# ML Report Lab Test 1

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

```
import csv
a=[]
with open('data.csv') as dataset:
    for x in csv.reader(dataset):
        a.append(x)
a.remove(a[0])
msh = ['0']*(len(a[0])-1)
for x in a:
    if x[len(x)-1]=='yes' or x[len(x)-1]=='Yes':
        for i in range(0,len(msh)):
        if msh[i]=='0' or msh[i]==x[i]:
            msh[i]=x[i]
        else:
        msh[i]='?'
print(msh)
```

### **Output:**

```
Finds ×

To:\College\Lab 5th\ML\Lab 1\venv\Scripts\python.exe" "D:/College/Lab 5th/ML/Lab 1/Find s.py"

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

Process finished with exit code 0
```

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

```
import numpy as np
import pandas as pd
# Reading the data from CSV file
data = pd.read_csv('data.csv')
concepts = np.array(data.iloc[:,:-1])
print("\nInstances are:\n",concepts)
target = np.array(data.iloc[:,-1])
print("\nTarget Values are: ",target)
def train(concepts, target):
  # Initializing general and specific hypothesis
  specific_h = concepts[0].copy()
  print("\nInitialization of specific hypothesis and general hypothesis")
  print("\nSpecific Boundary: ", specific_h)
  general_h = [["?" for i in range(len(specific_h))] for i in range(len(specific_h))]
  print("\nGeneric Boundary: ",general_h)
  for i, val in enumerate(concepts):
     print("\nInstance", i+1 , "is ", val)
     #positive example
     if target[i] == "yes":
       print("Instance is Positive ")
       for x in range(len(specific_h)):
          if val[x]!= specific_h[x]:
            specific_h[x] = '?'
```

```
general_h[x][x] = '?'
     #negative example
     if target[i] == "no":
       print("Instance is Negative ")
       for x in range(len(specific_h)):
          if val[x]!= specific_h[x]:
             general_h[x][x] = specific_h[x]
          else:
             general_h[x][x] = '?'
     print("Specific Bundary after ", i+1, "Instance is ", specific_h)
     print("Generic Boundary after ", i+1, "Instance is ", general_h)
     print("\n")
  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 = train(concepts, target)
# displaying Specific_hypothesis
print("Final Specific_h: ", s_final, sep="\n")
# displaying Generalized_Hypothesis
print("Final General_h: ", g_final, sep="\n")
```

#### **Output:**

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.

```
import pandas as pd
import numpy as np
import math
data=pd.read_csv('/content/data set-1.csv');
attributes=[feat for feat in data]
attributes.remove('answer')
# print(features)
class Node:
 def __init__(self):
  self.children=[];
  self.isLeaf=False;
  self.value="";
  self.pred="";
def main():
 res=ID3(data,attributes)
 printTree(res)
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)
# This function creates the decision tree recursively
def ID3(data_set,attributes):
 root=Node()
 max_gain=0.0;
 max_feat="";
 # Comparitively find out which attribute gives us the maximum information
 for attribute in attributes:
  gain=info_gain(data_set,attribute)
  if gain>max_gain:
   max_gain=gain
   max feat=attribute
 # once we find the max gain, that will be the attribute which we use.
 root.value=max feat
 # All types of a particular attribute. Ex: In outlook, we have sunny, rain, overcast
 types=np.unique(data_set[max_feat])
 for t in types:
  # Get all instances which match a particular type
  subdata=data_set[data_set[max_feat]==t]
  # In case we find instances where we have only one type of data result (yes/no). Entropy will be zero
(Obviously!!)
  if entropy(subdata)==0.0:
   newNode=Node()
   newNode.isLeaf=True
   newNode.value=t
   newNode.pred=np.unique(subdata["answer"])
   root.children.append(newNode)
```

```
else:
   # If even one instance has different type of data result, we still cannot come to conclusion,
   # hence go to the next attribute and create the node and apply the same algorithm on the next attribute.
   dummyNode=Node()
   dummyNode.value=t
   new_attr=attributes.copy()
   # We can remove the current attribute, only when we have come to a conclusion
   # that we cannot decide with this attribute, we have gone to the next attribute. Hence we don't want to
come back.
   # + we may get stuck in cycle.
   new_attr.remove(max_feat)
   # Apply the algorithm on the next attribute with same current attributes which have been deleted.
   child=ID3(subdata,new_attr)
   dummyNode.children.append(child)
   root.children.append(dummyNode)
 return root
def info_gain(data_set,feature):
 types=np.unique(data_set[feature])
 # We are trying to get the entropy for the entire data_set we have taken into consideration.
 gain=entropy(data_set)
 for u in types:
  subdata=data_set[data_set[feature]==u]
  sub_e=entropy(subdata)
  gain-=(float(len(subdata))/float(len(data_set))*sub_e)
 return gain
def entropy(data):
 pos=0
 neg=0
```

# For the formula of entropy we need to see for how many of the +ve samples (yes) we have and how many -ve samples(no).

```
for _, row in data.iterrows():
    if row['answer'] == "yes":
        pos += 1
    else:
        neg += 1

if pos==0.0 or neg==0.0:
    return 0.0

p=pos/(pos+neg)
n=neg/(pos+neg)
return -(p*math.log(p,2)+n*math.log(n,2))
main()
```

#### **Output:**

file. Compute the accuracy of the classifier, considering few test data set import math import csv import random # This make sures that the dataset is in an ordered format. If we have some arbirary 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 # 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))

4. Write a program to implement the naïve Bayesian classifier for a sample training data set stored as a .CSV

```
# 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
# 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)
# We will calculte the mean and std dev with respect to each attribute. Important while calculating gaussian
probablity
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
# As explained earlier why e 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)
```

return train, test

```
for key, value in dictionary.items():
  # dictionary[key]=meanStdDev(value)
  info[key]=meanStdDev(value) #Here value stands for a complete group.
 return info
# 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
# 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] *= calculateGaussianProbablity(x, mean, std_dev)
 return probablities
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
# 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
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
filename="/content/bayes.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
Output:
Out[ ]: 60.91954022988506
```

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

```
import csv
import math
def calculate(X,Y):
 sum_x=sum(X)
 sum_y=sum(Y)
 n=len(Y)
 sum_xy=0
 for i in range(len(X)):
  sum_xy+=X[i]*Y[i]
 sum_x2=sum([x**2 for x in X])
 denomin=float((n*sum_x2)-(sum_x**2))
 # y=y_intercept+slope*x
 y_intercept=float((sum_y*sum_x2)-(sum_x*sum_xy))/denomin
 slope=float((n*sum_xy)-(sum_x*sum_y))/denomin
 return slope,y_intercept
filename='/content/insurance.csv'
file=open(filename)
dataset=csv.reader(file)
dataset=list(dataset)
X=[]
Y=[]
for x in dataset:
 X.append(x[3])
 Y.append(x[len(x)-1])
print(dataset[0])
x_tag=str(X[0])
y_tag=str(Y[0])
X=X[1:200]
Y=Y[1:200]
X=[float(x) for x in X]
```

```
Y=[float(y) for y in Y]
# print(Y)
slope,y_intercept=calculate(X,Y)
print(slope,y_intercept)
```

```
['age', 'sex', 'bmi', 'children', 'smoker', 'region', 'charges']
299.40712303629675 12768.55599860939
```

import matplotlib.pyplot as plt
plt.scatter(X,Y,marker='o')
plt.xlabel(x\_tag)
plt.ylabel(y\_tag)
plt.title('Simple Linear Regression')
y\_pred=[slope\*x+y\_intercept for x in X]
plt.plot(X,y\_pred,color='red')
plt.show()

