

VISVESVARAYATECHNOLOGICAL UNIVERSITY

“JnanaSangama”, Belgaum-590014, Karnataka.



LABREPORT
on

MACHINELEARNING(20CS6PCMAL)

Submittedby

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inpartialfulfillmentfortheawardofthedegreeof
BACHELOROFENGINEERING

in
COMPUTERSCIENCEANDENGINEERING



B.M.S.COLLEGE OF ENGINEERING
(Autonomous Institution under VTU)

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B.M.S.CollegeofEngineering,
BullTempleRoad,Bangalore560019
(AffiliatedToVisvesvarayaTechnologicalUniversity,Belgaum)
DepartmentofComputerScienceandEngineering



CERTIFICATE

This is to certify that the Lab work entitled “**MACHINELEARNING**” carried out by **MOHDIRFAN(1BM20CS409)**, who is a bonafide 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** Visvesvaraya Technological University, Belgaum during the year 2022. The Lab report has been approved as it satisfies the academic requirements in respect of a **Machine Learning- (20CS6PCMAL)** work prescribed for the said degree.

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1.ImplementFINDSALGORITHM

In [5]:

```
import csv
```

```
a=[]
```

In [6]:

```
with open('enjoysports.csv','r') as csvfile:
```

```
    for row in csv.reader(csvfile):
```

```
        a.append(row)
```

```
    print(a)
```

```
print("\nThe Total Number of Training Instances are: ",len(a))
```

```
[['sunny', 'warm', 'normal', 'strong', 'warm', 'same', 'yes'], ['sunny', 'wa  
rm', 'high', 'strong', 'warm', 'same', 'yes'], ['rainy', 'cold', 'high', 'st  
rong', 'warm', 'change', 'no'], ['sunny', 'warm', 'high', 'strong', 'cool',  
'change', 'yes']]
```

The Total Number of Training Instances are: 4

In [7]:

```
num_attribute = len(a[0])-1
```

```
print("\nThe Initial Hypothesis is: ")
```

```
hypothesis = ['0']*num_attribute
```

```
print(hypothesis)
```

```
for i in range(0,len(a)):
```

```
    if a[i][num_attribute] == 'yes':
```

```
        for j in range(0,num_attribute):
```

```
            if hypothesis[j] == '0' or hypothesis[j] == a[i][j]:
```

```
    if hypothesis[j] == '0' or hypothesis[j] == a[i][j]:
        hypothesis[j] = a[i][j]
    else:
        hypothesis[j] = '?'

print("\nThe Hypothesis for the training instance {} is: {}".format(i+1, hypothesis))
```

The Initial Hypothesis is:

['0', '0', '0', '0', '0', '0']

The Hypothesis for the training instance 1 is:

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

The Hypothesis for the training instance 2 is:

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

The Hypothesis for the training instance 3 is:

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

The Hypothesis for the training instance 4 is:

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

In [8]:

```
print("\nThe Maximally Specific Hypothesis for the training instance is ")  
print(hypothesis)
```

The Maximally Specific Hypothesis for the training instance is
['sunny', 'warm', '?', 'strong', '?', '?']

2..ImplementCandidateeliminationAlgorithm

```
import numpy as np
import pandas as pd

data = pd.DataFrame(data = pd.read_csv('enjoysports.csv'))
concepts = np.array(data.iloc[:,0:-1])
print(concepts)

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

[['sunny' 'warm' 'normal' 'strong' 'warm' 'same']
 ['sunny' 'warm' 'high' 'strong' 'warm' 'same']
 ['rainy' 'cold' 'high' 'strong' 'warm' 'change']
 ['sunny' 'warm' 'high' 'strong' 'cool' 'change']]
['yes' 'yes' 'no' 'yes']

def learn(concepts, target):
    specific_h = concepts[0].copy()
    print("Initialization Of Specific Hypothesis & General Hypothesis : ")
    print(specific_h)
    general_h = [['?' for i in range(len(specific_h))] for i in range(len(specific_h))]
    print(general_h)

    for i, h in enumerate(concepts):
        if target[i] == "yes":
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    specific_h[x] = '?'
                    general_h[x][x] = '?'
            print(specific_h)
            print(specific_h)

        if target[i] == "no":
            for x in range(len(specific_h)):
                if h[x] != specific_h[x]:
                    general_h[x][x] = specific_h[x]
                else:
                    general_h[x][x] = '?'
            print("Steps of Candidate Elimination Algorithm: ",i+1)
            print(specific_h)
            print(general_h)
    indices = [i for i, val in enumerate(general_h) if val == ['?', '?', '?', '?', '?', '?']]
    for i in indices:
        general_h.remove(['?', '?', '?', '?', '?', '?'])
    return specific_h, general_h
```

```
print("Final Specific Hypothesis: ",s_final, sep="\n")
print("Final General Hypothesis: ",g_final, sep="\n")
```

Final Specific Hypothesis:

['sunny' 'warm' '?' 'strong' '?' '?']

Final General Hypothesis:

[['sunny', '?', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?']]

3.IMPLEMENTID-3ALGORITHM

```
In [ ]: import numpy as np

In [ ]: import pandas as pd
from sklearn.tree import DecisionTreeClassifier # Import Decision Tree Classifier
from sklearn.model_selection import train_test_split # Import train_test_split function
from sklearn import metrics #Import scikit-learn metrics module for accuracy calculation

In [4]: col_names = ['pregnant', 'glucose', 'bp', 'skin', 'insulin', 'bmi', 'pedigree', 'age', 'label']
pima = pd.read_csv("/content/drive/MyDrive/diabetes.csv", header=None, names=col_names)

In [5]: pima.head()

Out[5]:
```

	pregnant	glucose	bp	skin	insulin	bmi	pedigree	age	label
0	6	148	72	35	0	33.6	0.627	50	1
1	1	85	66	29	0	26.6	0.351	31	0
2	8	183	64	0	0	23.3	0.672	32	1
3	1	89	66	23	94	28.1	0.167	21	0
4	0	137	40	35	168	43.1	2.288	33	1

```
In [6]: feature_cols = ['pregnant', 'insulin', 'bmi', 'age', 'glucose', 'bp', 'pedigree']
X = pima[feature_cols] # Features
y = pima.label # Target variable

In [7]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=2)

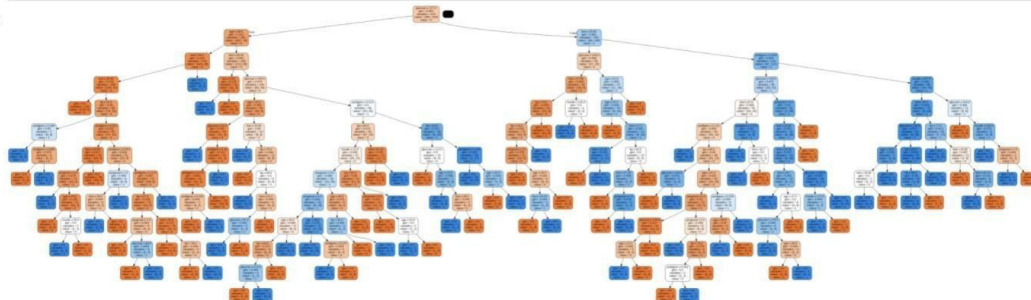
In [8]: clf = DecisionTreeClassifier()
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
```

```
clf = clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
print("Accuracy:", metrics.accuracy_score(y_test, y_pred))
```

Accuracy: 0.7467532467532467

```
In [9]: from sklearn.tree import export_graphviz
from six import StringIO
from IPython.display import Image
import pydotplus

dot_data = StringIO()
export_graphviz(clf, out_file=dot_data,
                filled=True, rounded=True,
                special_characters=True, feature_names = feature_cols, class_names=['0', '1'])
graph = pydotplus.graph_from_dot_data(dot_data.getvalue())
graph.write_png('diabetes.png')
Image(graph.create_png())
```



In []:

4.IMPLEMENTNAIVEBAYES

In [23]:



```
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.naive_bayes import GaussianNB
from sklearn import metrics
```

In [24]:



```
df = pd.read_csv("pima_indian.csv")
feature_col_names = ['num_preg', 'glucose_conc', 'diastolic_bp', 'thickness', 'insulin', 'b
predicted_class_names = ['diabetes']
```

In [25]:



```
X = df[feature_col_names].values
y = df[predicted_class_names].values
```

In [26]:



```
print(df.head)
xtrain,xtest,ytrain,ytest=train_test_split(X,y,test_size=0.33)

print ('\n the total number of Training Data :',ytrain.shape)
print ('\n the total number of Test Data :',ytest.shape)
```

<bound method NDFrame.head of	num_preg	glucose_conc	diastolic_bp	thi		
ckness insulin bmi \						
0	6	148	72	35	0	33.6
1	1	85	66	29	0	26.6
2	8	183	64	0	0	23.3
3	1	89	66	23	94	28.1
4	0	137	40	35	168	43.1
..
763	10	101	76	48	180	32.9
764	2	122	70	27	0	36.8
765	5	121	72	23	112	26.2
766	1	126	60	0	0	30.1
767	1	93	70	31	0	30.4

	diab_pred	age	diabetes
0	0.627	50	1
1	0.351	31	0
2	0.672	32	1
3	0.167	21	0
4	2.288	33	1
..
763	0.171	63	0
764	0.340	27	0
765	0.245	30	0
766	0.349	47	1
767	0.315	23	0

[768 rows x 9 columns]>

the total number of Training Data : (514, 1)

the total number of Test Data : (254, 1)

In [27]:



```
clf = GaussianNB().fit(xtrain,ytrain.ravel())
predicted = clf.predict(xtest)
predictTestData= clf.predict([[6,148,72,35,0,33.6,0.627,50]])
```

In [28]:



```
print('\n Confusion matrix')
print(metrics.confusion_matrix(ytest,predicted))

print('\n Accuracy of the classifier is',metrics.accuracy_score(ytest,predicted))

print('\n The value of Precision', metrics.precision_score(ytest,predicted))

print('\n The value of Recall', metrics.recall_score(ytest,predicted))

print(" Predicted Value for individual Test Data:", predictTestData)
```

```
Confusion matrix
[[139  20]
 [ 44  51]]
```

```
Accuracy of the classifier is 0.7480314960629921
```

```
The value of Precision 0.7183098591549296
```

```
The value of Recall 0.5368421052631579
```

```
Predicted Value for individual Test Data: [1]
```

5.IMPLEMENTLINEARREGRESSION

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

```
In [28]: dataset = pd.read_csv('Salary_Data.csv')
dataset.head()
```

```
Out[28]:
```

	YearsExperience	Salary
0	1.1	39343.0
1	1.3	46205.0
2	1.5	37731.0
3	2.0	43525.0
4	2.2	39891.0

```
In [19]: X = dataset.iloc[:, :-1].values
print(X)
```

```
<class 'numpy.ndarray'>
```

```
In [6]: y = dataset.iloc[:, -1].values
```

```
In [10]: dataset.head()
```

```
Out[10]:
```

	YearsExperience	Salary
0	1.1	39343.0
1	1.3	46205.0
2	1.5	37731.0
3	2.0	43525.0
4	2.2	39891.0

```
In [11]: from sklearn.model_selection import train_test_split
```

```
In [12]: X_train, X_test, y_train, y_test = train_test_split(X, y, test_size = 1/3, random_state = 0)
```

```
In [14]: from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
```

```
Out[14]: LinearRegression(copy_X=True, fit_intercept=True, n_jobs=None, normalize=False)
```

```
In [15]: y_pred = regressor.predict(X_test)
plt.scatter(X_train, y_train, color = 'red')
plt.plot(X_test, regressor.predict(X_test), color = 'blue')
plt.title('Salary vs Experience (Training set)')
plt.xlabel('Years of Experience')
plt.ylabel('Salary')
plt.show()
```

```
In [16]: pd.DataFrame(data={'Actuals': y_test, 'Predictions': y_pred})
```

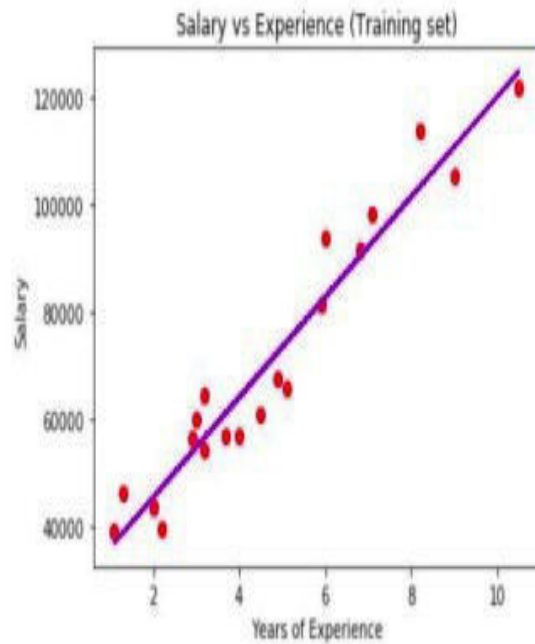
```
Out[16]:
```

	Actuals	Predictions
0	37731.0	40835.105909
1	122391.0	123079.399408
2	57081.0	65134.556261
3	63218.0	63265.367772
4	116969.0	115602.645454
5	109431.0	108125.891499
6	112635.0	116537.239698
7	55794.0	64199.962017
8	83088.0	76349.687193

row setup...

```
7 55794.0 64199.962017
8 83088.0 76349.687193
9 101302.0 100649.137545
```

```
In [17]: plt.scatter(X_train, y_train, color = 'red')
plt.plot(X_train, regressor.predict(X_train), color = 'blue')
plt.title('Salary vs Experience (Training set)')
plt.xlabel('Years of Experience')
plt.ylabel('Salary')
plt.show()
```



Program 6: K Means

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sns
sns.set()
from sklearn.cluster import KMeans
```

```
data = pd.read_csv("../input/K-Means/Dataset.csv")
```

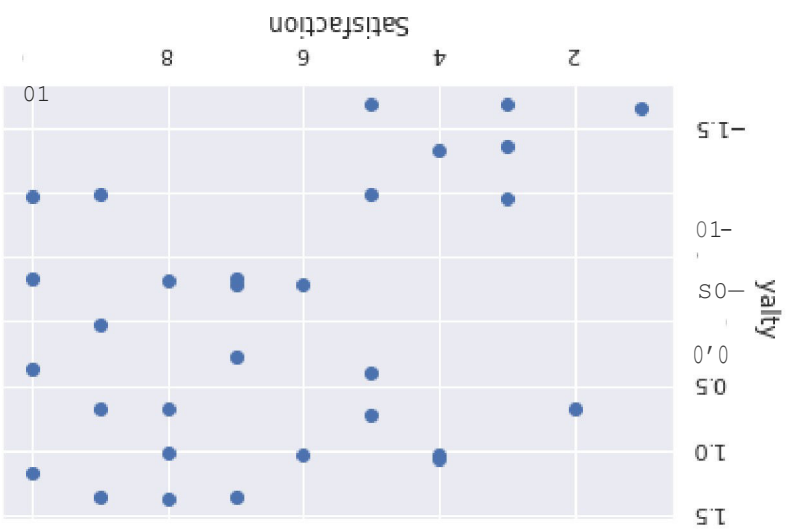
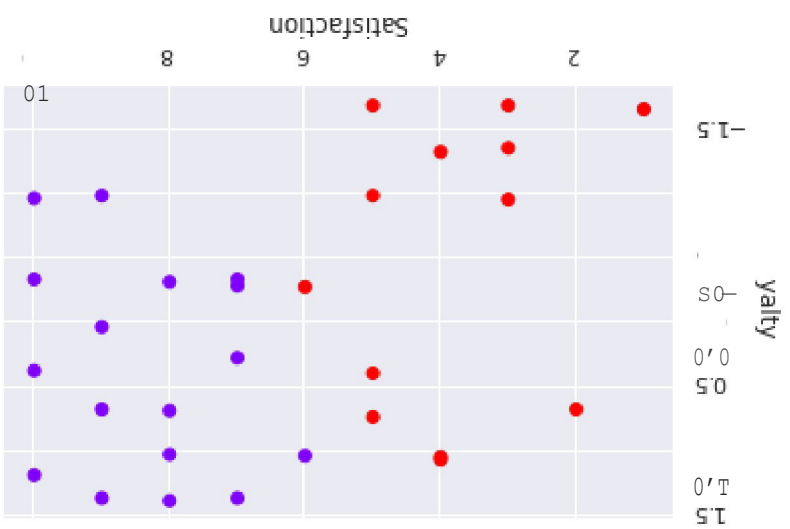
```
plt.scatter(data['Satisfaction'], data['Loyalty'])
plt.xlabel('Satisfaction')
plt.ylabel('Loyalty')
plt.show()
```

```
x = data.copy()
kmean = KMeans(2)
kmean.fit(x)
```

```
clusters = x.copy()
clusters['cluster_pred'] = kmean.fit_predict(x)
```

```
plt.scatter(clusters['Satisfaction'], clusters['Loyalty'], c=clusters['cluster_pred'], cmap='rainbow')
plt.xlabel('Satisfaction')
plt.ylabel('Loyalty')
plt.show()
```

Satisfaction	Loyalty
4	-1.33
6	-0.28
5	-0.99
7	-0.29
4	1.06
1	-1.66
10	-0.97
8	-0.32
8	1.02
8	0.68
10	-0.34
5	0.39
5	-1.69
2	0.67
7	0.27



Program 7: EM Algorithm

```
import matplotlib.pyplot as
plt from sklearn
import datasets
from sklearn.cluster import
KMeans import sklearn.metrics
asm
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:\n',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)
```

```

from sklearn.mixture import
GaussianMixture gmm =
GaussianMixture(n_components=3)
gmm.fit(xs)

y_gmm =

gmm.predict(xs)

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:\n',sm.confusion_matrix(y,
y_gmm))

```

<Figure size 1008x504 with 0 Axes>

Text(0, 0.5, 'Petal Width')



Text(0, 0.5, 'Petal Width')



The accuracy score of K-Mean: 0.8933333333333333

The Confusion matrix of K-Mean:

```
[[50  0  0]
```

```
[ 0 48  2]
```

```
[ 0 14 36]]
```

Text(0, 0.5, 'Petal Width')



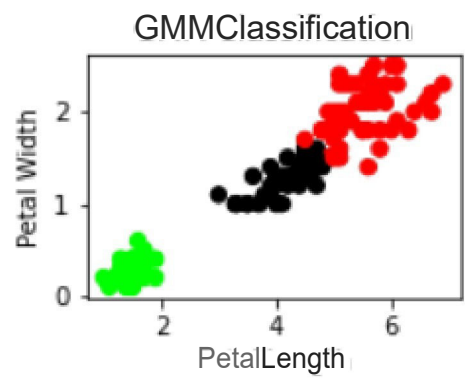
The accuracy score of EM: 0.0

The Confusion matrix of EM:

```
[[ 0 50  0]
```

```
[ 5  0 45]
```

```
[50  0  0]]
```



Program 8 : K Nearest Neighbor

```
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('target'
)print(Y)

x_train, x_test, y_train, y_test = train_test_split(X,Y,test_size=0.3)

classier =
KNeighborsClassifier(n_neighbors=5)
classier.fit(x_train, y_train)

y_pred=classier.predict(x_test)

print('Confusion Matrix')
print(confusion_matrix(y_test,y_pred)
)

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

[illegible]

Confusion Matrix

[[19 0 0]]

$$\begin{bmatrix} 0 & 16 & 1 \end{bmatrix}$$
$$\begin{bmatrix} 0 & 0 & 9 \end{bmatrix}$$

Accuracy

	precision	recall	f1-score	support
0	1.00	1.00	1.00	19
1	1.00	0.94	0.97	17
2	0.90	1.00	0.95	9
accuracy			0.98	45
macro avg	0.97	0.98	0.97	45
weighted avg	0.98	0.98	0.98	45

Program 9: Bayesian Network

```
import numpy as
np import pandas
as pd import csv
import pgmpy
from pgmpy.estimators import
MaximumLikelihoodEstimator from pgmpy.models import
BayesianNetwork
from pgmpy.inference import VariableElimination

#read Cleveland Heart Disease data
heartDisease =
pd.read_csv('../input/heartdisease/heartDiseaseDataset.csv') heartDisease
= heartDisease.replace('?',np.nan)

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

#display the Attributes names and
datatypes print('\n Attributes and
datatypes') print(heartDisease.dtypes)

#Create Model-Bayesian
Network model =
BayesianNetwork([('age','heartDisease'),('sex','heartDisease'),('exang','heartDisease'),('cp','heartDisease'),
('restecg','heartDisease'),('heartDisease','chol')])

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

#Inferencing with Bayesian Network
print('\n Inferencing with Bayesian Network:')
heartDiseasetest_infer =
VariableElimination(model)

#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)

#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)
```

age	sex	cp	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	heartDisease
63	1	1	145	233	1	2	150	0	2.3	3	0	6	0
67	1	4	160	286	0	2	108	1	1.5	2	3	3	2
67	1	4	120	229	0	2	129	1	2.6	2	2	7	1
37	1	3	130	250	0	0	187	0	3.5	3	0	3	0
41	0	2	130	204	0	2	172	0	1.4	1	0	3	0
56	1	2	120	236	0	0	178	0	0.8	1	0	3	0
62	0	4	140	268	0	2	160	0	3.6	3	2	3	3
57	0	4	120	354	0	0	163	1	0.6	1	0	3	0
63	1	4	130	254	0	2	147	0	1.4	2	1	7	2
53	1	4	140	203	1	2	155	1	3.1	3	0	7	1
57	1	4	140	192	0	0	148	0	0.4	2	0	6	0
56	0	2	140	294	0	2	153	0	1.3	2	0	3	0
56	1	3	130	256	1	2	142	1	0.6	2	1	6	2
44	1	2	120	263	0	0	173	0	0	1	0	7	0
52	1	3	172	199	1	0	162	0	0.5	1	0	7	0
57	1	3	150	168	0	0	174	0	1.6	1	0	3	0
48	1	2	110	229	0	0	168	0	1	3	0	7	1
54	1	4	140	239	0	0	160	0	1.2	1	0	3	0
48	0	3	130	275	0	0	139	0	0.2	1	0	3	0
49	1	2	130	266	0	0	171	0	0.6	1	0	3	0
64	1	1	110	211	0	2	144	1	1.8	2	0	3	0
58	0	1	150	283	1	2	162	0	1	1	0	3	0

Sample instances from the dataset are given below

```

age sex cp trestbps chol fbs restecg thalach exang oldpeak slope \
0 63 1 1 145 233 1 2 150 0 2.3 3
1 67 1 4 160 286 0 2 108 1 1.5 2
2 67 1 4 120 229 0 2 129 1 2.6 2
3 37 1 3 130 250 0 0 187 0 3.5 3
4 41 0 2 130 204 0 2 172 0 1.4 1

```

```

ca thal heartDisease
0 0 6 0
1 3 3 2
2 2 7 1
3 0 3 0
4 0 3 0

```

```

Attributes and datatypes
age int64
sex int64
cp int64
trestbps int64
chol int64
fbs int64
restecg int64
thalach int64
exang int64
oldpeak float64
slope int64
ca int64
thal int64
heartDisease int64
dtype: object

```

Learning CP using Maximum Likelihood Estimators Inference with B

Bayesian Network:

6. Probability of heartDisease given evidence = restecg

Finding

Elimination Order: 100% Elimination

g: exang: 10096

4/4[00:00<00:00,81.79it/s]

:1 4/4[00:DOEOD:0064.63it/s]

heartDiseaseh phi(heartDisease)

heartDisease(B) he

artDisease(1)heart	8.1978
D1sea se(2)	0.1976
heartDisease(3)	0.1972
heartDisease(4)	0.1976

7. Probab11ityofheartD1s ea se given evidence=cp:2

FindingEliminationOrder::100%Eliminating:e

xang:10096	0.2106
------------	--------

4/4[00:OOH00:00,69.B 5lt/s]

4/4[00:DOEOD:00 89.18it/s]

heartD1sea ph1heartDisease)

seheartDisease(B)

)

+heartDisease(1)-	
+heartDisease(2)-	0.1653
+heartD1sease(3)-	0.3138
+heartDisease(4)-	0.1527
	0.2150
	0.1552

Program 10: Locally Weighted Regression

```
from numpy import *
from os import listdir
import matplotlib
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np1
import numpy.linalg as np
from scipy.stats.stats import pearsonr

def kernel(point,xmat, k):
    m,n = np1.shape(xmat)
    weights = np1.mat(np1.eye((m)))
    for j in range(m):
        diff = point - X[j]
        weights[j,j] = np1.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 = np1.shape(xmat)
    ypred = np1.zeros(m)
    for i in range(m):
        ypred[i] = xmat[i]*localWeight(xmat[i],xmat,yamat,k)
    return ypred

#load data points
data = pd.read_csv('./input/tipsdata/tips.csv')
bill = np1.array(data.total_bill)
tip = np1.array(data.tip)

#preparing and add 1 in bill
mbill = np1.mat(bill)
mtip = np1.mat(tip)
# mat is used to convert to n dimensional to 2 dimensional array form m=
np1.shape(mbill)[1] # print(m) 244 data is stored in m
one = np1.mat(np1.ones(m))
X= np1.hstack((one.T,mbill.T)) # create a stack of bill from ONE
print(X)
```

```

#set k here
ypred = localWeightRegression(X,mtip,2)
SortIndex = X[:,1].argsort(0)

xsort = X[SortIndex][:,0]
fig = plt.figure()
ax = fig.add_subplot(1,1,1)
ax.scatter(bill,tip, color='blue')
ax.plot(xsort[:,1],ypred[SortIndex], color = 'red', linewidth=5)
plt.xlabel('Total bill')
plt.ylabel('Tip')
plt.show();

import numpy as np
from bokeh.plotting import figure, show, output_notebook
from bokeh.layouts import gridplot
from bokeh.io import push_notebook

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 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)]]))

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import matplotlib
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np1
import numpy.linalg as np
from scipy.stats.stats import pearsonr

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    m,n = np1.shape(xmat)
    weights = np1.mat(np1.eye((m)))
    for j in range(m):
        diff = point - X[j]
        weights[j,j] = np1.exp(diff*diff.T/(-2.0*k**2))
    return weights

def localWeight(point,xmat,yamat,k):
    wei = kernel(point,xmat,k)
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    W

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    m,n = np1.shape(xmat)
    ypred = np1.zeros(m)
    for i in range(m):
        ypred[i] = xmat[i]*localWeight(xmat[i],xmat,yamat,k)

```

```

return ypred

# load data points
data = pd.read_csv('../input/tipsdata/tips.csv')
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tip = np1.array(data.tip)

#preparing and add 1 in bill
mbill = np1.mat(bill)
mtip = np1.mat(tip) # mat is used to convert to n dimesiona to 2 dimensional array form m=
np1.shape(mbill)[1]
# print(m) 244 data is stored in m
one = np1.mat(np1.ones(m))
X= np1.hstack((one.T,mbill.T)) # create a stack of bill from ONE
#print(X)
#set k here
ypred = localWeightRegression(X,mtip,0.3)
SortIndex = X[:,1].argsort(0)
xsort = X[SortIndex][:,0]

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

```


total_bill	tip	sex	smoker	day	time	size
16.99	1.01	Female	No	Sun	Dinner	2
10.34	1.66	Male	No	Sun	Dinner	3
21.01	3.5	Male	No	Sun	Dinner	3
23.68	3.31	Male	No	Sun	Dinner	2
24.59	3.61	Female	No	Sun	Dinner	4
25.29	4.71	Male	No	Sun	Dinner	4
8.77	2.0	Male	No	Sun	Dinner	2
26.88	3.12	Male	No	Sun	Dinner	4
15.04	1.96	Male	No	Sun	Dinner	2
14.78	3.23	Male	No	Sun	Dinner	2
10.27	1.71	Male	No	Sun	Dinner	2
35.26	5.0	Female	No	Sun	Dinner	4
15.42	1.57	Male	No	Sun	Dinner	2
18.43	3.0	Male	No	Sun	Dinner	4
14.83	3.02	Female	No	Sun	Dinner	2
21.58	3.92	Male	No	Sun	Dinner	2
10.33	1.67	Female	No	Sun	Dinner	3
16.29	3.71	Male	No	Sun	Dinner	3
16.97	3.5	Female	No	Sun	Dinner	3
20.65	3.35	Male	No	Sat	Dinner	3
17.92	4.08	Male	No	Sat	Dinner	2

