append(row)
```

```
    return dict
```

```
----
```

```
+*In[5]:*+
```

```
[source, ipython3]
```

```
----
```

Standard formulae (just by-heart)

```
def mean(val):
```

```
    return sum(val)/float(len(val)) #Obvious
```

```
def stdDev(val):
```

```
avg=mean(val)
variance=sum([pow(x-avg,2) for x in val])/float(len(val)-1) # Especially this one
return math.sqrt(variance)
```

```
+*ln[6]:*+
```

```
[source, ipython3]
```

We will calculate the mean and std dev with respect to each attribute. Important while calculating gaussian probability

```
def meanStdDev(instances):
```

```
    info=[(mean(x),stdDev(x)) for x in zip(*instances)] # Here we are taking complete column's values of all instances.
```

```
    del info[-1]
```

```
    return info
```

```
+*ln[7]:*+
```

```
[source, ipython3]
```

As explained earlier why we need to group. We will be calculating the mean and std dev with respect each class.

```
def MeanAndStdDevForClass(train):
    info={}
    dictionary=groupUnderClass(train)
    # print(dictionary)
    for key,value in dictionary.items():
        # dictionary[key]=meanStdDev(value)
        info[key]=meanStdDev(value) #Here value stands for a complete group.
    return info
```

+*ln[8]:*+

[source, ipython3]

Its a formula by heart (no choice)

```
def calculateGaussianProbablity(x,mean,std_dev):
    expo = math.exp(-(math.pow(x - mean, 2) / (2 * math.pow(std_dev, 2))))
    return (1 / (math.sqrt(2 * math.pi) * std_dev)) * expo
```

+*ln[9]:*+

[source, ipython3]

After calculating mean and std dev w.r.t training data now its time to check if the logic will work on testing data

```
def calculateClassProbablities(info,ele):
```

```
    probablities={}
```

```
    for key,summaries in info.items(): # Info contains the groupName (key) and list of  
    (mean,std_dev) for each attribute of that group
```

```
        probablities[key]=1
```

```
        for i in range(len(summaries)): #Loop across all attributes
```

```
            mean,std_dev=summaries[i]
```

```
            x=ele[i] # Testing data's one instance's attribute value.
```

```
            probablities[key] *= calculateGaussianProbability(x, mean, std_dev)
```

```
    return probablities
```

```
    +*ln[10]:*+
```

```
    [source, ipython3]
```

```
def predict(info,ele):
```

```
    probablities=calculateClassProbablities(info,ele) # returns a dictionary of probablities for each  
    group
```

```
    bestLabel,bestProb=None,-1
```

```
    # Consider group name whichever gives you the highest probablities for this instance of  
    testing data
```

```
for key,prob in probablities.items():  
    if bestLabel==None or prob>bestProb:  
        bestProb=prob  
        bestLabel=key  
return bestLabel
```

```
+*In[11]:*+
```

```
[source, ipython3]
```

Loop across testing data and store the predicted result from our model in the list.

```
def getPredictions(info,test):
```

```
    predictions=[]
```

```
    for ele in test:
```

```
        result=predict(info,ele) # This will give you the group to which it will belong.
```

```
        predictions.append(result)
```

```
    return predictions
```

```
+*In[12]:*+
```

```
[source, ipython3]
```

```
def check_accuracy(predictions,test):  
    count=0  
    for i in range(len(test)):  
        if predictions[i]==test[i][-1]:  
            count+=1  
    return count/float(len(test))*100
```

+*ln[13]:*+

[source, ipython3]

```
filename=r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 4\pima-indians-  
diabetes.csv"  
dataset=csv.reader(open(filename))  
dataset=list(dataset)  
dataset=encode_class(dataset)  
for i in range(len(dataset)):  
    dataset[i]=[float(x) for x in dataset[i]]  
  
ratio=0.3  
print(len(dataset))  
train,test=train_test_split(dataset,ratio)  
info=MeanAndStdDevForClass(train)  
  
predictions=getPredictions(info,test)
```



```
accuracy=check_accuracy(predictions,test)
```

```
accuracy
```

```
----
```

```
+*Out[13]:*+
```

```
----
```

```
768
```

```
75.21739130434783----
```

```
+*In[ ]:*+
```

```
[source, ipython3]
```

```
----
```

```
----
```

Output screenshots :-



```
In [1]: import math
import csv
import random
```

```
In [2]: # This make sures that the dataset is in an ordered format. If we have some arbirary names in that column it difficult to deal wi

def encode_class(dataset):
    classes=[]
    for i in range(len(dataset)):
        if dataset[i][-1] not in classes:
            classes.append(dataset[i][-1])

    # Looping across the classes which we have derived above.This will make sure that we have definitive classes (numeric) and not
    for i in range(len(classes)):
        # Looping across all rows of dataset
        for j in range(len(dataset)):
            if dataset[j][-1] == classes[i]:
                dataset[j][-1]=i
    return dataset
```

```
In [3]: # Splitting the data between training set and testing set. Normally its a general understanding the training:testing=7:3
```

```
def train_test_split(dataset,ratio):
    test_num=int(ratio*len(dataset))
    train=list(dataset)
    test=[]
    for i in range(test_num):
        rand=random.randrange(len(train))
        test.append(train.pop(rand))
    return train,test
```

```
In [4]: # Now depending on resultant value (last column values), we need to group the rows. It will be usefult for calculating mean and s
```

```
def groupUnderClass(train):
    dict={}
    for row in train:
        if row[-1] not in dict:
            dict[row[-1]]=[]
        dict[row[-1]].append(row)
    return dict
```



In [5]: *# Standard formulae (just by-heart)*

```
def mean(val):  
    return sum(val)/float(len(val)) #Obvious  
  
def stdDev(val):  
    avg=mean(val)  
    variance=sum([pow(x-avg,2) for x in val])/float(len(val)-1) # Especially this one  
    return math.sqrt(variance)
```

In [6]: *# We will calculate the mean and std dev with respect to each attribute. Important while calculating gaussian probability*

```
def meanStdDev(instances):  
    info=[(mean(x),stdDev(x)) for x in zip(*instances)] # Here we are taking complete column's values of all instances.  
    del info[-1]  
    return info
```

In [7]: *# As explained earlier why we need to group. We will be calculating the mean and std dev with respect each class.*

```
def MeanAndStdDevForClass(train):  
    info={}  
    dictionary=groupUnderClass(train)  
    # print(dictionary)  
    for key,value in dictionary.items():  
        # dictionary[key]=meanStdDev(value)  
        info[key]=meanStdDev(value) #Here value stands for a complete group.  
    return info
```

In [8]: *# Its a formula by heart (no choice)*

```
def calculateGaussianProbability(x,mean,std_dev):  
    expo = math.exp(-(math.pow(x - mean, 2) / (2 * math.pow(std_dev, 2))))  
    return (1 / (math.sqrt(2 * math.pi) * std_dev)) * expo
```



File Edit View Insert Cell Kernel Widgets Help

Trusted

Python 3 (ipykernel)

Code

```
return (1 / (math.sqrt(2 * math.pi) * std_dev)) * expo
```

In [9]: *# After calculating mean and std dev w.r.t training data now its time to check if the logic will work on testing data*

```
def calculateClassProbabilities(info,ele):
    probabilities={}
    for key,summaries in info.items(): # Info contains the groupName (key) and List of (mean,std_dev) for each attribute of that group
        probabilities[key]=1
        for i in range(len(summaries)): #Loop across all attributes
            mean,std_dev=summaries[i]
            x=ele[i] # Testing data's one instance's attribute value.
            probabilities[key] *= calculateGaussianProbability(x, mean, std_dev)
    return probabilities
```

In [10]:

```
def predict(info,ele):
    probabilities=calculateClassProbabilities(info,ele) # returns a dictionary of probabilities for each group
    bestLabel,bestProb=None,-1
    # Consider group name whichever gives you the highest probabilities for this instance of testing data
    for key,prob in probabilities.items():
        if bestLabel==None or prob>bestProb:
            bestProb=prob
            bestLabel=key
    return bestLabel
```

In [11]: *# Loop across testing data and store the predicted result from our model in the list.*

```
def getPredictions(info,test):
    predictions=[]
    for ele in test:
        result=predict(info,ele) # This will give you the group to which it will belong.
        predictions.append(result)
    return predictions
```

In [12]:

```
def check_accuracy(predictions,test):
    count=0
    for i in range(len(test)):
        if predictions[i]==test[i]:
            count+=1
    return count/float(len(test))*100
```

In [11]: # Loop across testing data and store the predicted result from our model in the list.

```
def getPredictions(info,test):  
    predictions=[]  
    for ele in test:  
        result=predict(info,ele) # This will give you the group to which it will belong.  
        predictions.append(result)  
    return predictions
```

In [12]: def check_accuracy(predictions,test):

```
    count=0  
    for i in range(len(test)):  
        if predictions[i]!=test[i][-1]:  
            count+=1  
    return count/float(len(test))*100
```

In [13]: filename=r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 4\pima-indians-diabetes.csv"

```
dataset=csv.reader(open(filename))  
dataset=list(dataset)  
dataset=encode_class(dataset)  
for i in range(len(dataset)):  
    dataset[i]=[float(x) for x in dataset[i]]  
  
ratio=0.3  
print(len(dataset))  
train,test=train_test_split(dataset,ratio)  
info=MeanAndStdDevForClass(train)  
  
predictions=getPredictions(info,test)  
accuracy=check_accuracy(predictions,test)  
accuracy
```

768

Out[13]: 75.21739130434783

In []:

Lab Program -5:-

Implement the Linear Regression algorithm in order to fit data points. Select appropriate data set for your experiment and draw graphs.

Source code and output :-

```
+*In[1]:*+
```

```
[source, ipython3]
```

```
----
```

```
import numpy as np
```

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

```
import pandas as pd
```

```
----
```

```
+*In[11]:*+
```

```
[source, ipython3]
```

```
----
```

```
dataset = pd.read_csv(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 5\Lr-Salary Dataset.csv")
```

```
X = dataset.iloc[:, :-1].values
```

```
y = dataset.iloc[:, 1].values
```

```
----
```

```
+*In[13]:*+
```

```
[source, ipython3]
```

```
----
```

```
from sklearn.model_selection import train_test_split
```

```
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=1/3, random_state=0)
```

```
----
```

```
+*In[14]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Fitting Simple Linear Regression to the Training set
```

```
from sklearn.linear_model import LinearRegression
```

```
regressor = LinearRegression()
```

```
regressor.fit(X_train, y_train)
```

```
----
```

```
+*Out[14]:*+
```

```
----LinearRegression()----
```

```
+*In[15]:*+
```

```
[source, ipython3]
```

```
----
```

```
# Predicting the Test set results
```

```
y_pred = regressor.predict(X_test)
```

+*In[19]:*+

[source, ipython3]

Visualizing the Training set results

viz_train = plt

viz_train.scatter(X_train, y_train, color='red')

viz_train.plot(X_train, regressor.predict(X_train), color='blue')

viz_train.title('Salary VS Experience (Training set)')

viz_train.xlabel('Year of Experience')

viz_train.ylabel('Salary')

viz_train.show()

+*Out[19]:*+

![png](output_5_0.png)

+*In[17]:*+

[source, ipython3]

Visualizing the Test set results

viz_test = plt

viz_test.scatter(X_test, y_test, color='red')

viz_test.plot(X_train, regressor.predict(X_train), color='blue')

viz_test.title('Salary VS Experience (Test set)')

viz_test.xlabel('Year of Experience')

viz_test.ylabel('Salary')

viz_test.show()

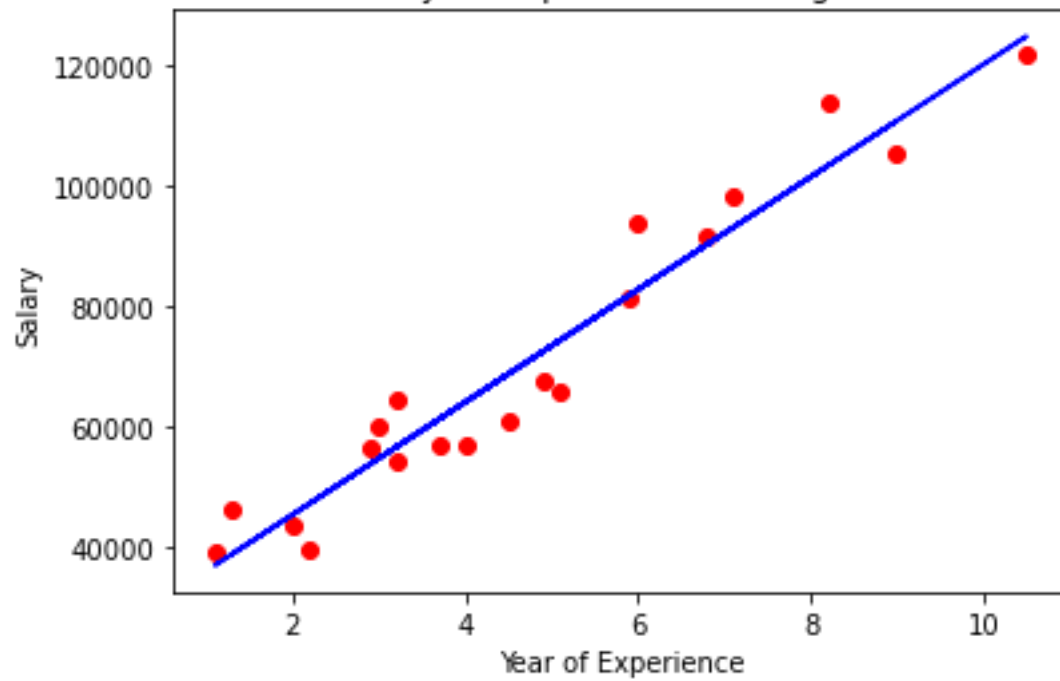
+*Out[17]:*+

![png](output_6_0.png)

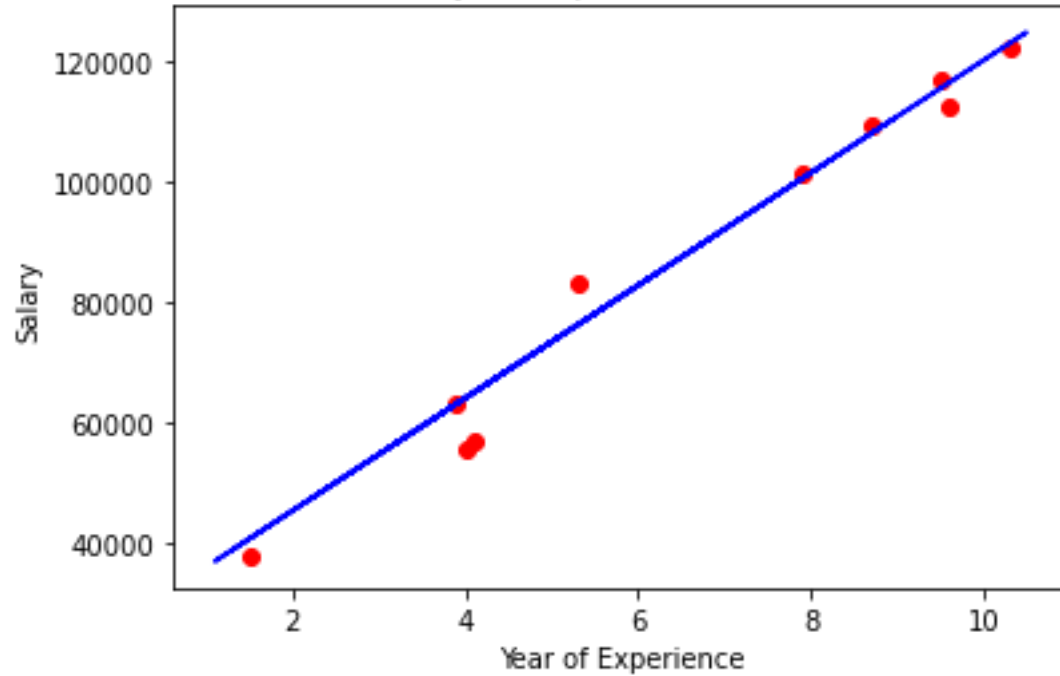
+*In []:*+

[source, ipython3]


Salary VS Experience (Training set)



Salary VS Experience (Test set)



Output screenshots :-

jupyter Lab-5 Linear regression 1BM19CS159 Last Checkpoint: 4 minutes ago (autosaved)  Logout

File Edit View Insert Cell Kernel Widgets Help Not Trusted Python 3 (ipykernel)

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
import pandas as pd


In [11]: dataset = pd.read_csv(r"C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 5\Lr-Salary Dataset.csv")
X = dataset.iloc[:, :-1].values
y = dataset.iloc[:, 1].values

In [13]: from sklearn.model_selection import train_test_split
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=1/3, random_state=0)

In [14]: # Fitting Simple Linear Regression to the Training set
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)

Out[14]: LinearRegression()


In [15]: # Predicting the Test set results
y_pred = regressor.predict(X_test)
```

jupyter Lab-5 Linear regression 1BM19CS159 Last Checkpoint: 4 minutes ago (autosaved)  Logout

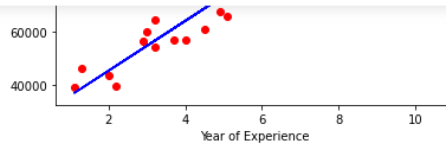
File Edit View Insert Cell Kernel Widgets Help Not Trusted Python 3 (ipykernel)

```
y_pred = regressor.predict(X_test)

In [19]: # Visualizing the Training set results
viz_train = plt
viz_train.scatter(X_train, y_train, color='red')
viz_train.plot(X_train, regressor.predict(X_train), color='blue')
viz_train.title('Salary VS Experience (Training set)')
viz_train.xlabel('Year of Experience')
viz_train.ylabel('Salary')
viz_train.show()
```



| Year of Experience | Salary |
|--------------------|--------|
| 1 | 40000 |
| 2 | 45000 |
| 3 | 55000 |
| 4 | 60000 |
| 5 | 65000 |
| 6 | 80000 |
| 7 | 90000 |
| 8 | 100000 |
| 9 | 105000 |
| 10 | 115000 |



```
In [17]: # Visualizing the Test set results
viz_test = plt
viz_test.scatter(X_test, y_test, color='red')
viz_test.plot(X_train, regressor.predict(X_train), color='blue')
viz_test.title('Salary VS Experience (Test set)')
viz_test.xlabel('Year of Experience')
viz_test.ylabel('Salary')
viz_test.show()
```



In []:

| format. | | | | |
|-------------------|-----------------|--------|---|-------|
| A1 | : | ✕ | ✓ | f_x |
| | | | | Year |
| | A | B | C | |
| 1 | YearsExperience | Salary | | |
| 2 | 1.1 | 39343 | | |
| 3 | 1.3 | 46205 | | |
| 4 | 1.5 | 37731 | | |
| 5 | 2 | 43525 | | |
| 6 | 2.2 | 39891 | | |
| 7 | 2.9 | 56642 | | |
| 8 | 3 | 60150 | | |
| 9 | 3.2 | 54445 | | |
| 10 | 3.2 | 64445 | | |
| 11 | 3.7 | 57189 | | |
| 12 | 3.9 | 63218 | | |
| 13 | 4 | 55794 | | |
| 14 | 4 | 56957 | | |
| 15 | 4.1 | 57081 | | |
| 16 | 4.5 | 61111 | | |
| 17 | 4.9 | 67938 | | |
| 18 | 5.1 | 66029 | | |
| 19 | 5.3 | 83088 | | |
| 20 | 5.9 | 81363 | | |
| 21 | 6 | 93940 | | |
| 22 | 6.8 | 91738 | | |
| 23 | 7.1 | 98273 | | |
| 24 | 7.9 | 101302 | | |
| 25 | 8.2 | 113812 | | |
| Lr-Salary Dataset | | | | |

Lab Program -6 :-

Write a program to construct a Bayesian network considering training data. Use this model to make predictions.

Source code and output :-

```
 -*-In[1]:*+
```

```
[source, ipython3]
```

```
----
```

```
!pip install pgmpy
```

```
----
```

```
 -*-Out[1]:*+
```

```
----
```

Defaulting to user installation because normal site-packages is not writeable

Collecting pgmpy

Downloading pgmpy-0.1.18-py3-none-any.whl (1.9 MB)

Requirement already satisfied: scipy in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.7.3)

Requirement already satisfied: pyparsing in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (3.0.4)

Requirement already satisfied: pandas in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.4.2)

Collecting torch

Downloading torch-1.11.0-cp39-cp39-win_amd64.whl (157.9 MB)

Requirement already satisfied: scikit-learn in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.0.2)

Requirement already satisfied: numpy in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.21.5)

Requirement already satisfied: tqdm in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (4.64.0)

Requirement already satisfied: networkx in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (2.7.1)

Requirement already satisfied: joblib in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.1.0)

Requirement already satisfied: statsmodels in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (0.13.2)

Requirement already satisfied: python-dateutil>=2.8.1 in c:\programdata\anaconda3\lib\site-packages (from pandas->pgmpy) (2.8.2)

Requirement already satisfied: pytz>=2020.1 in c:\programdata\anaconda3\lib\site-packages (from pandas->pgmpy) (2021.3)

Requirement already satisfied: six>=1.5 in c:\programdata\anaconda3\lib\site-packages (from python-dateutil>=2.8.1->pandas->pgmpy) (1.16.0)

Requirement already satisfied: threadpoolctl>=2.0.0 in c:\programdata\anaconda3\lib\site-packages (from scikit-learn->pgmpy) (2.2.0)

Requirement already satisfied: patsy>=0.5.2 in c:\programdata\anaconda3\lib\site-packages (from statsmodels->pgmpy) (0.5.2)

Requirement already satisfied: packaging>=21.3 in c:\programdata\anaconda3\lib\site-packages (from statsmodels->pgmpy) (21.3)

Requirement already satisfied: typing-extensions in c:\programdata\anaconda3\lib\site-packages (from torch->pgmpy) (4.1.1)

Requirement already satisfied: colorama in c:\programdata\anaconda3\lib\site-packages (from tqdm->pgmpy) (0.4.4)

Installing collected packages: torch, pgmpy

Successfully installed pgmpy-0.1.18 torch-1.11.0

WARNING: The scripts convert-caffe2-to-onnx.exe, convert-onnx-to-caffe2.exe and torchrun.exe are installed in 'C:\Users\Admin\AppData\Roaming\Python\Python39\Scripts' which is not on PATH.

Consider adding this directory to PATH or, if you prefer to suppress this warning, use --no-warn-script-location.

```
+*In[1]:*+
```

```
[source, ipython3]
```

```
import numpy as np
```

```
import pandas as pd
```

```
import csv
```

```
import pgmpy
```



```

from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination

----

+*In[6]:*+
[source, ipython3]
----

#read Cleveland Heart Disease data
heartDisease = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 6\heart.csv')
heartDisease = heartDisease.replace('?',np.nan)

----

+*In[7]:*+
[source, ipython3]
----

#display the data
print('Sample instances from the dataset are given below')
print(heartDisease.head())

----

+*Out[7]:*+
----

Sample instances from the dataset are given below

   Unnamed: 0  age  sex  cp  trestbps  chol  fbs  restecg  thalach  exang \
0      NaN  63.0  1.0  1.0   145.0  233.0  1.0    2.0   150.0   0.0

```

| | | | | | | | | | | |
|---|-----|------|-----|-----|-------|-------|-----|-----|-------|-----|
| 1 | NaN | 67.0 | 1.0 | 4.0 | 160.0 | 286.0 | 0.0 | 2.0 | 108.0 | 1.0 |
| 2 | NaN | 67.0 | 1.0 | 4.0 | 120.0 | 229.0 | 0.0 | 2.0 | 129.0 | 1.0 |
| 3 | NaN | 37.0 | 1.0 | 3.0 | 130.0 | 250.0 | 0.0 | 0.0 | 187.0 | 0.0 |
| 4 | NaN | 41.0 | 0.0 | 2.0 | 130.0 | 204.0 | 0.0 | 2.0 | 172.0 | 0.0 |

... slope ca thal heartdisease Unnamed: 15 Unnamed: 16 Unnamed: 17 \

| | | | | | | | |
|-------|-----|---|---|-----|-----|-----|-----|
| 0 ... | 3.0 | 0 | 6 | 0.0 | NaN | NaN | NaN |
| 1 ... | 2.0 | 3 | 3 | 2.0 | NaN | NaN | NaN |
| 2 ... | 2.0 | 2 | 7 | 1.0 | NaN | NaN | NaN |
| 3 ... | 3.0 | 0 | 3 | 0.0 | NaN | NaN | NaN |
| 4 ... | 1.0 | 0 | 3 | 0.0 | NaN | NaN | NaN |

Unnamed: 18 Unnamed: 19 Unnamed: 20

| | | | |
|---|-----|-----|-----|
| 0 | NaN | NaN | NaN |
| 1 | NaN | NaN | NaN |
| 2 | NaN | NaN | NaN |
| 3 | NaN | NaN | NaN |
| 4 | NaN | NaN | NaN |

[5 rows x 21 columns]

.*In[8]:.*

[source, ipython3]

#display the Attributes names and datatypes

print('\n Attributes and datatypes')

print(heartDisease.dtypes)

+*Out[8]:*+

Attributes and datatypes

Unnamed: 0 float64

age float64

sex float64

cp float64

trestbps float64

chol float64

fbs float64

restecg float64

thalach float64

exang float64

oldpeak float64

slope float64

ca object

thal object

heartdisease float64

Unnamed: 15 float64

Unnamed: 16 float64

Unnamed: 17 float64

Unnamed: 18 float64

Unnamed: 19 float64

Unnamed: 20 float64

dtype: object

+*In[9]:*+

[source, ipython3]

#Creat Model-Bayesian Network

model =

BayesianModel([('age','heartdisease'),('sex','heartdisease'),('exang','heartdisease'),('cp','heartdisease'),('heartdisease','restecg'),('heartdisease','chol')])

+*Out[9]:*+

C:\Users\Admin\AppData\Roaming\Python\Python39\site-packages\pgmpy\models\BayesianModel.py:8:
FutureWarning: BayesianModel has been renamed to BayesianNetwork. Please use BayesianNetwork class,
BayesianModel will be removed in future.

warnings.warn(

+*In[10]:*+

[source, ipython3]

#Learning CPDs using Maximum Likelihood Estimators

print('\n Learning CPD using Maximum likelihood estimators')

model.fit(heartDisease,estimator=MaximumLikelihoodEstimator)

```
+*Out[10]:*+
```

```
----
```

```
Learning CPD using Maximum likelihood estimators
```

```
----
```

```
+*In[11]:*+
```

```
[source, ipython3]
```

```
----
```

```
#Inferencing with Bayesian Network
```

```
print('\n Inferencing with Bayesian Network:')
```

```
HeartDiseasetest_infer = VariableElimination(model)
```

```
----
```

```
+*Out[11]:*+
```

```
----
```

```
Inferencing with Bayesian Network:
```

```
----
```

```
+*In[12]:*+
```

```
[source, ipython3]
```

```
----
```

```
#computing the Probability of HeartDisease given restecg
```

```
print('\n 1.Probability of HeartDisease given evidence= restecg :1')
```

```
q1=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'restecg':1})
```

```
print(q1)
```

```
----
```

```
+*Out[12]:*+
```

```
----
```

1.Probability of HeartDisease given evidence= restecg :1

0%| | 0/4 [00:00<?, ?it/s] 0%| | 0/4 [00:00<?, ?it/s]

```
+-----+-----+
```

```
| heartdisease | phi(heartdisease) |
```

```
+=====+
```

```
| heartdisease(0.0) |      0.2000 |
```

```
+-----+-----+
```

```
| heartdisease(1.0) |      0.2000 |
```

```
+-----+-----+
```

```
| heartdisease(2.0) |      0.2000 |
```

```
+-----+-----+
```

```
| heartdisease(3.0) |      0.2000 |
```

```
+-----+-----+
```

```
| heartdisease(4.0) |      0.2000 |
```

```
+-----+-----+
```

```
----
```

```
+*In[14]:*+
```

```
[source, ipython3]
```

```
----
```

```
#computing the Probability of HeartDisease given cp
print('\n 2.Probability of HeartDisease given evidence= cp:2 ')
q2=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

```
----
```

```
+*Out[14]:*+
```

```
----
```


```
2.Probability of HeartDisease given evidence= cp:2
0%|      | 0/3 [00:00<?, ?it/s] 0%|      | 0/3 [00:00<?, ?it/s]
```

```
+-----+-----+
| heartdisease | phi(heartdisease) |
+=====+=====+
| heartdisease(0.0) |      0.2000 |
+-----+-----+
| heartdisease(1.0) |      0.2000 |
+-----+-----+
| heartdisease(2.0) |      0.2000 |
+-----+-----+
| heartdisease(3.0) |      0.2000 |
+-----+-----+
| heartdisease(4.0) |      0.2000 |
+-----+-----+
```

```
----
```


```
+*In[ ]:*+
```

[source, ipython3]

 jupyter









Lab-6 Bayesian Network (heart dataset) 1BM19CS159

Last Checkpoint: 6 minutes ago (autosaved)

 Logout

FileEditViewInsertCellKernelWidgetsHelp

TrustedPython 3 (ipykernel)



Code

```
In [1]: !pip install pgmpy

Defaulting to user installation because normal site-packages is not writeable
Collecting pgmpy
  Downloading pgmpy-0.1.18-py3-none-any.whl (1.9 MB)
Requirement already satisfied: scipy in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.7.3)
Requirement already satisfied: pyparsing in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (3.0.4)
Requirement already satisfied: pandas in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.4.2)
Collecting torch
  Downloading torch-1.11.0-cp39-cp39-win_amd64.whl (157.9 MB)
Requirement already satisfied: scikit-learn in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.0.2)
Requirement already satisfied: numpy in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.21.5)
Requirement already satisfied: tqdm in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (4.64.0)
Requirement already satisfied: networkx in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (2.7.1)
Requirement already satisfied: joblib in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (1.1.0)
Requirement already satisfied: statsmodels in c:\programdata\anaconda3\lib\site-packages (from pgmpy) (0.13.2)
Requirement already satisfied: python-dateutil>=2.8.1 in c:\programdata\anaconda3\lib\site-packages (from pandas->pgmpy) (2.8.2)
Requirement already satisfied: pytz>=2020.1 in c:\programdata\anaconda3\lib\site-packages (from pandas->pgmpy) (2021.3)
Requirement already satisfied: six>=1.5 in c:\programdata\anaconda3\lib\site-packages (from python-dateutil>=2.8.1->pandas->pgmpy) (1.16.0)
Requirement already satisfied: threadpoolctl>=2.0.0 in c:\programdata\anaconda3\lib\site-packages (from scikit-learn->pgmpy) (2.2.0)
Requirement already satisfied: patsy>=0.5.2 in c:\programdata\anaconda3\lib\site-packages (from statsmodels->pgmpy) (0.5.2)
Requirement already satisfied: packaging>=21.3 in c:\programdata\anaconda3\lib\site-packages (from statsmodels->pgmpy) (21.3)
Requirement already satisfied: typing-extensions in c:\programdata\anaconda3\lib\site-packages (from torch->pgmpy) (4.1.1)
Requirement already satisfied: colorama in c:\programdata\anaconda3\lib\site-packages (from tqdm->pgmpy) (0.4.4)
Installing collected packages: torch, pgmpy
Successfully installed pgmpy-0.1.18 torch-1.11.0

WARNING: The scripts convert-caffe2-to-onnx.exe, convert-onnx-to-caffe2.exe and torchrun.exe are installed in 'C:\Users\Admin\AppData\Roaming\Python\Python39\Scripts' which is not on PATH.
Consider adding this directory to PATH or, if you prefer to suppress this warning, use --no-warn-script-location.
```

```
In [1]: import numpy as np
import pandas as pd
import csv
import pgmpy
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
```




```
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
```

```
In [6]: #read Cleveland Heart Disease data
heartDisease = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 6\heart.csv')
heartDisease = heartDisease.replace('?', np.nan)
```

```
In [7]: #display the data
print('Sample instances from the dataset are given below')
print(heartDisease.head())
```

```
Sample instances from the dataset are given below
   Unnamed: 0  age  sex  cp  trestbps  chol  fbs  restecg  thalach  exang  \
0         NaN  63.0  1.0  1.0    145.0  233.0  1.0     2.0    150.0   0.0
1         NaN  67.0  1.0  4.0    160.0  286.0  0.0     2.0    108.0   1.0
2         NaN  67.0  1.0  4.0    120.0  229.0  0.0     2.0    129.0   1.0
3         NaN  37.0  1.0  3.0    130.0  250.0  0.0     0.0    187.0   0.0
4         NaN  41.0  0.0  2.0    130.0  204.0  0.0     2.0    172.0   0.0

   ...  slope  ca  thal  heartdisease  Unnamed: 15  Unnamed: 16  Unnamed: 17  \
0  ...    3.0  0   6             0.0          NaN          NaN          NaN
1  ...    2.0  3   3             2.0          NaN          NaN          NaN
2  ...    2.0  2   7             1.0          NaN          NaN          NaN
3  ...    3.0  0   3             0.0          NaN          NaN          NaN
4  ...    1.0  0   3             0.0          NaN          NaN          NaN

   Unnamed: 18  Unnamed: 19  Unnamed: 20
0         NaN          NaN          NaN
1         NaN          NaN          NaN
2         NaN          NaN          NaN
3         NaN          NaN          NaN
4         NaN          NaN          NaN
```

[5 rows x 21 columns]

```
In [8]: #display the Attributes names and datatypes
print('\n Attributes and datatypes')
print(heartDisease.dtypes)
```

```
Attributes and datatypes
Unnamed: 0      float64
age            float64
sex            float64
cp            float64
trestbps      float64
chol          float64
fbs           float64
restecg       float64
thalach       float64
exang         float64
oldpeak       float64
slope         float64
ca            object
thal          object
heartdisease   float64
Unnamed: 15    float64
Unnamed: 16    float64
Unnamed: 17    float64
Unnamed: 18    float64
Unnamed: 19    float64
Unnamed: 20    float64
dtype: object
```

```
In [9]: #Creat Model-Bayesian Network
model = BayesianModel([('age', 'heartdisease'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'heartdisease'), ('heartdisease', 'restecg')])

C:\Users\Admin\AppData\Roaming\Python\Python39\site-packages\pgmpy\models\BayesianModel.py:8: FutureWarning: BayesianModel has
been renamed to BayesianNetwork. Please use BayesianNetwork class, BayesianModel will be removed in future.
warnings.warn()
```

```
In [10]: #Learning CPDs using Maximum Likelihood Estimators
print('\n Learning CPD using Maximum likelihood estimators')
model.fit(heartDisease, estimator=MaximumLikelihoodEstimator)
```

Learning CPD using Maximum likelihood estimators

```
In [11]: #Inferencing with Bayesian Network
print('\n Inferencing with Bayesian Network:')
HeartDiseaseTest_infer = VariableElimination(model)
```

Inferencing with Bayesian Network:

```
In [12]: #computing the Probability of HeartDisease given restecg
print('\n 1.Probability of HeartDisease given evidence= restecg :1')
q1=HeartDiseaseTest_infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)
```

1.Probability of HeartDisease given evidence= restecg :1

Finding Elimination Order: : 100% 4/4 [00:00<00:00, 142.85it/s]

Eliminating: exang: 100% 4/4 [00:00<00:00, 129.02it/s]

| heartdisease | phi(heartdisease) |
|-------------------|-------------------|
| heartdisease(0.0) | 0.2000 |
| heartdisease(1.0) | 0.2000 |
| heartdisease(2.0) | 0.2000 |
| heartdisease(3.0) | 0.2000 |
| heartdisease(4.0) | 0.2000 |

```
In [13]: #computing the Probability of HeartDisease given exang
```

```
In [14]: #computing the Probability of HeartDisease given cp
print('\n 2.Probability of HeartDisease given evidence= cp:2 ')
q2=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

2.Probability of HeartDisease given evidence= cp:2

Finding Elimination Order: : 0% 0/3 [00:00<?, ?it/s]

Eliminating: exang: 100% 3/3 [00:00<00:00, 157.74it/s]

```
+-----+-----+
| heartdisease | phi(heartdisease) |
+-----+-----+
| heartdisease(0.0) | 0.2000 |
+-----+-----+
| heartdisease(1.0) | 0.2000 |
+-----+-----+
| heartdisease(2.0) | 0.2000 |
+-----+-----+
| heartdisease(3.0) | 0.2000 |
+-----+-----+
| heartdisease(4.0) | 0.2000 |
+-----+-----+
```

In []:

| A | B | C | D | E | F | G | H | I | J | K | L | M |
|---|-----|-----|----|----------|------|-----|---------|---------|-------|---------|-------|----|
| | age | sex | cp | trestbps | chol | fbs | restecg | thalach | exang | oldpeak | slope | ca |
| | 63 | 1 | 1 | 145 | 233 | 1 | 2 | 150 | 0 | 2.3 | 3 | |
| | 67 | 1 | 4 | 160 | 286 | 0 | 2 | 108 | 1 | 1.5 | 2 | |
| | 67 | 1 | 4 | 120 | 229 | 0 | 2 | 129 | 1 | 2.6 | 2 | |
| | 37 | 1 | 3 | 130 | 250 | 0 | 0 | 187 | 0 | 3.5 | 3 | |
| | 41 | 0 | 2 | 130 | 204 | 0 | 2 | 172 | 0 | 1.4 | 1 | |
| | 56 | 1 | 2 | 120 | 236 | 0 | 0 | 178 | 0 | 0.8 | 1 | |
| | 62 | 0 | 4 | 140 | 268 | 0 | 2 | 160 | 0 | 3.6 | 3 | |
| | 57 | 0 | 4 | 120 | 354 | 0 | 0 | 163 | 1 | 0.6 | 1 | |
| | 63 | 1 | 4 | 130 | 254 | 0 | 2 | 147 | 0 | 1.4 | 2 | |
| | 53 | 1 | 4 | 140 | 203 | 1 | 2 | 155 | 1 | 3.1 | 3 | |
| | 57 | 1 | 4 | 140 | 192 | 0 | 0 | 148 | 0 | 0.4 | 2 | |
| | 56 | 0 | 2 | 140 | 294 | 0 | 2 | 153 | 0 | 1.3 | 2 | |
| | 56 | 1 | 3 | 130 | 256 | 1 | 2 | 142 | 1 | 0.6 | 2 | |
| | 44 | 1 | 2 | 120 | 263 | 0 | 0 | 173 | 0 | 0 | 1 | |
| | 52 | 1 | 3 | 172 | 199 | 1 | 0 | 162 | 0 | 0.5 | 1 | |
| | 57 | 1 | 3 | 150 | 168 | 0 | 0 | 174 | 0 | 1.6 | 1 | |
| | 48 | 1 | 2 | 110 | 229 | 0 | 0 | 168 | 0 | 1 | 3 | |
| | 54 | 1 | 4 | 140 | 239 | 0 | 0 | 160 | 0 | 1.2 | 1 | |
| | 48 | 0 | 3 | 130 | 275 | 0 | 0 | 139 | 0 | 0.2 | 1 | |
| | 49 | 1 | 2 | 130 | 266 | 0 | 0 | 171 | 0 | 0.6 | 1 | |
| | 64 | 1 | 1 | 110 | 211 | 0 | 2 | 144 | 1 | 1.8 | 2 | |
| | 58 | 0 | 1 | 150 | 283 | 1 | 2 | 162 | 0 | 1 | 1 | |
| | 58 | 1 | 2 | 120 | 284 | 0 | 2 | 160 | 0 | 1.8 | 2 | |

Lab Program -7 :-

Apply k-Means algorithm to cluster a set of data stored in a .CSV file.

Source code and output :-

```
#!/usr/bin/env python
```

```
# coding: utf-8
```

```
# In[18]:
```

```
import pandas as pd
```

```
import matplotlib
```

```
import numpy as np
```

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

```
from sklearn.cluster import KMeans
```

```
from sklearn.preprocessing import MinMaxScaler
```

```
from matplotlib import pyplot as plt
```

```
get_ipython().run_line_magic('matplotlib', 'inline')
```

```
# In[19]:
```

```
df = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 7\income.csv')
```

```
df.head(10)
```

```
# In[20]:
```

```
scaler = MinMaxScaler()
scaler.fit(df[['Age']])
df[['Age']] = scaler.transform(df[['Age']])

scaler.fit(df[['Income($)']])
df[['Income($)']] = scaler.transform(df[['Income($)']])
df.head(10)
```

```
# In[21]:
```

```
plt.scatter(df['Age'], df['Income($)'])
```

```
# In[22]:
```

```
k_range = range(1, 11)
sse = []
for k in k_range:
    kmc = KMeans(n_clusters=k)
    kmc.fit(df[['Age', 'Income($)']])
    sse.append(kmc.inertia_)
sse
```

```
# In[23]:
```

```
plt.xlabel = 'Number of Clusters'  
plt.ylabel = 'Sum of Squared Errors'  
plt.plot(k_range, sse)
```

Therefore, the elbow point is 3

```
# In[24]:
```

```
km = KMeans(n_clusters=3)  
km
```

```
# In[25]:
```

```
y_predict = km.fit_predict(df[['Age', 'Income($)']])  
y_predict
```

```
# In[26]:
```

```
df['cluster'] = y_predict  
df.head()
```

```
# In[27]:
```

```
df0 = df[df.cluster == 0]
```

```
df0
```

```
# In[28]:
```

```
df1 = df[df.cluster == 1]
```

```
df1
```

```
# In[29]:
```

```
df2 = df[df.cluster == 2]
```

```
df2
```

```
# In[30]:
```

```
km.cluster_centers_
```

```
# In[34]:
```

```

p1 = plt.scatter(df0['Age'], df0['Income($)', marker='+', color='red')
p2 = plt.scatter(df1['Age'], df1['Income($)', marker='*', color='blue')
p3 = plt.scatter(df2['Age'], df2['Income($)', marker='^', color='green')
c = plt.scatter(km.cluster_centers[:,0], km.cluster_centers[:,1], color='black')
plt.legend((p1, p2, p3, c),
           ('Cluster 1', 'Cluster 2', 'Cluster 3', 'Centroid'))

```

| | A | B | C | D |
|----|----------|-----|------------|---|
| 1 | Name | Age | Income(\$) | |
| 2 | Rob | 27 | 70000 | |
| 3 | Michael | 29 | 90000 | |
| 4 | Mohan | 29 | 61000 | |
| 5 | Ismail | 28 | 60000 | |
| 6 | Kory | 42 | 150000 | |
| 7 | Gautam | 39 | 155000 | |
| 8 | David | 41 | 160000 | |
| 9 | Andrea | 38 | 162000 | |
| 10 | Brad | 36 | 156000 | |
| 11 | Angelina | 35 | 130000 | |
| 12 | Donald | 37 | 137000 | |
| 13 | Tom | 26 | 45000 | |
| 14 | Arnold | 27 | 48000 | |
| 15 | Jared | 28 | 51000 | |
| 16 | Stark | 29 | 49500 | |
| 17 | Ranbir | 32 | 53000 | |
| 18 | Dipika | 40 | 65000 | |
| 19 | Priyanka | 41 | 63000 | |
| 20 | Nick | 43 | 64000 | |
| 21 | Alia | 39 | 80000 | |
| 22 | Sid | 41 | 82000 | |
| 23 | Abdul | 39 | 58000 | |

In[]:


```
In [18]: import pandas as pd
import matplotlib
import numpy as np
import matplotlib.pyplot as plt
from sklearn.cluster import KMeans
from sklearn.preprocessing import MinMaxScaler
from matplotlib import pyplot as plt
%matplotlib inline
```

```
In [19]: df = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 7\income.csv')
df.head(10)
```

Out[19]:

| | Name | Age | Income(\$) |
|---|----------|-----|------------|
| 0 | Rob | 27 | 70000 |
| 1 | Michael | 29 | 90000 |
| 2 | Mohan | 29 | 61000 |
| 3 | Ismail | 28 | 60000 |
| 4 | Kory | 42 | 150000 |
| 5 | Gautam | 39 | 155000 |
| 6 | David | 41 | 160000 |
| 7 | Andrea | 38 | 162000 |
| 8 | Brad | 36 | 156000 |
| 9 | Angelina | 35 | 130000 |

```
In [20]: scaler = MinMaxScaler()
scaler.fit(df[['Age']])
df[['Age']] = scaler.transform(df[['Age']])

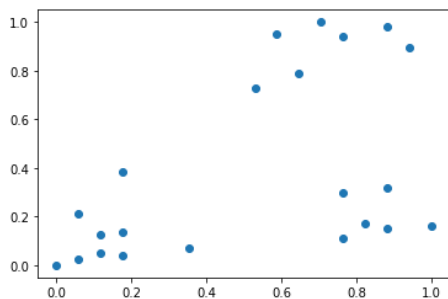
scaler.fit(df[['Income($)']])
df[['Income($)']] = scaler.transform(df[['Income($)']])
df.head(10)
```

Out[20]:

| | Name | Age | Income(\$) |
|---|---------|----------|------------|
| 0 | Rob | 0.058824 | 0.213675 |
| 1 | Michael | 0.176471 | 0.384615 |

```
In [21]: plt.scatter(df['Age'], df['Income($)'])
```

```
Out[21]: <matplotlib.collections.PathCollection at 0x298b0f99760>
```



```
In [22]: k_range = range(1, 11)
sse = []
for k in k_range:
    kmc = KMeans(n_clusters=k)
    kmc.fit(df[['Age', 'Income($)']])
    sse.append(kmc.inertia_)
sse
```

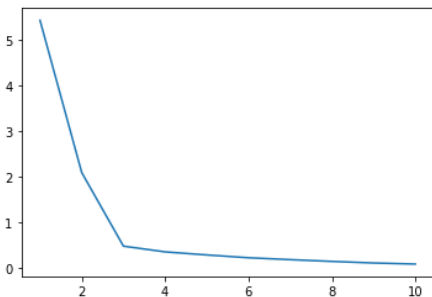
C:\ProgramData\Anaconda3\lib\site-packages\sklearn\cluster_kmeans.py:1036: UserWarning: KMeans is known to have a memory leak on Windows with MKL, when there are less chunks than available threads. You can avoid it by setting the environment variable OMP_NUM_THREADS=1.
warnings.warn(

```
Out[22]: [5.434011511988178,
2.0911363886990775,
0.4750783498553096,
0.34910470944195654,
0.2818479744366238,
0.22020960864009398,
0.17840674931327938,
0.1397684499538816,
0.1049748860620909,
0.08139933135681814]
```

```
In [23]: plt.xlabel = 'Number of Clusters'
```

```
In [23]: plt.xlabel = 'Number of Clusters'
plt.ylabel = 'Sum of Squared Errors'
plt.plot(k_range, sse)
```

```
Out[23]: <matplotlib.lines.Line2D at 0x298b1fc7eb0>
```



Therefore, the elbow point is 3

```
In [24]: km = KMeans(n_clusters=3)
km
```

```
Out[24]: KMeans(n_clusters=3)
```

```
In [25]: y_predict = km.fit_predict(df[['Age', 'Income($)']])
y_predict
```

```
Out[25]: array([0, 0, 0, 0, 1, 1, 1, 1, 1, 1, 1, 0, 0, 0, 0, 0, 2, 2, 2, 2, 2])
```

```
In [26]: df['cluster'] = y_predict
df.head()
```

```
In [28]: df1 = df[df.cluster == 1]
df1
```

```
Out[28]:
```

| | Name | Age | Income(\$) | cluster |
|----|----------|----------|------------|---------|
| 4 | Kory | 0.941176 | 0.897436 | 1 |
| 5 | Gautam | 0.764706 | 0.940171 | 1 |
| 6 | David | 0.882353 | 0.982906 | 1 |
| 7 | Andrea | 0.705882 | 1.000000 | 1 |
| 8 | Brad | 0.588235 | 0.948718 | 1 |
| 9 | Angelina | 0.529412 | 0.726496 | 1 |
| 10 | Donald | 0.647059 | 0.786325 | 1 |

```
In [29]: df2 = df[df.cluster == 2]
df2
```

```
Out[29]:
```

| | Name | Age | Income(\$) | cluster |
|----|----------|----------|------------|---------|
| 16 | Dipika | 0.823529 | 0.170940 | 2 |
| 17 | Priyanka | 0.882353 | 0.153846 | 2 |
| 18 | Nick | 1.000000 | 0.162393 | 2 |
| 19 | Alia | 0.764706 | 0.299145 | 2 |
| 20 | Sid | 0.882353 | 0.316239 | 2 |
| 21 | Abdul | 0.764706 | 0.111111 | 2 |

```
In [30]: km.cluster_centers_
```

```
Out[30]: array([[0.1372549 , 0.11633428],
 [0.72268908, 0.8974359 ],
 [0.85294118, 0.2022792 ]])
```

```
In [34]: p1 = plt.scatter(df0['Age'], df0['Income($)', marker='+', color='red')
p2 = plt.scatter(df1['Age'], df1['Income($)', marker='*', color='blue')
p3 = plt.scatter(df2['Age'], df2['Income($)', marker='^', color='green')
c = plt.scatter(km.cluster_centers_[0], km.cluster_centers_[1], color='black')
```

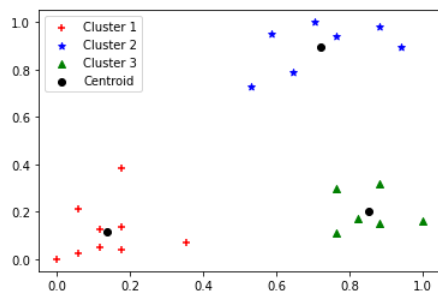
```
21 Abdul 0.764706 0.111111 2
```

```
In [30]: km.cluster_centers_
```

```
Out[30]: array([[0.1372549 , 0.11633428],
 [0.72268908, 0.8974359 ],
 [0.85294118, 0.2022792 ]])
```

```
In [34]: p1 = plt.scatter(df0['Age'], df0['Income($)', marker='+', color='red')
p2 = plt.scatter(df1['Age'], df1['Income($)', marker='*', color='blue')
p3 = plt.scatter(df2['Age'], df2['Income($)', marker='^', color='green')
c = plt.scatter(km.cluster_centers_[0], km.cluster_centers_[1], color='black')
plt.legend((p1, p2, p3, c),
           ('Cluster 1', 'Cluster 2', 'Cluster 3', 'Centroid'))
```

```
Out[34]: <matplotlib.legend.Legend at 0x298b47d5970>
```



```
In [ ]:
```

Lab Program -8 :-

Apply EM algorithm to cluster a set of data stored in a .CSV file. Compare the results of k-Means algorithm and EM algorithm.

Source code and output :-

```
+*In[1]:*+
```

```
[source, ipython3]
```

```
----
```

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

```
from sklearn import datasets
```

```
from sklearn.cluster import KMeans
```

```
import sklearn.metrics as sm
```

```
import pandas as pd
```

```
import numpy as np
```

```
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']
```

```
model = KMeans(n_clusters=3)
```

```
model.fit(X)
```

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

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

```
# Plot the Original Classifications
```

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

```
# Plot the Models Classifications
```

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

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

```
plt.title('K Mean Classification')
```

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

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

```
print('The accuracy score of K-Mean: ', sm.accuracy_score(y, model.labels_))
```

```
print('The Confusion matrix of K-Mean: ', sm.confusion_matrix(y, model.labels_))
```

```
from sklearn import preprocessing
```

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

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

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

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

```
#xs.sample(5)
```

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

```
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)

y_gmm = gmm.predict(xs)
#y_cluster_gmm

plt.subplot(2, 2, 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('The accuracy score of EM: ',sm.accuracy_score(y, y_gmm))
print('The Confusion matrix of EM: ',sm.confusion_matrix(y, y_gmm))
```

```
+*Out[1]:*+
```

The accuracy score of K-Mean: 0.24

The Confusion matrix of K-Mean: [[0 50 0]

[48 0 2]

[14 0 36]]

The accuracy score of EM: 0.3533333333333333

The Confusion matrix of EM: [[5 0 45]

[2 48 0]

[0 50 0]]

![png](output_0_1.png)

+*In[]:*+

[source, ipython3]



File Edit View Insert Cell Kernel Widgets Help

Trusted

Python 3 (ipykernel) C

Code

```
In [1]: import matplotlib.pyplot as plt
from sklearn import datasets
from sklearn.cluster import KMeans
import sklearn.metrics as sm
import pandas as pd
import numpy as np

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']

model = KMeans(n_clusters=3)
model.fit(X)

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

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

# Plot the Original Classifications
plt.subplot(1, 2, 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')

# Plot the Models Classifications
plt.subplot(1, 2, 2)
plt.scatter(X.Petal_Length, X.Petal_Width, c=colormap[model.labels_], s=40)
plt.title('K Mean Classification')
plt.xlabel('Petal Length')
plt.ylabel('Petal Width')
print('The accuracy score of K-Mean: ', sm.accuracy_score(y, model.labels_))
print('The Confusion matrix of K-Mean: ', sm.confusion_matrix(y, model.labels_))

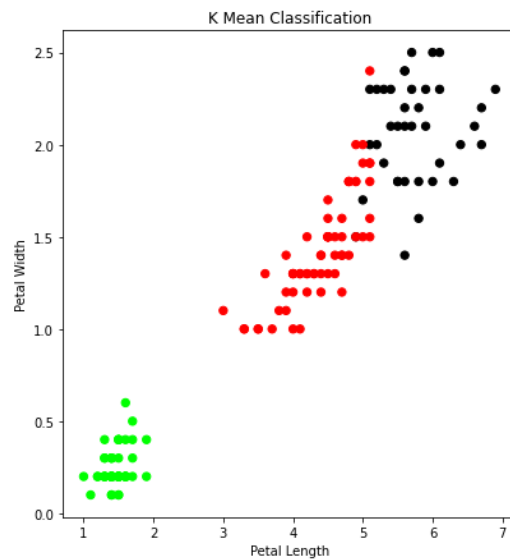
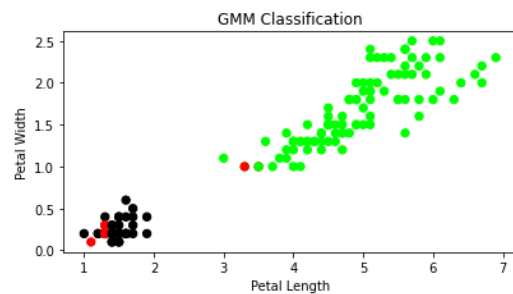
from sklearn import preprocessing
scaler = preprocessing.StandardScaler()
scaler.fit(X)
```



```
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('The accuracy score of EM: ',sm.accuracy_score(y, y_gmm))
print('The Confusion matrix of EM: ',sm.confusion_matrix(y, y_gmm))
```

```
The accuracy score of K-Mean: 0.24
The Confusion matrix of K-Mean: [[ 0 50  0]
 [48  0  2]
 [14  0 36]]
The accuracy score of EM: 0.35333333333333333
The Confusion matrix of EM: [[ 5  0 45]
 [ 2 48  0]
 [ 0 50  0]]
```



In []:

Lab Program -9:-

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

Source code and output :-

```
+#ln[1]:*+
```

```
[source, ipython3]
```

```
----
```

```
from sklearn.model_selection import train_test_split
```

```
from sklearn.neighbors import KNeighborsClassifier
```

```
from sklearn.metrics import classification_report, confusion_matrix
```

```
from sklearn import datasets
```

```
iris=datasets.load_iris()
```

```
x = iris.data
```

```
y = iris.target
```

```
print ('sepal-length', 'sepal-width', 'petal-length', 'petal-width')
```

```
print(x)
```

```
print('class: 0-Iris-Setosa, 1- Iris-Versicolour, 2- Iris-Virginica')
```

```
print(y)
```

```
x_train, x_test, y_train, y_test = train_test_split(x,y,test_size=0.3)
```

```
#To Training the model and Nearest nighbors K=5
```

```
classifier = KNeighborsClassifier(n_neighbors=5)
```

```
classifier.fit(x_train, y_train)
```

```
#To make predictions on our test data
```

```
y_pred=classifier.predict(x_test)
```

```
print('Confusion Matrix')
```

```
print(confusion_matrix(y_test,y_pred))
```

```
print('Accuracy Metrics')
```

```
print(classification_report(y_test,y_pred))
```

```
----
```

```
+*Out[1]:*+
```

```
----
```

```
sepal-length sepal-width petal-length petal-width
```

```
[[5.1 3.5 1.4 0.2]
```

```
[4.9 3. 1.4 0.2]
```

```
[4.7 3.2 1.3 0.2]
```

```
[4.6 3.1 1.5 0.2]
```

```
[5. 3.6 1.4 0.2]
```

```
[5.4 3.9 1.7 0.4]
```

```
[4.6 3.4 1.4 0.3]
```

```
[5. 3.4 1.5 0.2]
```

[4.4 2.9 1.4 0.2]

[4.9 3.1 1.5 0.1]

[5.4 3.7 1.5 0.2]

[4.8 3.4 1.6 0.2]

[4.8 3. 1.4 0.1]

[4.3 3. 1.1 0.1]

[5.8 4. 1.2 0.2]

[5.7 4.4 1.5 0.4]

[5.4 3.9 1.3 0.4]

[5.1 3.5 1.4 0.3]

[5.7 3.8 1.7 0.3]

[5.1 3.8 1.5 0.3]

[5.4 3.4 1.7 0.2]

[5.1 3.7 1.5 0.4]

[4.6 3.6 1. 0.2]

[5.1 3.3 1.7 0.5]

[4.8 3.4 1.9 0.2]

[5. 3. 1.6 0.2]

[5. 3.4 1.6 0.4]

[5.2 3.5 1.5 0.2]

[5.2 3.4 1.4 0.2]

[4.7 3.2 1.6 0.2]

[4.8 3.1 1.6 0.2]

[5.4 3.4 1.5 0.4]

[5.2 4.1 1.5 0.1]

[5.5 4.2 1.4 0.2]

[4.9 3.1 1.5 0.2]

[5. 3.2 1.2 0.2]

[5.5 3.5 1.3 0.2]

[4.9 3.6 1.4 0.1]

[4.4 3. 1.3 0.2]

[5.1 3.4 1.5 0.2]

[5. 3.5 1.3 0.3]

[4.5 2.3 1.3 0.3]

[4.4 3.2 1.3 0.2]

[5. 3.5 1.6 0.6]

[5.1 3.8 1.9 0.4]

[4.8 3. 1.4 0.3]

[5.1 3.8 1.6 0.2]

[4.6 3.2 1.4 0.2]

[5.3 3.7 1.5 0.2]

[5. 3.3 1.4 0.2]

[7. 3.2 4.7 1.4]

[6.4 3.2 4.5 1.5]

[6.9 3.1 4.9 1.5]

[5.5 2.3 4. 1.3]

[6.5 2.8 4.6 1.5]

[5.7 2.8 4.5 1.3]

[6.3 3.3 4.7 1.6]

[4.9 2.4 3.3 1.]

[6.6 2.9 4.6 1.3]

[5.2 2.7 3.9 1.4]

[5. 2. 3.5 1.]

[5.9 3. 4.2 1.5]

[6. 2.2 4. 1.]

[6.1 2.9 4.7 1.4]

[5.6 2.9 3.6 1.3]

[6.7 3.1 4.4 1.4]

[5.6 3. 4.5 1.5]

[5.8 2.7 4.1 1.]

[6.2 2.2 4.5 1.5]

[5.6 2.5 3.9 1.1]

[5.9 3.2 4.8 1.8]

[6.1 2.8 4. 1.3]

[6.3 2.5 4.9 1.5]

[6.1 2.8 4.7 1.2]

[6.4 2.9 4.3 1.3]

[6.6 3. 4.4 1.4]

[6.8 2.8 4.8 1.4]

[6.7 3. 5. 1.7]

[6. 2.9 4.5 1.5]

[5.7 2.6 3.5 1.]

[5.5 2.4 3.8 1.1]

[5.5 2.4 3.7 1.]

[5.8 2.7 3.9 1.2]

[6. 2.7 5.1 1.6]

[5.4 3. 4.5 1.5]

[6. 3.4 4.5 1.6]

[6.7 3.1 4.7 1.5]

[6.3 2.3 4.4 1.3]

[5.6 3. 4.1 1.3]

[5.5 2.5 4. 1.3]

[5.5 2.6 4.4 1.2]

[6.1 3. 4.6 1.4]

[5.8 2.6 4. 1.2]

[5. 2.3 3.3 1.]

[5.6 2.7 4.2 1.3]

[5.7 3. 4.2 1.2]

[5.7 2.9 4.2 1.3]

[6.2 2.9 4.3 1.3]

[5.1 2.5 3. 1.1]

[5.7 2.8 4.1 1.3]

[6.3 3.3 6. 2.5]

[5.8 2.7 5.1 1.9]

[7.1 3. 5.9 2.1]

[6.3 2.9 5.6 1.8]

[6.5 3. 5.8 2.2]

[7.6 3. 6.6 2.1]

[4.9 2.5 4.5 1.7]

[7.3 2.9 6.3 1.8]

[6.7 2.5 5.8 1.8]

[7.2 3.6 6.1 2.5]

[6.5 3.2 5.1 2.]

[6.4 2.7 5.3 1.9]

[6.8 3. 5.5 2.1]

[5.7 2.5 5. 2.]

[5.8 2.8 5.1 2.4]

[6.4 3.2 5.3 2.3]

[6.5 3. 5.5 1.8]

[7.7 3.8 6.7 2.2]

[7.7 2.6 6.9 2.3]

[6. 2.2 5. 1.5]

[6.9 3.2 5.7 2.3]

[5.6 2.8 4.9 2.]

[7.7 2.8 6.7 2.]

[6.3 2.7 4.9 1.8]

[6.7 3.3 5.7 2.1]

[7.2 3.2 6. 1.8]

[6.2 2.8 4.8 1.8]

[6.1 3. 4.9 1.8]

[6.4 2.8 5.6 2.1]

[7.2 3. 5.8 1.6]

[7.4 2.8 6.1 1.9]

[7.9 3.8 6.4 2.]

[6.4 2.8 5.6 2.2]

[6.3 2.8 5.1 1.5]

[6.1 2.6 5.6 1.4]

[7.7 3. 6.1 2.3]

[6.3 3.4 5.6 2.4]

[6.4 3.1 5.5 1.8]

[6. 3. 4.8 1.8]

[6.9 3.1 5.4 2.1]

[6.7 3.1 5.6 2.4]

[6.9 3.1 5.1 2.3]

[5.8 2.7 5.1 1.9]

[6.8 3.2 5.9 2.3]

[6.7 3.3 5.7 2.5]

[6.7 3. 5.2 2.3]

[6.3 2.5 5. 1.9]

[6.5 3. 5.2 2.]

[6.2 3.4 5.4 2.3]

[5.9 3. 5.1 1.8]]

class: 0-Iris-Setosa, 1- Iris-Versicolour, 2- Iris-Virginica

```
[0000000000000000000000000000000000000000000000000
```

0000000000000011111111111111111111111111

111

222

 $2\ 2]$

Confusion Matrix

[[12 0 0]]

$$[0 \ 14 \ 0]$$
$$\begin{bmatrix} 0 & 0 & 19 \end{bmatrix}$$

Accuracy Metrics

```
precision  recall  f1-score  support
```

| | | | | |
|---|------|------|------|----|
| 0 | 1.00 | 1.00 | 1.00 | 12 |
|---|------|------|------|----|

| | | | | |
|---|------|------|------|----|
| 1 | 1.00 | 1.00 | 1.00 | 14 |
|---|------|------|------|----|

| | | | | |
|---|------|------|------|----|
| 2 | 1.00 | 1.00 | 1.00 | 19 |
|---|------|------|------|----|

| | | | | |
|----------|--|--|------|----|
| accuracy | | | 1.00 | 45 |
|----------|--|--|------|----|

| | | | | |
|-----------|------|------|------|----|
| macro avg | 1.00 | 1.00 | 1.00 | 45 |
|-----------|------|------|------|----|

| | | | | |
|--------------|------|------|------|----|
| weighted avg | 1.00 | 1.00 | 1.00 | 45 |
|--------------|------|------|------|----|

+*ln[]:*+

[source, ipython3]

```
In [1]: from sklearn.model_selection import train_test_split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import classification_report, confusion_matrix
from sklearn import datasets

iris=datasets.load_iris()

x = iris.data
y = iris.target

print ('sepal-length', 'sepal-width', 'petal-length', 'petal-width')
print(x)
print('class: 0- Iris-Setosa, 1- Iris-Versicolour, 2- Iris-Virginica')
print(y)

x_train, x_test, y_train, y_test = train_test_split(x,y,test_size=0.3)

#To Training the model and Nearest neighbors K=5
classifier = KNeighborsClassifier(n_neighbors=5)
classifier.fit(x_train, y_train)

#To make predictions on our test data
y_pred=classifier.predict(x_test)

print('Confusion Matrix')
print(confusion_matrix(y_test,y_pred))
print('Accuracy Metrics')
print(classification_report(y_test,y_pred))
```

```
sepal-length sepal-width petal-length petal-width
[[5.1 3.5 1.4 0.2]
 [4.9 3.  1.4 0.2]
 [4.7 3.2 1.3 0.2]
 [4.6 3.1 1.5 0.2]
 [5.  3.6 1.4 0.2]
 [5.4 3.9 1.7 0.4]
 [4.6 3.4 1.4 0.3]
 [5.  3.4 1.5 0.2]
 [4.4 2.9 1.4 0.2]
 [4.9 3.1 1.5 0.1]
 [5.4 3.7 1.5 0.2]
 [4.8 3.4 1.6 0.2]
 [4.8 3.  1.4 0.1]
 [4.3 3.  1.1 0.1]]
```

| | | | | |
|--------------|------|------|------|----|
| weighted avg | 1.00 | 1.00 | 1.00 | 45 |
|--------------|------|------|------|----|

Lab Program -10 :-

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

Source code and output :-

```
+*In[2]:*+
```

```
[source, ipython3]
```

```
----
```

```
import numpy as np
```

```
from bokeh.plotting import figure, show, output_notebook
```

```
from bokeh.layouts import gridplot
```

```
from bokeh.io import push_notebook
```

```
from matplotlib import pyplot as plt
```

```
def local_regression(x0, X, Y, tau):# add bias term
```

```
    x0 = np.r_[1, x0] # Add one to avoid the loss in information
```

```
    X = np.c_[np.ones(len(X)), X]
```

```
    # fit model: normal equations with kernel
```

```
    xw = X.T * radial_kernel(x0, X, tau) # XTranspose * W
```

```
    beta = np.linalg.pinv(xw @ X) @ xw @ Y #@ Matrix Multiplication or Dot Product
```

```

# predict value
return x0 @ beta # @ Matrix Multiplication or Dot Product for prediction
def radial_kernel(x0, X, tau):
    return np.exp(np.sum((X - x0) ** 2, axis=1) / (-2 * tau * tau))
# Weight or Radial Kernal Bias Function

n = 1000
# generate dataset
X = np.linspace(-3, 3, num=n)
print("The Data Set ( 10 Samples) X :\n",X[1:10])
Y = np.log(np.abs(X ** 2 - 1) + .5)
print("The Fitting Curve Data Set (10 Samples) Y :\n",Y[1:10])
# jitter X
X += np.random.normal(scale=.1, size=n)
print("Normalised (10 Samples) X :\n",X[1:10])

domain = np.linspace(-3, 3, num=300)
print(" Xo Domain Space(10 Samples) :\n",domain[1:10])
def plot_lwr(tau):
    # prediction through regression
    prediction = [local_regression(x0, X, Y, tau) for x0 in domain]
    plot = figure(plot_width=400, plot_height=400)
    plot.title.text='tau=%g' % tau
    plot.scatter(X, Y, alpha=.3)
    plot.line(domain, prediction, line_width=2, color='red')
    return plot

```

```
show(gridplot([
    [plot_lwr(10.), plot_lwr(1.)],
    [plot_lwr(0.1), plot_lwr(0.01)]]))
plt.title('K Mean Classification')
plt.xlabel('Petal Length')
```

```
+*Out[2]:*+
```

The Data Set (10 Samples) X :

```
[-2.99399399 -2.98798799 -2.98198198 -2.97597598 -2.96996997 -2.96396396
-2.95795796 -2.95195195 -2.94594595]
```

The Fitting Curve Data Set (10 Samples) Y :

```
[2.13582188 2.13156806 2.12730467 2.12303166 2.11874898 2.11445659
2.11015444 2.10584249 2.10152068]
```

Normalised (10 Samples) X :

```
[-2.88440998 -2.97461063 -2.97639127 -2.9042727 -3.1194782 -3.06506157
-2.8349021 -2.90676221 -2.92454458]
```

Xo Domain Space(10 Samples) :

```
[-2.97993311 -2.95986622 -2.93979933 -2.91973244 -2.89966555 -2.87959866
-2.85953177 -2.83946488 -2.81939799]
```

```
Text(0.5, 0, 'Petal Length')
```

```
![png](output_0_2.png)
```

```
+*In[3]:*+
```

```
[source, ipython3]
```

```
----
```

```
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))) # eye - identity matrix
```

```
    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,yamat,k):
```

```
    wei = kernel(point,xmat,k)
```

```
    W = (X.T*(wei*X)).I*(X.T*(wei*yamat.T))
```

```
    return W
```

```
def localWeightRegression(xmat,yamat,k):
```

```
    m,n = np.shape(xmat)
```

```
    ypred = np.zeros(m)
```

```
    for i in range(m):
```



```
ypred[i] = xmat[i]*localWeight(xmat[i],xmat,yamat,k)
```

```
return ypred
```

```
def graphPlot(X,ypred):
```

```
    sortindex = X[:,1].argsort(0) #argsort - index of the smallest
```

```
    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();
```

```
# load data points
```

```
data = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 10\tips.csv')
```

```
bill = np.array(data.total_bill) # We use only Bill amount and Tips data
```

```
tip = np.array(data.tip)
```

```
mbill = np.mat(bill) # .mat will convert nd array is converted in 2D array
```

```
mtip = np.mat(tip)
```

```
m= np.shape(mbill)[1]
```

```
one = np.mat(np.ones(m))
```

```
X = np.hstack((one.T,mbill.T)) # 244 rows, 2 cols
```

```
# increase k to get smooth curves
```

```
ypred = localWeightRegression(X,mtip,3)
```

```
graphPlot(X,ypred)
```

```
----
```

```
+*Out[3]:*+
```

```
----
```

```
![png](output_1_0.png)
```

```
----
```

```
+*In[ ]:*+
```

```
[source, ipython3]
```

```
----
```

| | A | B | C | D | E | F | G | H | I |
|----|------------|------|--------|--------|-----|--------|------|---|---|
| 1 | total_bill | tip | sex | smoker | day | time | size | | |
| 2 | 16.99 | 1.01 | Female | No | Sun | Dinner | 2 | | |
| 3 | 10.34 | 1.66 | Male | No | Sun | Dinner | 3 | | |
| 4 | 21.01 | 3.5 | Male | No | Sun | Dinner | 3 | | |
| 5 | 23.68 | 3.31 | Male | No | Sun | Dinner | 2 | | |
| 6 | 24.59 | 3.61 | Female | No | Sun | Dinner | 4 | | |
| 7 | 25.29 | 4.71 | Male | No | Sun | Dinner | 4 | | |
| 8 | 8.77 | 2 | Male | No | Sun | Dinner | 2 | | |
| 9 | 26.88 | 3.12 | Male | No | Sun | Dinner | 4 | | |
| 10 | 15.04 | 1.96 | Male | No | Sun | Dinner | 2 | | |
| 11 | 14.78 | 3.23 | Male | No | Sun | Dinner | 2 | | |
| 12 | 10.27 | 1.71 | Male | No | Sun | Dinner | 2 | | |
| 13 | 35.26 | 5 | Female | No | Sun | Dinner | 4 | | |
| 14 | 15.42 | 1.57 | Male | No | Sun | Dinner | 2 | | |
| 15 | 18.43 | 3 | Male | No | Sun | Dinner | 4 | | |
| 16 | 14.83 | 3.02 | Female | No | Sun | Dinner | 2 | | |
| 17 | 21.58 | 3.92 | Male | No | Sun | Dinner | 2 | | |
| 18 | 10.33 | 1.67 | Female | No | Sun | Dinner | 3 | | |
| 19 | 16.29 | 3.71 | Male | No | Sun | Dinner | 3 | | |
| 20 | 16.97 | 3.5 | Female | No | Sun | Dinner | 3 | | |
| 21 | 20.65 | 3.35 | Male | No | Sat | Dinner | 3 | | |
| 22 | 17.92 | 4.08 | Male | No | Sat | Dinner | 2 | | |
| 23 | 20.29 | 2.75 | Female | No | Sat | Dinner | 2 | | |
| 24 | 15.77 | 2.23 | Female | No | Sat | Dinner | 2 | | |
| 25 | 39.42 | 7.58 | Male | No | Sat | Dinner | 4 | | |

tips



```

In [2]: import numpy as np
from bokeh.plotting import figure, show, output_notebook
from bokeh.layouts import gridplot
from bokeh.io import push_notebook
from matplotlib import pyplot as plt

def local_regression(x0, X, Y, tau):# add bias term
    x0 = np.r_[1, x0] # Add one to avoid the loss in information
    X = np.c_[np.ones(len(X)), X]

    # fit model: normal equations with kernel
    xw = X.T * radial_kernel(x0, X, tau) # XTranspose * W

    beta = np.linalg.pinv(xw @ X) @ xw @ Y #@ Matrix Multiplication or Dot Product

    # predict value
    return x0 @ beta # @ Matrix Multiplication or Dot Product for prediction
def radial_kernel(x0, X, tau):
    return np.exp(np.sum((X - x0) ** 2, axis=1) / (-2 * tau * tau))
# Weight or Radial Kernel Bias Function

n = 1000
# generate dataset
X = np.linspace(-3, 3, num=n)
print("The Data Set ( 10 Samples) X :\n",X[1:10])
Y = np.log(np.abs(X ** 2 - 1) + .5)
print("The Fitting Curve Data Set (10 Samples) Y :\n",Y[1:10])
# jitter X
X += np.random.normal(scale=.1, size=n)
print("Normalised (10 Samples) X :\n",X[1:10])

domain = np.linspace(-3, 3, num=300)
print(" Xo Domain Space(10 Samples) :\n",domain[1:10])
def plot_lwr(tau):
    # prediction through regression
    prediction = [local_regression(x0, X, Y, tau) for x0 in domain]
    plot = figure(plot_width=400, plot_height=400)
    plot.title.text='tau=%g' % tau
    plot.scatter(X, Y, alpha=.3)
    plot.line(domain, prediction, line_width=2, color='red')
    return plot

```

```

plot.title.text= tau=%g % tau
plot.scatter(X, Y, alpha=.3)
plot.line(domain, prediction, line_width=2, color='red')
return plot

show(gridplot([
    [plot_lwr(10.), plot_lwr(1.)],
    [plot_lwr(0.1), plot_lwr(0.01)]]))
plt.title('K Mean Classification')
plt.xlabel('Petal Length')

```

The Data Set (10 Samples) X :

```

[-2.99399399 -2.98798799 -2.98198198 -2.97597598 -2.96996997 -2.96396396
-2.95795796 -2.95195195 -2.94594595]

```

The Fitting Curve Data Set (10 Samples) Y :

```

[2.13582188 2.13156806 2.12730467 2.12303166 2.11874898 2.11445659
2.11015444 2.10584249 2.10152068]

```

Normalised (10 Samples) X :

```

[-2.88440998 -2.97461063 -2.97639127 -2.9042727 -3.1194782 -3.06506157
-2.8349021 -2.90676221 -2.92454458]

```

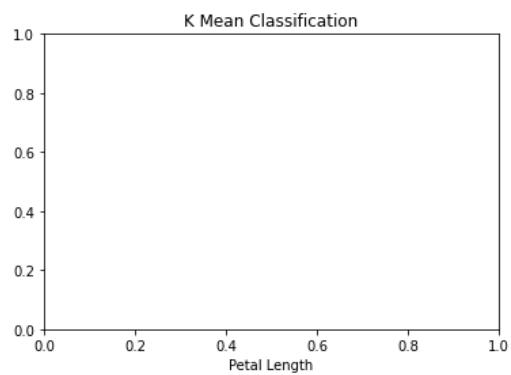
Xo Domain Space(10 Samples) :

```

[-2.97993311 -2.95986622 -2.93979933 -2.91973244 -2.89966555 -2.87959866
-2.85953177 -2.83946488 -2.81939799]

```

Out[2]: Text(0.5, 0, 'Petal Length')



```

In [3]: 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))) # eye - identity matrix
    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,yamat,k):
    wei = kernel(point,xmat,k)
    W = (X.T*(wei*X)).I*(X.T*(wei*yamat.T))
    return W

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

def graphPlot(X,ypred):
    sortindex = X[:,1].argsort(0) #argsort - index of the smallest
    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();

# Load data points
data = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 10\tips.csv')
bill = np.array(data.total_bill) # We use only Bill amount and Tips data
tip = np.array(data.tip)

mbill = np.mat(bill) # .mat will convert nd array is converted in 2D array
mtip = np.mat(tip)
m = np.shape(mbill)[1]
one = np.mat(np.ones(m))
X = np.hstack((one.T,mbill.T)) # 244 rows, 2 cols
plt.ylabel('Tip')
plt.show();

# Load data points
data = pd.read_csv(r'C:\Users\Admin\OneDrive\Desktop\6th sem\ML\lab-ml\Lab 10\tips.csv')
bill = np.array(data.total_bill) # We use only Bill amount and Tips data
tip = np.array(data.tip)

mbill = np.mat(bill) # .mat will convert nd array is converted in 2D array
mtip = np.mat(tip)
m = np.shape(mbill)[1]
one = np.mat(np.ones(m))
X = np.hstack((one.T,mbill.T)) # 244 rows, 2 cols

# increase k to get smooth curves
ypred = localWeightRegression(X,mtip,3)
graphPlot(X,ypred)

```

