# VISVESVARAYA TECHNOLOGICAL UNIVERSITY

"JnanaSangama", Belgaum -590014, Karnataka.



# LAB REPORT on

# MACHINE LEARNING (20CS6PCMAL)

Submitted by

Ishan Bhandari (1BM19CS198)

in partial fulfillment for the award of the degree of BACHELOR OF ENGINEERING

in

COMPUTER SCIENCE AND ENGINEERING



B.M.S. COLLEGE OF ENGINEERING
(Autonomous Institution under VTU)
BENGALURU-560019
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# B. M. S. College of Engineering,

Bull Temple Road, Bangalore 560019
(Affiliated To Visvesvaraya Technological University, Belgaum)

Department of Computer Science and Engineering



#### **CERTIFICATE**

This is to certify that the Lab work entitled "MACHINE LEARNING" carried out by ISHAN BHANDARI (1BM19CS198), 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 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 a Machine Learning- (20CS6PCMAL) work prescribed for the said degree.

**Dr. Asha G R**Assistant Professor
Department of CSE
BMSCE, Bengaluru

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

```
import pandas as pd
import numpy as np
d=pd.read csv("data.csv")
print(d)
att=np.array(d)[:,:-1]
print(att)
tar=np.array(d)[:,-1]
print(tar)
def finds(att, tar):
  for i, val in enumerate(tar):
     if val == "yes":
       res=att[i].copy()
       break
  for i, val in enumerate(att):
     if tar[i] == "yes":
       for x in range (len(res)):
          if val[x] != res[x]:
             res[x] = "?"
          else:
             pass
  return res
 print(finds(att,tar))
                 return res
  In [10]:
             print(finds(att,tar))
            ['sunny' 'warm' '?' 'strong' '?' '?']
```

```
sky air temp humidity
                            wind water forecast enjoy sport
0 sunny
            warm
                   normal strong warm
                                           same
                                                        yes
1 sunny
                     high strong warm
            warm
                                           same
                                                        yes
2 rainy
            cold
                     high strong warm
                                         change
                                                        no
                     high strong cool
3 sunny
            warm
                                         change
                                                       yes
```

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.

```
import numpy as np
import pandas as pd
data= pd.read csv("data.csv")
concepts=np.array(data.iloc[:,0:-1])
target=np.array(data.iloc[:,-1])
def learn(concepts, target):
  specific h=concepts[0].copy()
  general h=[["?" for i in range(len(specific h))] for i in range(len(specific 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]=specific h[x]
     if target[i]=="no":
       for x in range(len(specific h)):
          if h[x]!=specific h[x]:
             general h[x][x]=specific h[x]
             general h[x][x]='?'
  indices=[i for i,val in enumerate(general h) if val==['?','?','?','?','?','?']]
  for i in indices:
     general h.remove(['?','?','?','?','?'])
  return specific h, general h
 s final, f final = learn(concepts, target)
 s final
 f final
```

```
In [37]: s_final
Out[37]: array(['sunny', 'warm', '?', 'strong', '?', '?'], dtype=object)

In [38]: f_final
Out[38]: [['sunny', '?', '?', '?', '?'], ['?', 'warm', '?', '?', '?', '?']]
```

	sky	air temp	humidity	wind	water	forecast	enjoy sport
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

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.

```
import pandas as pd
import math
import numpy as np
data = pd.read_csv("3-dataset.csv")
features=[feat for feat in data]
features.remove("answer")
class Node:
  def init (self):
     self.children=[]
     self.value=""
     self.isLeaf=False
     self.pred=""
def entropy(examples):
  pos=0.0
  neg=0.0
  for , row in examples.iterrows():
    if row["answer"]=="yes":
       pos+=1
     else:
       neg+=1
  if pos==0.0 or neg==0.0:
    return 0.0
  else:
     p=pos/(pos+neg)
    n=neg/(pos+neg)
     return -(p * math.log(p,2) + n * math.log(n,2))
def info gain(examples, attr):
  uniq = np.unique(examples[attr])
  gain=entropy(examples)
  for u in uniq:
     subdata=examples[examples[attr] == u]
     sub e = entropy(subdata)
     gain -=(float(len(subdata))/float(len(examples)))*sub e
  return gain
def ID3(examples, attrs):
```

```
root = Node()
  max gain = 0
  max feat = ""
  for feature in attrs:
     gain = info gain(examples, feature)
    if gain > max gain:
       max gain = gain
       max feat = feature
  root.value = max feat
  uniq = np.unique(examples[max feat])
  for u in uniq:
    subdata = examples[examples[max feat] == u]
    if entropy(subdata)==0.0:
       newNode = Node()
       newNode.isLeaf = True
       newNode.value = u
       newNode.pred = np.unique(subdata["answer"])
       root.children.append(newNode)
    else:
       dummyNode = Node()
       dummyNode.value = u
       new attrs = attrs.copy()
       new attrs.remove(max feat)
       child = ID3(subdata, new attrs)
       dummyNode.children.append(child)
       root.children.append(dummyNode)
  return root
def printTree(root: Node, depth=0):
  for i in range(depth):
    print("\t", end=" ")
  print(root.value, end=" ")
  if root.isLeaf:
    print("->", root.pred)
  print()
  for child in root.children:
    printTree(child, depth+1)
root=ID3(data, features)
printTree(root)
```

	outlook	temperature	humidity	wind	answer
0	sunny	hot	high	weak	no
1	sunny	hot	high	strong	no
2	overcast	hot	high	weak	yes
3	rain	mild	high	weak	yes
4	rain	cool	normal	weak	yes
5	rain	cool	normal	strong	no
6	overcast	cool	normal	strong	yes
7	sunny	mild	high	weak	no
8	sunny	cool	normal	weak	yes
9	rain	mild	normal	weak	yes
10	sunny	mild	normal	strong	yes
11	overcast	mild	high	strong	yes
12	overcast	hot	normal	weak	yes
13	rain	mild	high	strong	no

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

```
import pandas as pd
import numpy as np
from sklearn import linear model
import matplotlib.pyplot as plt
df = pd.read csv('/content/drive/MyDrive/Colab Notebooks/canada per capita income -
canada per capita income.csv')
df
%matplotlib inline
plt.xlabel('year')
plt.ylabel('income')
plt.scatter(df.year,df.income,color='red',marker='+')
new df = df.drop('income',axis='columns')
new df
income = df.income
income
reg = linear model.LinearRegression()
reg.fit(new df,income)
reg.predict([[2021]])
reg.coef
reg.intercept
reg.predict([[2020]])
plt.xlabel('year',fontsize=20)
plt.ylabel('income',fontsize=20)
plt.scatter(df.year,df.income,color='red',marker='+')
plt.plot(df.year,reg.predict(df[['year']]),color='blue')
```

```
reg.predict([[2021]])
               reg.coef_
               reg.intercept_
              /usr/local/lib/python3.7/dist-packages/sklearn/base.py:451: UserWarning: X does not have valid feature names, but LinearRegression was fitted with feature names
"X does not have valid feature names, but"
              -1632210.7578554575
In [ ]:
               reg.predict([[2020]])
              /usr/local/lib/python3.7/dist-packages/sklearn/base.py:451: UserWarning: X does not have valid feature names, but LinearRegression was fitted with feature names
"X does not have valid feature names, but"
array([41288.69409442])
               plt.xlabel('year',fontsize=20)
plt.ylabel('income',fontsize=20)
               plt.scatter(df.year,df.income,color='red',marker='+')
plt.plot(df.year,reg.predict(df[['year']]),color='blue')
Out[ ]: [<matplotlib.lines.Line2D at 0x7f1d402a2590>]
                    40000
                    30000
               income
                    20000
                    10000
                              1970
                                                                                              2010
                                                              1990
                                                                              2000
                                                                year
```

	year	income
0	1970	3399.299037
1	1971	3768.297935
2	1972	4251.175484
3	1973	4804.463248
4	1974	5576.514583
5	1975	5998.144346
6	1976	7062.131392
7	1977	7100.126170
8	1978	7247.967035
9	1979	7602.912681
10	1980	8355.968120
11	1981	9434.390652
12	1982	9619.438377

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

```
from sklearn.datasets import fetch 20newsgroups
data = fetch 20newsgroups()
data.target names
categories = ['talk.religion.misc', 'soc.religion.christian',
        'sci.space', 'comp.graphics']
train = fetch 20newsgroups(subset='train', categories=categories)
test = fetch 20newsgroups(subset='test', categories=categories)
print(train.data[5])
from sklearn.feature extraction.text import TfidfVectorizer
from sklearn.naive bayes import MultinomialNB
from sklearn.pipeline import make pipeline
model = make pipeline(TfidfVectorizer(), MultinomialNB())
model.fit(train.data, train.target)
labels = model.predict(test.data)
from sklearn.metrics import confusion matrix
import seaborn as sns
import matplotlib.pyplot as plt
mat = confusion matrix(test.target, labels)
sns.heatmap(mat.T, square=True, annot=True, fmt='d', cbar=False,
       xticklabels=train.target names, yticklabels=train.target names)
plt.xlabel('true label')
plt.ylabel('predicted label');
def predict category(s, train=train, model=model):
  pred = model.predict([s])
  return train.target names[pred[0]]
predict category('Rocket launch in 3 months')
  In [22]:
           predict category('Rocket launch in 3 months')
  Out[22]: 'sci.space'
```

```
def predict_category(s, train=train, model=model):
    pred = model.predict([s])
    return train.target_names[pred[0]]

predict_category('determining the screen resolution')

'comp.graphics'

predict_category('what is 650 cc?')

'rec.motorcycles'

predict_category('launching payload')

'sci.space'
```

```
['alt.atheism',
 'comp.graphics',
 'comp.os.ms-windows.misc',
 'comp.sys.ibm.pc.hardware',
 'comp.sys.mac.hardware',
 'comp.windows.x',
 'misc.forsale',
 'rec.autos',
 'rec.motorcycles',
 'rec.sport.baseball',
 'rec.sport.hockey',
 'sci.crypt',
 'sci.electronics',
 'sci.med',
 'sci.space',
 'soc.religion.christian',
 'talk.politics.guns',
 'talk.politics.mideast',
 'talk.politics.misc',
 'talk.religion.misc']
```

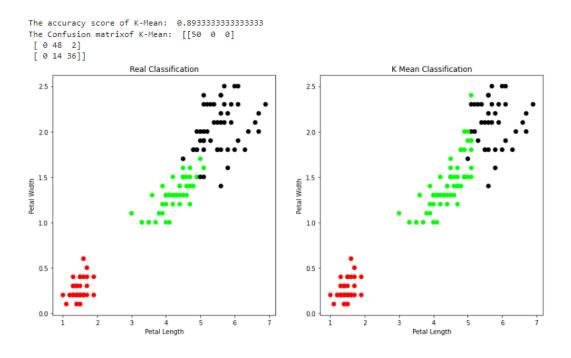
#### Apply k-Means algorithm to cluster a set of data stored in a .CSV file.

```
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)
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))
```



```
from sklearn import preprocessing
scaler = preprocessing.StandardScaler()
scaler.fit(X)
xsa = scaler.transform(X)
xs = pd.DataFrame(xsa, columns = X.columns)
#xs.sample(5)
```

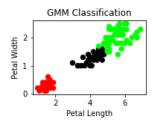
```
from sklearn.mixture import GaussianMixture
gmm = GaussianMixture(n_components=3)
gmm.fit(xs)

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')
```

Text(0, 0.5, '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 EM: 0.36666666666664
The Confusion matrix of EM: [[50 0 0]
  [ 0 5 45]
  [ 0 50 0]]
```

	Sepal_Length	Sepal_Width	Petal_Length	Petal_Width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

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

```
import numpy as np
import pandas as pd
import csv
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
heartDisease = pd.read csv('heart.csv')
heartDisease = heartDisease.replace('?',np.nan)
print('Sample instances from the dataset are given below')
print('\n Attributes and datatypes')
print(heartDisease.dtypes)
model=
BayesianModel([('age', 'heartdisease'), ('sex', 'heartdisease'), ('exang', 'heartdisease'), ('cp', 'heartdisease')
e'),('heartdisease','restecg'),('heartdisease','chol')])
print('\nLearning CPD using Maximum likelihood estimators')
model.fit(heartDisease,estimator=MaximumLikelihoodEstimator)
print('\n Inferencing with Bayesian Network:')
HeartDiseasetest infer = VariableElimination(model)
print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=HeartDiseasetest infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)
print('\n 2. Probability of HeartDisease given evidence= cp ')
g2=HeartDiseasetest infer.guery(variables=['heartdisease'],evidence={'cp':2})
print(q2)
```

```
model= BayesianModel([('age','heartdisease'),('sex','heartdisease'),('exang','heartdisease'),('cp','heartdi
 print('\nLearning CPD using Maximum likelihood estimators')
model.fit(heartDisease,estimator=MaximumLikelihoodEstimator)
 print('\n Inferencing with Bayesian Network:')
Learning CPD using Maximum likelihood estimators
 Inferencing with Bayesian Network:
HeartDiseasetest_infer = VariableElimination(model)
print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=HeartDiseasetest_infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)
Finding Elimination Order: : 100%
500.78it/s]
Eliminating: chol: 100%| 5/5 [00:00<00:00,
185.63it/s]
 1. Probability of HeartDisease given evidence= restecg
+----+
| heartdisease | phi(heartdisease) |
+========+
| heartdisease(0) | 0.1012 |
| heartdisease(1) | 0.0000 |
+----+
| heartdisease(2) |
                    0.2392
+----+
| heartdisease(3) |
                    0.2015
+----+
| heartdisease(4) | 0.4581 |
+----+
print('\n 2. Probability of HeartDisease given evidence= cp ')
q2=HeartDiseasetest infer.query(variables=['heartdisease'],evidence={'cp':2})
print(q2)
507.06it/s]
Eliminating: restecg: 100%
                                   5/5 [00:00<00:00,
179.06it/sl
2. Probability of HeartDisease given evidence= cp
+----+
| heartdisease | phi(heartdisease) |
+========+===++======++
| heartdisease(0) | 0.3610 |
+----+
| heartdisease(1) |
                 0.2159
| heartdisease(2) |
+-----
| heartdisease(3) | 0.1537 |
+----+
| heartdisease(4) | 0.1321 |
+----+
```

age	sex	ср	trestbps	chol	fbs	restecg	thalach	exang	oldpeak	slope	ca	thal	heartdisease
63			145	233	1	2	150	0	2.3		0	6	0
67		4	160	286	0		108		1.5	2			2
67		4	120	229	0	2	129		2.6	2	2		1
37			130	250	0	0	187	0	3.5		0		0
41		2	130	204	0	2	172	0	1.4		0		0
56		2	120	236	0	0	178	0	0.8		0		0
62	0	4	140	268	0	2	160	0	3.6		2		
57	0	4	120	354	0	0	163		0.6		0		0
63		4	130	254	0	2	147	0	1.4	2	1	7	2
53		4	140	203	1	2	155		3.1		0	7	1
57		4	140	192	0	0	148	0	0.4	2	0	6	0
56		2	140	294	0	2	153	0	1.3	2	0		0
56			130	256	1	2	142		0.6	2	1	6	2

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

```
from sklearn import datasets
from sklearn.cluster import KMeans
from sklearn.utils import shuffle
import numpy as np
import pandas as pd
iris=datasets.load iris()
X=iris.data
Y=iris.target
#Shuffle of Data
X,Y = \text{shuffle}(X,Y)
model=KMeans(n clusters=3,init='k-means++',max iter=10,n init=1,random state=3425)
model.fit(X)
# This is what KMeans thought (Prediction)
Y Pred=model.labels
from sklearn.metrics import confusion matrix
cm=confusion matrix(Y,Y Pred)
print(cm)
from sklearn.metrics import accuracy score
print(accuracy score(Y,Y Pred))
from sklearn.mixture import GaussianMixture
model2=GaussianMixture(n components=3,random state=3425)
#Training of the model
model2.fit(X)
Y predict2= model2.predict(X)
#Accuracy of EM Model
from sklearn.metrics import confusion matrix
cm=confusion_matrix(Y,Y_predict2)
print(cm)
from sklearn.metrics import accuracy score
```

# print(accuracy\_score(Y,Y\_predict2))

```
#Predicting classes for our data
Y_predict2= model2.predict(X)

#Accuracy of EM Model
from sklearn.metrics import confusion_matrix

cm=confusion_matrix(Y,Y_predict2)
print(cm)

from sklearn.metrics import accuracy_score
print(accuracy_score(Y,Y_predict2))

[[50 0 0]
[ 0 5 45]
[ 0 50 0]]
0.366666666666666664
```

				Petal_Width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8
 145 146 147 148	6.7 6.3 6.5 6.2	3.6  3.0 2.5 3.0 3.4	5.2 5.0 5.2 5.4	2 1. 2. 2.

from sklearn.model selection import train test split

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

```
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,\text{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))
 y_pred=classier.predict(x_test)
 print('confusion matrix')
 print(confusion_matrix(y_test,y_pred))
confusion matrix
[[16 0 0]
 [ 0 12 1]
 [0 0 16]]
 print('accuracy')
 print(classification_report(y_test,y_pred))
accuracy
            precision recall f1-score support
                1.00 1.00
                                 1.00
                1.00 0.92 0.96
                                           13
         1
                0.94 1.00 0.97
                                           16
                                0.98
   accuracy
   macro avg 0.98 0.97
ighted avg 0.98 0.98
                                 0.98
                                           45
weighted avg
                                 0.98
                                           45
```

	Sepal_Length	Sepal_Width	Petal_Length	Petal_Width
0	5.1	3.5	1.4	0.2
1	4.9	3.0	1.4	0.2
2	4.7	3.2	1.3	0.2
3	4.6	3.1	1.5	0.2
4	5.0	3.6	1.4	0.2
145	6.7	3.0	5.2	2.3
146	6.3	2.5	5.0	1.9
147	6.5	3.0	5.2	2.0
148	6.2	3.4	5.4	2.3
149	5.9	3.0	5.1	1.8

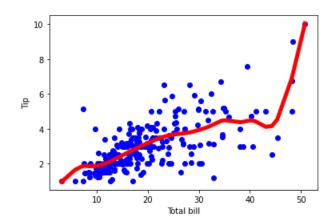
Implement the non-parametric Locally Weighted Regression algorithm in order to fit data points. Select the appropriate data set for your experiment and draw graphs.

```
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 i in range(m):
    diff = point - X[i]
     weights[j,j] = np1.exp(diff*diff.T/(-2.0*k**2))
  return weights
def localWeight(point,xmat,ymat,k):
  wei = kernel(point,xmat,k)
  W = (X.T*(wei*X)).I*(X.T*(wei*ymat.T))
  return W
def localWeightRegression(xmat,ymat,k):
  m,n = np1.shape(xmat)
  ypred = np1.zeros(m)
  for i in range(m):
    ypred[i] = xmat[i]*localWeight(xmat[i],xmat,ymat,k)
  return ypred
data = pd.read csv('tips.csv')
bill = np1.array(data.total bill)
tip = np1.array(data.tip)
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]
one = np1.mat(np1.ones(m))
X= np1.hstack((one.T,mbill.T)) # create a stack of bill from ONE
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):
  x0 = np.r [1, x0]
  X = np.c [np.ones(len(X)), X]
  xw = X.T * radial kernel(x0, X, tau)
  beta = np.linalg.pinv(xw @ X) @ xw @ Y
  return x0 @ beta
def radial kernel(x0, X, tau):
  return np.exp(np.sum((X - x0) ** 2, axis=1) / (-2 * tau * tau))
n = 1000
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])
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 = [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)]]))
```



```
def plot_lwr(tau):
    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)]]))
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 :
 [-2.98256634 -2.99368144 -3.05914505 -3.03174286 -3.07963801 -2.85954046
 -2.92988067 -2.958209 -2.96962333]
 Xo Domain Space(10 Samples) :
 [-2.97993311 -2.95986622 -2.93979933 -2.91973244 -2.89966555 -2.87959866
 -2.85953177 -2.83946488 -2.81939799]
```

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