ML Lab Record

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Lab 1: Find S Algorithm

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

Dataset:

| 1 | Weather | Temprature | Humidity | Goes |
|---|---------|------------|----------|------|
| 2 | Sunny | Warm | Mild | Yes |
| 3 | Rainy | Cold | Mild | No |
| 4 | Sunny | Moderate | Normal | Yes |
| 5 | Sunny | Cold | High | Yes |

```
import csv
a = []
with open('edata.csv', 'r') as csvfile:
    for row in csv.reader(csvfile):
         a.append(row)
    print(a)
print("\n The total number of training instances are : ",len(a))
num_attribute = len(a[0])-1
print("\n The initial hypothesis is : ")
hypothesis = ['0']*num_attribute
print(hypothesis)
for i in range(0, len(a)):
     if a[i][num_attribute] == 'positive':
        for j in range(0, num_attribute):
             if hypothesis[j] == '0' or hypothesis[j] == a[i][j]:
                hypothesis[j] = a[i][j]
             else:
                 hypothesis[j] = '?'
        print("\n The hypothesis for the training instance {} is : \n" .format(i+1),hypoth
esis)
print("\n The Maximally specific hypothesis for the training instance is ")
print(hypothesis)
```

```
### Superdiction of the properties of the properties of the training instance is in the properties of the training instance is into the training instanc
```

Lab 2: Candidate Elimination Algorithm

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.

| 1 | sky | airtemp | humidity | wind | water | forcast | enjoysport |
|---|-------|---------|----------|--------|-------|---------|------------|
| 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 |

```
import numpy as np
import pandas as pd
data = pd.DataFrame(data=pd.read_csv('edata.csv'))
concepts = np.array(data.iloc[:,0:-1])
print(concepts)
target = np.array(data.iloc[:,-1])
print(target)
def learn(concepts, target):
    specific_h = concepts[0].copy()
    print("initialization of specific_h and general h")
    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
s_final, g_final = learn(concepts, target)
print("Final Specific_h:", s_final, sep="\n")
print("Final General_h:", g_final, sep="\n")
```

Lab 3: Decision Tree

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.

Dataset:

| 1 | sky | airtemp | humidity | wind | water | forcast | enjoysport |
|---|-------|---------|----------|--------|-------|---------|------------|
| 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 |

```
import math
import csv

def load_csv(filename):
    lines = csv.reader(open(filename, "r"))
    dataset = list(lines)
    headers = dataset.pop(0)
    return dataset, headers
```

```
class Node:
    def __init__(self, attribute):
        self.attribute = attribute
        self.children = []
        self.answer = ""
def subtables(data, col, delete):
    dic = \{\}
    coldata = [row[col] for row in data]
    attr = list(set(coldata))
    counts = [0] * len(attr)
    r = len(data)
    c = len(data[0])
    for x in range(len(attr)):
        for y in range(r):
            if data[y][col] == attr[x]:
                counts[x] += 1
    for x in range(len(attr)):
        dic[attr[x]] = [[0 for i in range(c)] for j in range(counts[x])]
        pos = 0
        for y in range(r):
            if data[y][col] == attr[x]:
                if delete:
                    del data[y][col]
                dic[attr[x]][pos] = data[y]
                pos += 1
    return attr, dic
def entropy(S):
    attr = list(set(S))
    if len(attr) == 1:
        return 0
    counts = [0, 0]
    for i in range(2):
        counts[i] = sum([1 for x in S if attr[i] == x]) / (len(S) * 1.0)
    sums = 0
    for cnt in counts:
        sums += -1 * cnt * math.log(cnt, 2)
    return sums
def compute_gain(data, col):
    attr, dic = subtables(data, col, delete=False)
    total_size = len(data)
    entropies = [0] * len(attr)
    ratio = [0] * len(attr)
```

```
total entropy = entropy([row[-1] for row in data])
    for x in range(len(attr)):
        ratio[x] = len(dic[attr[x]]) / (total_size * 1.0)
        entropies[x] = entropy([row[-1] for row in dic[attr[x]]])
        total_entropy -= ratio[x] * entropies[x]
    return total entropy
def build_tree(data, features):
    lastcol = [row[-1] for row in data]
    if (len(set(lastcol))) == 1:
        node = Node("")
        node.answer = lastcol[0]
        return node
   n = len(data[0]) - 1
   gains = [0] * n
   for col in range(n):
        gains[col] = compute_gain(data, col)
    split = gains.index(max(gains))
   node = Node(features[split])
   fea = features[:split] + features[split + 1:]
    attr, dic = subtables(data, split, delete=True)
    for x in range(len(attr)):
        child = build_tree(dic[attr[x]], fea)
        node.children.append((attr[x], child))
    return node
def print_tree(node, level):
    if node.answer != "":
        print("---" * level, node.answer)
        return
    print("---" * level, node.attribute)
    for value, n in node.children:
        print("---" * (level + 1), value)
        print_tree(n, level + 2)
def classify(node, x_test, features):
    if node.answer != "":
        print(node.answer)
   pos = features.index(node.attribute)
   for value, n in node.children:
        if x_test[pos] == value:
            classify(n, x_test, features)
'''Main Program'''
dataset, features = load_csv("edata.csv")
model = build_tree(dataset, features)
```

```
print("-----THE DECISION TREE-----")
print_tree(model, 0)
testdata, features = load_csv("test.csv")
for xtest in testdata:
    print("-----test instance: ", xtest)
    print("-----label for test instance: ", end=" ")
    classify(model, xtest, features)
```

```
attr, dic = subtables(data, split, delete=True)
                                                                                                                       agred@ROG-ZEPHYRUS:/mnt/c/Users/sagred/Desktop/War/College/6/ML/LAB-3$ ls
               x in range(len(attr)):
child = build_tree(dic[attr[x]], fea)
node.children.append((attr[x], child))
                                                                                                                     sagred@ROG-ZEPHYRUS:/mnt/c/Users/sagred/Desktop/War/College/6/ML/LAB-3$ python3 dec-tree.py
--------THE DECISION TREE--------
                                                                                                                      Outlook
         return node
                                                                                                                      Outlook
--- sunny
---- Humidity
----- normal
----- yes
----- high
def print_tree(node, level):
    if node.answer != "":
        print("---" * level, node.answer)
                                                                                                                      --- overcast
----- yes
       print("---" * level, node.attribute)
for value, n in node.children:
    print("---" * (level + 1), value)
    print_tree(n, level + 2)
                                                                                                                      --- rain
----- Wind
   lef classify(node, x_test, features):
    if node.answer != "":
        print(node.answer)
                                                                                                                     ------ yes
------- test instance: ['rain', 'cool', 'normal', 'strong']
------label for test instance: no
------test instance: ['sunny', 'mild', 'normal', 'strong']
------label for test instance: yes
        pos = features.index(node.attribute)
                                                                                                                    ------label for test instance: yes sagred@ROG-ZEPHYRUS:/mnt/c/Users/sagred/Desktop/War/College/6/ML/LAB-3$
          for value, n in node.children:
    if x_test[pos] == value:
        classify(n, x_test, features)
dataset, features = load_csv("data.csv'
model = build_tree(dataset, features)
classify(model, xtest, features)
```

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

Naïve Bayesian Classifier Example 1:

| 1 | num_preg | glucose_conc | diastolic_bp | thickness | insulin | bmi | diab_pred | age | diabetes |
|----|----------|--------------|--------------|-----------|---------|------|-----------|-----|----------|
| 2 | 6 | 148 | 72 | 35 | 0 | 33.6 | 0.627 | 50 | 1 |
| 3 | 1 | 85 | 66 | 29 | 0 | 26.6 | 0.351 | 31 | 0 |
| 4 | 8 | 183 | 64 | 0 | 0 | 23.3 | 0.672 | 32 | 1 |
| 5 | 1 | 89 | 66 | 23 | 94 | 28.1 | 0.167 | 21 | 0 |
| 6 | 0 | 137 | 40 | 35 | 168 | 43.1 | 2.288 | 33 | 1 |
| 7 | 5 | 116 | 74 | 0 | 0 | 25.6 | 0.201 | 30 | 0 |
| 8 | 3 | 78 | 50 | 32 | 88 | 31 | 0.248 | 26 | 1 |
| 9 | 10 | 115 | 0 | 0 | 0 | 35.3 | 0.134 | 29 | 0 |
| 10 | 2 | 197 | 70 | 45 | 543 | 30.5 | 0.158 | 53 | 1 |
| 11 | 8 | 125 | 96 | 0 | 0 | 0 | 0.232 | 54 | 1 |
| 12 | 4 | 110 | 92 | 0 | 0 | 37.6 | 0.191 | 30 | 0 |
| 13 | 10 | 168 | 74 | 0 | 0 | 38 | 0.537 | 34 | 1 |
| 14 | 10 | 139 | 80 | 0 | 0 | 27.1 | 1.441 | 57 | 0 |
| 15 | 1 | 189 | 60 | 23 | 846 | 30.1 | 0.398 | 59 | 1 |
| 16 | 5 | 166 | 72 | 19 | 175 | 25.8 | 0.587 | 51 | 1 |
| 17 | 7 | 100 | 0 | 0 | 0 | 30 | 0.484 | 32 | 1 |
| 18 | 0 | 118 | 84 | 47 | 230 | 45.8 | 0.551 | 31 | 1 |
| 19 | 7 | 107 | 74 | 0 | 0 | 29.6 | 0.254 | 31 | 1 |
| 20 | 1 | 103 | 30 | 38 | 83 | 43.3 | 0.183 | 33 | 0 |
| 21 | 1 | 115 | 70 | 30 | 96 | 34.6 | 0.529 | 32 | 1 |
| 22 | 3 | 126 | 88 | 41 | 235 | 39.3 | 0.704 | 27 | 0 |
| 23 | 8 | 99 | 84 | 0 | 0 | 35.4 | 0.388 | 50 | 0 |
| 24 | 7 | 196 | 90 | 0 | 0 | 39.8 | 0.451 | 41 | 1 |
| 25 | 9 | 119 | 80 | 35 | 0 | 29 | 0.263 | 29 | 1 |
| 26 | 11 | 143 | 94 | 33 | 146 | 36.6 | 0.254 | 51 | 1 |
| 27 | 10 | 125 | 70 | 26 | 115 | 31.1 | 0.205 | 41 | 1 |
| 28 | 7 | 147 | 76 | 0 | 0 | 39.4 | 0.257 | 43 | 1 |
| 29 | 1 | 97 | 66 | 15 | 140 | 23.2 | 0.487 | 22 | 0 |
| 30 | 12 | 1/15 | 22 | 10 | 110 | 22.2 | 0.245 | 57 | 0 |

```
import csv
import random
import math
def loadcsv(filename):
    lines = csv.reader(open(filename, "r"))
    dataset = list(lines)
    for i in range(len(dataset)):
        dataset[i] = [float(x) for x in dataset[i]]
    return dataset
def splitdataset(dataset, splitratio):
    trainsize = int(len(dataset) * splitratio)
    trainset = []
    copy = list(dataset);
    while len(trainset) < trainsize:</pre>
        index = random.randrange(len(copy));
        trainset.append(copy.pop(index))
    return [trainset, copy]
def separatebyclass(dataset):
    separated = {} #dictionary of classes 1 and 0
    for i in range(len(dataset)):
        vector = dataset[i]
        if (vector[-1] not in separated):
            separated[vector[-1]] = []
        separated[vector[-1]].append(vector)
    return separated
def mean(numbers):
    return sum(numbers)/float(len(numbers))
def stdev(numbers):
    avg = mean(numbers)
    variance = sum([pow(x-avg,2) for x in numbers])/float(len(numbers)-1)
    return math.sqrt(variance)
def summarize(dataset): #creates a dictionary of classes
    summaries = [(mean(attribute), stdev(attribute)) for attribute in zip(*dataset)]
    del summaries[-1] #excluding labels +ve or -ve
    return summaries
def summarizebvclass(dataset):
```

```
separated = separatebyclass(dataset);
    summaries = {}
    for classvalue, instances in separated.items():
        summaries[classvalue] = summarize(instances) #summarize is used to cal to mean and
    return summaries
def calculateprobability(x, mean, stdev):
    exponent = math.exp(-(math.pow(x-mean,2)/(2*math.pow(stdev,2))))
    return (1 / (math.sqrt(2*math.pi) * stdev)) * exponent
def calculateclassprobabilities(summaries, inputvector):
    probabilities = {} # probabilities contains the all prob of all class of test data
    for classvalue, classsummaries in summaries.items():#class and attribute information a
       probabilities[classvalue] = 1
        for i in range(len(classsummaries)):
            mean, stdev = classsummaries[i] #take mean and sd of every attribute for class
            x = inputvector[i] #testvector's first attribute
            probabilities[classvalue] *= calculateprobability(x, mean, stdev); #use normal
   return probabilities
def predict(summaries, inputvector): #training and test data is passed
    probabilities = calculateclassprobabilities(summaries, inputvector)
    bestLabel, bestProb = None, -1
    for classvalue, probability in probabilities.items():#assigns that class which has he
        if bestLabel is None or probability > bestProb:
            bestProb = probability
            bestLabel = classvalue
    return bestLabel
def getpredictions(summaries, testset):
   predictions = []
   for i in range(len(testset)):
        result = predict(summaries, testset[i])
        predictions.append(result)
    return predictions
def getaccuracy(testset, predictions):
    correct = 0
    for i in range(len(testset)):
        if testset[i][-1] == predictions[i]:
            correct += 1
    return (correct/float(len(testset))) * 100.0
def main():
    filename = 'edata.csv'
    splitratio = 0.67
```

```
dataset = loadcsv(filename);

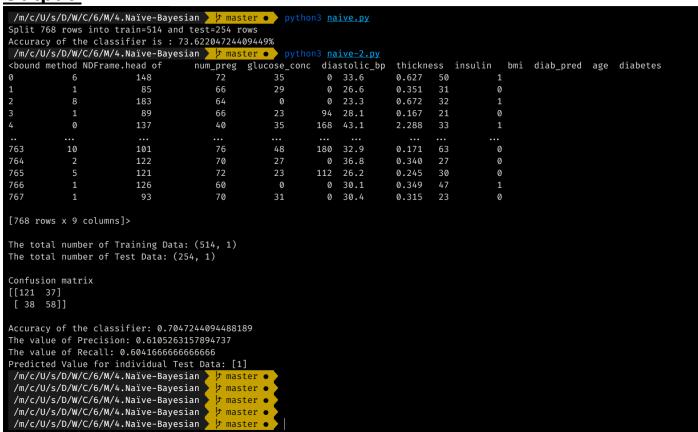
trainingset = dataset
  testset = [['sunny','cool','high','strong']]
  print('Split {0} rows into train={1} and test={2} rows'.format(len(dataset), len(train ingset), len(testset)))
  # prepare model
  summaries = summarizebyclass(trainingset);
  #print(summaries)
  # test model
  predictions = getpredictions(summaries, testset) #find the predictions of test data wi
th the training data
  accuracy = getaccuracy(testset, predictions)
  print('Accuracy of the classifier is : {0}%'.format(accuracy))
main()
```

Naïve Bayesian Classifier Example 2:

Dataset:

| 1 | day | outlook | temp | humidity | wind | play |
|----|-----|----------|------|----------|--------|------|
| 2 | D1 | Sunny | Hot | High | Weak | No |
| 3 | D2 | Sunny | Hot | High | Strong | No |
| 4 | D3 | Overcast | Hot | High | Weak | Yes |
| 5 | D4 | Rain | Mild | High | Weak | Yes |
| 6 | D5 | Rain | Cool | Normal | Weak | Yes |
| 7 | D6 | Rain | Cool | Normal | Strong | No |
| 8 | D7 | Overcast | Cool | Normal | Strong | Yes |
| 9 | D8 | Sunny | Mild | High | Weak | No |
| 10 | D9 | Sunny | Cool | Normal | Weak | Yes |
| 11 | D10 | Rain | Mild | Normal | Weak | Yes |
| 12 | D11 | Sunny | Mild | Normal | Strong | Yes |
| 13 | D12 | Overcast | Mild | High | Strong | Yes |
| 14 | D13 | Overcast | Hot | Normal | Weak | Yes |
| 15 | D14 | Rain | Mild | High | Strong | No |

```
import pandas as pd
import numpy as np
from sklearn.model selection import train test split
from sklearn.naive_bayes import GaussianNB
from sklearn import metrics
from sklearn import preprocessing
dataf = pd.read_csv("./edata.csv")
feature col names = ['outlook','temp','humidity','wind']
predicted_class_names = ['play']
def MultiLabelEncoder(columnlist,dataframe):
    for i in columnlist:
        labelencoder_X=preprocessing.LabelEncoder()
        dataframe[i]=labelencoder X.fit transform(dataframe[i])
    return dataframe
le = preprocessing.LabelEncoder()
feature_col = ['outlook','temp','humidity','wind','play']
Xdata = MultiLabelEncoder(feature_col,dataf)
X = Xdata[feature_col_names]
yy = dataf[predicted_class_names]
y = Xdata[predicted_class_names]
print(dataf.head)
xtrain,xtest,ytrain,ytest=train_test_split(X,y,test_size=0.33)
print ('\nThe total number of Training Data:',ytrain.shape)
print ('The total number of Test Data:',ytest.shape)
print(xtrain,ytrain)
classif = GaussianNB().fit(xtrain,ytrain)
print(classif)
predicted = classif.predict(xtest)
pri_enc = le.fit_transform(['sunny','cool','high','strong'])
predictTestData= classif.predict([pri_enc])
print('\nConfusion matrix')
print(metrics.confusion_matrix(ytest,predicted))
print('\nAccuracy of the classifier:',metrics.accuracy_score(ytest,predicted))
print('The value of Precision:', metrics.precision_score(ytest,predicted))
print('The value of Recall:', metrics.recall_score(ytest,predicted))
print("Predicted Value for individual Test Data:", predictTestData)
```



Lab 5: Bayesian Network

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

| 1 | age | sex | ср | trestbps | chol | fbs | restecg | thalach | exang | oldpeak | slope | ca | thal | heartdisease |
|----|-----|-----|----|----------|------|-----|---------|---------|-------|---------|-------|----|------|--------------|
| 2 | 63 | 1 | 1 | 145 | 233 | 1 | 2 | 150 | 0 | 2.3 | 3 | 0 | 6 | 0 |
| 3 | 67 | 1 | 4 | 160 | 286 | 0 | 2 | 108 | 1 | 1.5 | 2 | 3 | 3 | 2 |
| 4 | 67 | 1 | 4 | 120 | 229 | 0 | 2 | 129 | 1 | 2.6 | 2 | 2 | 7 | 1 |
| 5 | 37 | 1 | 3 | 130 | 250 | 0 | 0 | 187 | 0 | 3.5 | 3 | 0 | 3 | 0 |
| 6 | 41 | 0 | 2 | 130 | 204 | 0 | 2 | 172 | 0 | 1.4 | 1 | 0 | 3 | 0 |
| 7 | 56 | 1 | 2 | 120 | 236 | 0 | 0 | 178 | 0 | 0.8 | 1 | 0 | 3 | 0 |
| 8 | 62 | 0 | 4 | 140 | 268 | 0 | 2 | 160 | 0 | 3.6 | 3 | 2 | 3 | 3 |
| 9 | 57 | 0 | 4 | 120 | 354 | 0 | 0 | 163 | 1 | 0.6 | 1 | 0 | 3 | 0 |
| 10 | 63 | 1 | 4 | 130 | 254 | 0 | 2 | 147 | 0 | 1.4 | 2 | 1 | 7 | 2 |
| 11 | 53 | 1 | 4 | 140 | 203 | 1 | 2 | 155 | 1 | 3.1 | 3 | 0 | 7 | 1 |
| 12 | 57 | 1 | 4 | 140 | 192 | 0 | 0 | 148 | 0 | 0.4 | 2 | 0 | 6 | 0 |
| 13 | 56 | 0 | 2 | 140 | 294 | 0 | 2 | 153 | 0 | 1.3 | 2 | 0 | 3 | 0 |
| 14 | 56 | 1 | 3 | 130 | 256 | 1 | 2 | 142 | 1 | 0.6 | 2 | 1 | 6 | 2 |
| 15 | 44 | 1 | 2 | 120 | 263 | 0 | 0 | 173 | 0 | 0 | 1 | 0 | 7 | 0 |
| 16 | 52 | 1 | 3 | 172 | 199 | 1 | 0 | 162 | 0 | 0.5 | 1 | 0 | 7 | 0 |
| 17 | 57 | 1 | 3 | 150 | 168 | 0 | 0 | 174 | 0 | 1.6 | 1 | 0 | 3 | 0 |
| 18 | 48 | 1 | 2 | 110 | 229 | 0 | 0 | 168 | 0 | 1 | 3 | 0 | 7 | 1 |
| 19 | 54 | 1 | 4 | 140 | 239 | 0 | 0 | 160 | 0 | 1.2 | 1 | 0 | 3 | 0 |
| 20 | 48 | 0 | 3 | 130 | 275 | 0 | 0 | 139 | 0 | 0.2 | 1 | 0 | 3 | 0 |
| 21 | 49 | 1 | 2 | 130 | 266 | 0 | 0 | 171 | 0 | 0.6 | 1 | 0 | 3 | 0 |
| 22 | 64 | 1 | 1 | 110 | 211 | 0 | 2 | 144 | 1 | 1.8 | 2 | 0 | 3 | 0 |
| 23 | 58 | 0 | 1 | 150 | 283 | 1 | 2 | 162 | 0 | 1 | 1 | 0 | 3 | 0 |

```
import numpy as np # linear algebra
import pandas as pd # data processing, CSV file I/O (e.g. pd.read_csv)
from pgmpy.estimators import MaximumLikelihoodEstimator
from pgmpy.models import BayesianModel
from pgmpy.inference import VariableElimination
heart_Disease = pd.read_csv('./heart.csv')
heart_Disease = heart_Disease.replace('?',np.nan)

print('Sample instances from the dataset are given below')
print(heart_Disease.head())

print('\n Attributes and datatypes')
print(heart_Disease.dtypes)
```

```
model= BayesianModel([('age', 'heartdisease'),('sex', 'heartdisease'),('exang', 'heartdisease
'),('cp', 'heartdisease'),('heartdisease', 'restecg'),('heartdisease', 'chol')])
print('\nLearning CPD using Maximum likelihood estimators')
model.fit(heart_Disease, estimator=MaximumLikelihoodEstimator)

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

print('\n 1. Probability of HeartDisease given evidence= restecg')
q1=Heart_Disease_test_infer.query(variables=['heartdisease'],evidence={'restecg':1})
print(q1)

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

/m/c/U/s/D/W/C/6/M/5.Bayesian_Network ② ② master ● ② python3 Bayesian_Network.py Sample instances from the dataset are given below

age sex cp trestbps chol fbs restecg thalach exang oldpeak slope ca thal heartdisease

| 0 | 63 | 1 | 1 | 145 | 233 | 1 | 2 | 150 | 0 | 2.3 | 3 0 | 6 | 0 |
|---|----|---|---|-----|-----|---|---|-----|---|-----|-----|---|---|
| 1 | 67 | 1 | 4 | 160 | 286 | 0 | 2 | 108 | 1 | 1.5 | 2 3 | 3 | 2 |
| 2 | 67 | 1 | 4 | 120 | 229 | 0 | 2 | 129 | 1 | 2.6 | 2 2 | 7 | 1 |
| 3 | 37 | 1 | 3 | 130 | 250 | 0 | 0 | 187 | 0 | 3.5 | 3 0 | 3 | 0 |
| 4 | 41 | 0 | 2 | 130 | 204 | 0 | 2 | 172 | 0 | 1.4 | 1 0 | 3 | 0 |

Attributes and datatypes

age int64 int64 sex int64 trestbps int64 chol int64 int64 restecg int64 thalach int64 int64 exang oldpeak float64 slope int64 object ca object thal heartdisease int64

dtype: object

Learning CPD using Maximum likelihood estimators

Inferencing with Bayesian Network:

1. Probability of HeartDisease given evidence= restecg

Finding Elimination Order: : 100%

| Eliminating: cp: 100% | |
|---|--------------------------------|
| | 5/5 [00:00<00:00, 251.50it/s] |
| ++ | 0/5 [00:00 , ?it/s]</td |
| heartdisease phi(heartdisease) | |
| +=======+ | |
| heartdisease(0) 0.1012 | |
| ++ | |
| heartdisease(1) 0.0000 | |
| ++ | |
| heartdisease(2) 0.2392 | |
| <u>+</u> | |
| heartdisease(3) 0.2015 | |
| <u>+</u> | |
| heartdisease(4) 0.4581 | |
| + <u>+</u> | |
| 2. Probability of HeartDisease given evidence= cp | |
| Finding Elimination Order: : 100% | |
| | 5/5 [00:00<00:00, 3916.25it/s] |
| Eliminating: restecg: 100% | |
| | 5/5 [00:00<00:00, 487.52it/s] |
| ++ | 0/5 [00:00 , ?it/s]</td |
| heartdisease phi(heartdisease) | |
| +======+ | |
| heartdisease(0) 0.3610 | |
| <u>+</u> | |
| heartdisease(1) 0.2159 | |
| <u>+</u> | |
| heartdisease(2) 0.1373 | |
| | |
| <u>+</u> | |

| heartdisease(4) | 0.1321 |

Lab 6: K Means

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

| one | two |
|----------------------|---------------------|
| 0.22767982399693698 | 0.8582041480574577 |
| 0.9791882160551239 | 0.07715064988053028 |
| 0.504576604695406 | 0.5531144137299899 |
| 0.058132400743383585 | 0.52809798025712 |
| 0.7753430178214513 | 0.2179216898195512 |
| 0.5504238310550534 | 0.4708598154998745 |
| 0.04578653961976978 | 0.9185789498889001 |
| 0.5857699421693808 | 0.05803225463485838 |
| 0.7090721735923948 | 0.5818736617699065 |
| 0.018503930375096615 | 0.8865229185953829 |
| 0.8860650735174704 | 0.2395640162180548 |
| 0.6036387317795797 | 0.665583852216638 |
| 0.06942298413661226 | 0.858127672145648 |
| 0.6047387405995206 | 0.2781095847447219 |
| 0.7589891224957259 | 0.5120267751911348 |
| 0.08497507872469842 | 0.9911224601360906 |
| 0.6442858551230015 | 0.10730211335716414 |
| 0.6033125864818462 | 0.5364792610891768 |
| 0.0356326816753208 | 0.5874738213240243 |
| 0.0007170456636103 | 0.04656070001470005 |

```
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'])
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')
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 matrixof K-Mean: ',sm.confusion_matrix(y, model.labels_))
```

Lab 7: EM Algorithm

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

```
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'])
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))
```

```
/m/c/U/s/D/W/C/6/ML
/m/c/U/s/D/W/C/6/Ml
/m/c/U/s/D/W/C/6/M/6.K-Means
                                                                                           /m/c/U/s/D/W/C/6/ML
                                                                                                                                     7.EM-Algo/
                                                                                           /m/c/U/s/D/W/C/6/M/7.EM-Algo
/m/c/U/s/D/W/C/6/M/6.K-Means
                                                 python3 k-means.py
                                                                                          The accuracy score of EM: 0.966666666666667

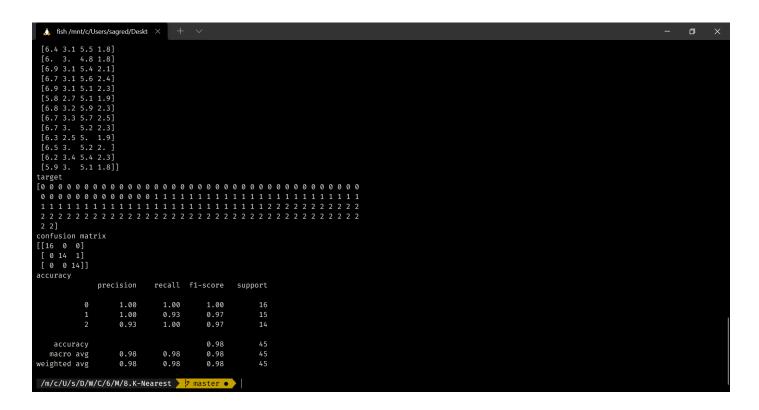
The Confusion matrix of EM: [[50 0 0]]
The accuracy score of K-Mean: 0.24
The Confusion matrixof K-Mean: [[ 0 50 0]
                                                                                           [ 0 45 5]
[ 0 0 50]]
/m/c/U/s/D/W/C/6/M/6.K-Means
                                                                                          The accuracy score of K-Mean: 0.09333333333333334
The Confusion matrixof K-Mean: [[ 0 50 0]
                                                                                                                                            python3 em.py
[ 2 0 48]
[36 0 14]]
                                                                                           [ 0 5 45]
[ 0 50 0]]
/m/c/U/s/D/W/C/6/M/7.EM-Algo <mark>} ♭</mark> π
 /m/c/U/s/D/W/C/6/M/6.K-Means // master // python3 k-means.py
The accuracy score of K-Mean: 0.24
The Confusion matrixof K-Mean: [[ 0 50 0]
                                                                                                                                            python3 em.py
                                                                                          The accuracy score of EM:
[48 0 2]
[14 0 36]]
                                                                                          The Confusion matrix of EM: [[50 0 0]
/m/c/U/s/D/W/C/6/M/6.K-Means
                                                                                           /m/c/U/s/D/W/C/6/M/7.EM-Algo > ├/> master ● >
```

Lab 8: K Nearest

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

```
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))
```

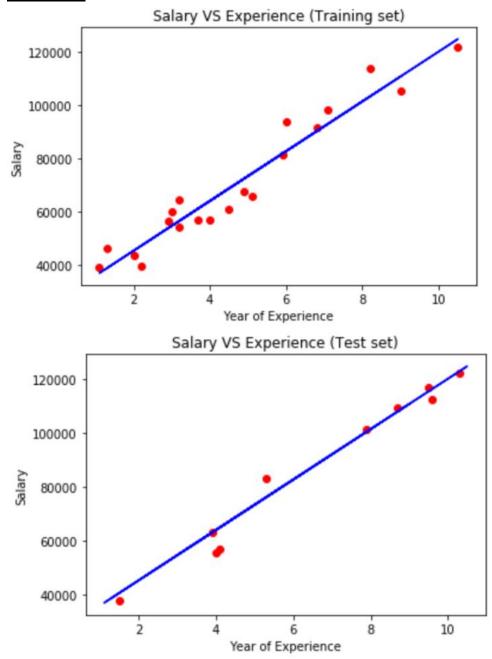


Lab 9: Linear Regression

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

| YearsExperience | Salary |
|-----------------|--------|
| 1.1 | 39343 |
| 1.3 | 46205 |
| 1.5 | 37731 |
| 2.0 | 43525 |
| 2.2 | 39891 |
| 2.9 | 56642 |
| 3.0 | 60150 |
| 3.2 | 54445 |
| 3.2 | 64445 |
| 3.7 | 57189 |
| 3.9 | 63218 |
| 4.0 | 55794 |
| 4.0 | 56957 |
| 4.1 | 57001 |

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
dataset = pd.read_csv('./data.csv')
X = dataset.iloc[:, :-1].values
y = dataset.iloc[:, 1].values
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)
from sklearn.linear_model import LinearRegression
regressor = LinearRegression()
regressor.fit(X_train, y_train)
y_pred = regressor.predict(X_test)
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()
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()
```



Lab 9: Locally Weighted Regression

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

| 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 02 | 2.02 | Fomalo | No | Sup | Dinner | 2 |

```
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,ymat,k):
wei = kernel(point,xmat,k)
W = (X.T*(wei*X)).I*(X.T*(wei*ymat.T))
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(∅)
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)]]))
```

