Artificial Intelligence Project 2 Decision Tree

Class 1, Team 6

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1 Team Members

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2 Problem Description

The dataset contains the records of the United States congressional votes in 1984. There are 435 records, each containing the party (goal attribute) of the congressman/congresswoman and his/her votes on 16 bills. There are only two types of values for the goal attribute: 'democrat' and 'republican', and three types for other attributes: 'n' (can be considered as nay), 'y' (can be considered as yea), and '?' (can be considered as missing).

Given this dataset, the goal is to build a decision tree to classify a record (that is, to find out the party of this congressman/congresswoman) according to its known attributes (that is, his/her votes on these 16 bills). Both the training set and the testing set will be sampled from this dataset.

3 Algorithm and Implementation

For this project we implemented the C4.5 algorithm to build the decision tree. Since the dataset only have two effective values ('y' and 'n') for all attributes, there is no need to use the information gain ratio as the measure of attributes. The information gain alone is more suitable for this dataset.

To handle the missing data, when building the tree, we only consider the records with defined value for the attribute being measured. When classifying the record, we use the probability of each possible value to weigh the calculation of each branch.

The data set is split by sampling the training set with a given probability p. The rest of the data will be put in to the testing set.

3.1 Decision tree building

The algorithm for building the decision tree is described in Algorithm 1.

Since the decision tree built with this data set won't be very deep (after all, there are only 16 attributes, and the data set is fairly skewed), a recursive implementation is sufficient. Records with missing values of the chosen attribute will be ignored when calculating the information gain.

Algorithm 1 Decision tree bulding

```
1: function Build(data, attributes)
       if data is empty then
 3:
           return an empty node
       end if
 4:
 5:
       gain_{best} = 0.0, attr_{best} = None
       value_{best} = None, sets_{best} = None
 6:
 7:
       for each attribute a in attributes do
           Divide data based on values of a into set1, set2
 8:
           Calculate the information gain of spliting data into set1 and set2, ignoring missing values
 9:
           if The new information gain > gain_{best} then
10:
11:
               Update gain_{best}, attr_{best}, sets_{best}, value_{best}
           end if
12:
       end for
13:
       if attributes is not empty and gain_{best} \neq 0.0 then
14:
           remove attr_{best} from attributes
15:
           left\_branch = Build(set1 insets_{best}, attributes)
16:
           right\_branch = Build(set2 insets_{best}, attributes)
17:
           Create a node node
18:
           node.attr = attr_{best}, node.value = value_{best}
19:
           node.left = left\_branch, node.right = right\_branch, node.count = size of data
20:
21:
           return node
22:
       else
           Create a node node
23:
           node.leaves = count of each parties in data
24:
           node.count = size of data
25:
           return node
26:
       end if
27:
28: end function
```

Remark 1.

The entropy of a attribute V in the data set D is defined as:

$$H(V, D) = -\sum_{k} P(v_k) \log_2 \frac{1}{P(v_k)}$$

where v_k are values of V in D, $P(v_k)$ is the probability of v_k in D.

Remark 2.

When a data set D with goal attribute V is split into subsets D_k using attribute A, each with n_k records, the *information gain* is defined as:

$$Gain(A) = H(V, D) - Remainder(A)$$

where

$$Remainder(A) = \sum_{k} P(n_k)H(V, D_k)$$

 $P(n_k)$ is the proportion of E_k in D.

3.2 Decision tree pruning

The algorithm for pruning the decision tree is described in Algorithm 2. For the same reason as the tree building, a recursive implementation is enough. Nonetheless, since the data set is quite skewed, there is seldom a need for pruning the tree.

Algorithm 2 Decision tree pruning

```
1: function Prune(tree, min_gain)
       if tree.left is not a leave then
           Prune(tree.left, min\_gain)
 3:
       end if
 4:
 5:
       if tree.right is not a leave then
           Prune(tree.right, min\_gain)
 6:
 7:
       end if
       if Both tree.left and tree.right are leaves then
 8:
           Temprorily combine tree.left and tree.right into new_node
 9:
           delta = \text{Entropy}(new\_node) - (\text{Entropy}(tree.left) + \text{Entropy}(tree.right))/2
10:
11:
           if delta < min_{-}qain then
              Replace tree with new_node
19.
           end if
13:
       end if
15: end function
```

3.3 Classification

The algorithm for classifying an observation is described in Algorithm 3. It follows all branches at any node for which a value is missing and multiply the weights along each path.

Algorithm 3 Classification

```
1: function Classify(tree, observation)
       if tree is a leave then
2:
3:
          return tree.leaves
       end if
4:
       value = observation[tree.attr]
5:
       if value is missing then
6:
          result\_left = Classify(tree.left, observation)
 7:
8:
          result\_right = Classify(tree.right, observation)
          weight\_left = tree.left.count/tree.count
9:
           weight\_right = tree.left.count/tree.count
10:
11:
           for each pair of goal, count in result_left do
12:
              result[goal] + = count * weight\_left
          end for
13:
           for each pair of goal, count in result_right do
14:
              result[goal] + = count * weight\_right
15:
          end for
16:
          return result
17:
18:
       else
          if value = tree.left.value then
19:
              return CLASSIFY(tree.left, observation)
20:
21:
22:
              return Classify(tree.right, observation)
          end if
23:
       end if
24:
25: end function
```

4 Experiment Result

To evaluate the implementation, we sample the data with probability p and use the sample as training set. The rest of the data set will be the testing set. One of the decision tree built with p=0.2 is visualized in Figure 1. In addition, we take training sets sampled with various p to generate the learning curve for evaluation, as shown in Figure 2. With training sets of different sizes, the precisions of cross validation fluctuate between 0.9 and 1.0, with mean $\mu=0.9462$ and standard variation $\sigma=0.0175$.

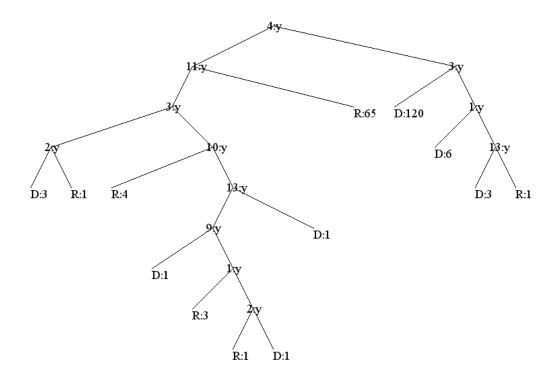


Figure 1: One of the decision trees built with training set sampled with p=0.5

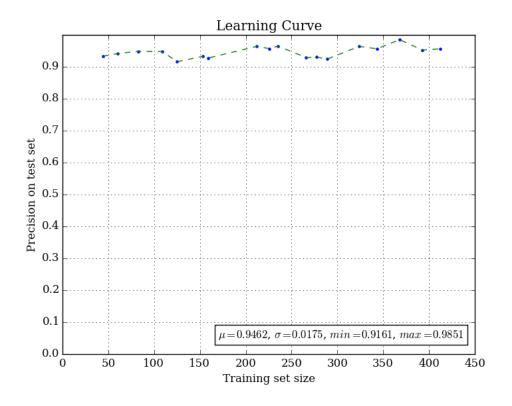


Figure 2: Learning Curve