

Bharatiya Vidya Bhavan's Sardar Patel Institute of Technology

Bhavan's Campus, Munshi Nagar, Andheri (West), Mumbai-400058-India (Autonomous College Affiliated to University of Mumbai)

BE-ETRX B Sub- AIML Lab
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Name of the Experiment: Decision Tree (ID3) algorithm

Objective:

Write Python program to demonstrate the working of the decision tree based ID3 algorithm by using appropriate data set for building the decision tree and apply this knowledge to classify a new sample.

Outcomes:

- 1. Find entropy of data and follow steps of the algorithm to construct a tree.
- 2. Representation of hypothesis using decision tree.
- 3. Apply Decision Tree algorithm to classify the given data.
- 4. Interpret the output of Decision Tree.

System Requirements: Windows with MATLAB

Data Set Link: https://www.kaggle.com/code/kralmachine/analyzing-the-heart-

Dataset Description:

Number of Instances: 14

Number of Attributes (including the class attribute): 8

Attribute Information:

- Age (age in years)
- Sex (1 = male; 0 = female)
- Height (Below 150 = Low, 150 180 = Normal, Above ,180 = High)
- Weight (Below 40 = Low, 40 80 = Normal, Above 80 = Over, Above 120 = Obese)
- FPS (fasting blood sugar > 120 mg/dl) (1 = true; 0= false)
- TRESTBPS (resting blood pressure (in mm Hg on admission to the hospital))
- CHOL (serum cholesterol in mg/dl)
- ACTIVE(1or0)

Algorithm:

The decision tree builds classification or regression models in the form of a tree structure. It breaks down a dataset into smaller and smaller subsets while at the same time an associated decision tree is incrementally developed. The final result is a tree with decision nodes and leaf nodes. A decision node (e.g., Outlook) has two or more branches (e.g., Sunny, Overcast and Rainy). Leaf node (e.g., Play) represents a classification or decision. The topmost decision node in a tree which corresponds to the best predictor called root node. Decision trees can handle both categorical and numerical data.

Entropy:

A decision tree is built top-down from a root node and involves partitioning the data into subsets that contain instances with similar values (homogenous). ID3 algorithm uses entropy to calculate the homogeneity of a sample. If the sample is completely homogeneous the entropy is zero and if the sample is an equally divided it has entropy of one.

E(S) is the Entropy of the entire set, while the second term E(S, A) relates to an Entropy of an attribute A.



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$$E(S) = \sum_{x \in X} -P(x) \log_2 P(x) \quad E(S,A) = \sum_{x \in X} [P(x)*E(S)]$$

Information Gain:

The information gain is based on the decrease in entropy after a dataset is split on an attribute. Constructing a decision tree is all about finding attribute that returns the highest information gain (i.e., the most homogeneous branches).

$$IG(S,A) = E(S) - E(S,A)$$

Code:

```
import numpy as np
   import matplotlib.pyplot as plt
   dataset = pd.read_csv(r'C:\Users\Dell\Desktop\Shubham\SEM7\AIML\EXP3\cardio_train.csv')
   print(X)
   attribute = ['age', 'sex', 'height', 'weight', 'trtbps', 'bloodsugar', 'chol', 'active']
[['young' 'm' 'short' 'normal' 'normal' 'active' 'Yes']
 ['young' 'f' 'tall' 'normal' 'normal' 'active' 'Yes']
 ['old' 'f' 'average' 'normal' 'low' 'normal' 'not-active' 'Yes']
 ['adult' 'm' 'average' 'over' 'low' 'high' 'not-active' 'No']
 ['adult' 'f' 'short' 'over' 'low' 'high' 'not-active' 'No']
 ['adult' 'f' 'short' 'normal' 'normal' 'low' 'not-active' 'No']
 ['old' 'f' 'short' 'normal' 'high' 'high' 'active' 'No']
 ['young' 'm' 'tall' 'obese' 'high' 'normal' 'active' 'Yes']
 ['young' 'f' 'short' 'normal' 'low' 'normal' 'active' 'No']
 ['adult' 'f' 'average' 'obese' 'low' 'normal' 'not-active' 'No']
 ['young' 'f' 'average' 'obese' 'low' 'low' 'not-active' 'No']
 ['old' 'm' 'tall' 'normal' 'low' 'low' 'active' 'No']
 ['old' 'm' 'average' 'normal' 'low' 'low' 'not-active' 'No']
 ['adult' 'f' 'short' 'over' 'high' 'high' 'active' 'No']]
 class Node(object):
      def init (self):
            self.value = None
            self.decision = None
            self.childs = None
```



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```
def findEntropy(data, rows):
    yes = 0
    no = 0
    ans = -1
    idx = len(data[0]) - 1
    entropy = 0
    for i in rows:
        if data[i][idx] == 'Yes':
            yes = yes + 1
        else:
            no = no + 1

        x = yes/(yes+no)
        y = no/(yes+no)
        if x != 0 and y != 0:
            entropy = -1 * (x*math.log2(x) + y*math.log2(y))
        if x == 1:
            ans = 1
        if y == 1:
            ans = 0
        return entropy, ans
```

```
def findMaxGain(data, rows, columns):
   maxGain = 0
   retidx = -1
   entropy, ans = findEntropy(data, rows)
    if entropy == 0:
           print("No")"""
       return maxGain, retidx, ans
    for j in columns:
       mydict = {}
        for i in rows:
           key = data[i][idx]
           if key not in mydict:
               mydict[key] = 1
               mydict[key] = mydict[key] + 1
       gain = entropy
        for key in mydict:
            for k in rows:
                if data[k][j] == key:
                   if data[k][-1] == 'Yes':
                       yes = yes + 1
```



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```
x = yes/(yes+no)
           y = no/(yes+no)
              gain += (mydict[key] * (x*math.log2(x) + y*math.log2(y)))/14
        if gain > maxGain:
           maxGain = gain
   return maxGain, retidx, ans
def buildTree(data, rows, columns):
   maxGain, idx, ans = findMaxGain(X, rows, columns)
   if maxGain == 0:
       if ans == 1:
           root.value = 'No'
   root.value = attribute[idx]
   mydict = {}
       key = data[i][idx]
       if key not in mydict:
           mydict[key] = 1
           mydict[key] += 1
```

```
newcolumns = copy.deepcopy(columns)
   newcolumns.remove(idx)
   for key in mydict:
       newrows = []
       for i in rows:
          if data[i][idx] == key:
               newrows.append(i)
       temp = buildTree(data, newrows, newcolumns)
       temp.decision = key
       root.childs.append(temp)
   return root
def traverse(root):
   print(root.decision)
   print(root.value)
   n = len(root.childs)
   if n > 0:
           traverse(root.childs[i])
   columns = [i for i in range(0, 6)]
   root = buildTree(X, rows, columns)
   traverse(root)
calculate()
```



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Interpretation of output:

```
Person is fit ?
└─ BloodSugar
      - high
        L_ NO
        L- NO
        └─ BloodPressure
            ├─ high
                └─ YES
                └─ Age
                    ├─ adult
                       L__ NO
                       old
                         --- NO
                        L_ YES
                       young
               normal
                └─ YES
```

Conclusion:

- We learned how to make a decision tree out of the given Dataset
- We learned to identify the root node, leaf node and connecting node
- We learned to calculate the entropy of the dataset and information gain of each attribute to decide the root node and subsequently the leaf nodes