

# INF264 Project 1

## IMPLEMENTING DECISION TREES

*Sophie Blum and Benjamin Friedl — 18.09.2020*

About the document: The names behind every function description point put, who initially wrote the code for this functionality. The largest amount of time however went into debugging and correcting the code, as well as re-evaluating the design choices and making the algorithm work. We did all this together, so that it is hard to point out, who did what exactly.

### 1 Implementation of the tree data structure (Sophie)

The decision tree is implemented as a node data structure in a semi-object-oriented way. A tree is viewed as a root with children connected to it. That means that every node can be the root of a tree (and every node is indeed the root of its subtree). As every node could be a tree, the decision tree is not encapsulated in a data structure itself and has to be given as a parameter for every function that uses it (see also: `learn()`) The nodes themselves are implemented in the node class, which contains variables for data and the majority label in the subtree denoted by this node. The data variable holds either the feature for the corresponding decision or a label, if the node is a leaf in the tree. The majority label is used for pruning. The node also has a reference to its parent and its children. The latter are organized in a list of tuples. Every tuple has a reference to the child node as well as the value that is used for the decision.

```
1  class node:
2      def __init__(self, data, parent = None,
3          majority_label = None):
4          self.data = data
5          self.parent = parent
6          self.majority_label = majority_label
7          self.children = []
8      def __str__(self):
9          return f"{self.data} - {self.children}"
10     def addChild(self, child):
11         self.children.append(child)
```

The `printTree()` is an inorder traversal of a tree denoted by a root-node. Remark: We designed the data structure before we knew we only had to compute binary trees, so this could have been implemented easier by just storing each child node separately and having only one variable to denote the decision value.

## 2 Implementation of learn (Sophie)

The `learn` function is splitted into to separate functions, as the integration of pruning is easier that way. In fact the `learn_rec()` has been `learn()` in the first place. As we implemented pruning and needed to integrate it, the easiest way was to split the function into two parts.

`Learn_rec()` is the powerhouse of the `learn` function and outputs the root node of the decision tree, that is learned on the input data.

It is important to note, that we considered a lot of different options for this function. We tried basically two different options concerning the decision tree:

1. On a given path in the tree one feature is used at most one time
2. A feature can be used multiple times to split on a given path in the tree

Both variants result in valid decision trees. Of course the decision tree that is return by the first one is a lot smaller, but it also has a significantly better accuracy. We chose accuracy over performance and therefore decided to allow a label to be used more than once. We know that this solution is also more prone to overfitting, but the use of pruning should reduce this risk.

The function first checks all base cases and outputs the majority label as a leaf.

```
1     if same_value(y):
2         return node(y[0])
3     elif len(X[0]) == len(used_feature_list) or same_value(X)
4         :
5         label = get_maj_label(y)
6         return node(label)
```

When there is no base case present, a feature gets selected based on the selected impurity measure (separately implemented). The dataset then gets splitted based on the mean of the selected feature. These separate datasets are then used for a recursive call of the same function. All the created nodes get assigned their respective parent and/or children. All this functionality is subdivided in different functions doing smaller tasks.

`Learn()` is just the packaging for the recursion and the pruning in one function. It splits the given dataset, trains a decision tree with `learn` and uses pruning afterwards. All of this according to the input, whether pruning is wanted or not.

```

1  def learn(X, y, impurity_measure, pruning):
2      seed = 432
3      if(pruning):
4          X_train, X_prun, Y_train, Y_prun =
model_selection.train_test_split(X, y, test_size= 0.3,
shuffle=True,
                                random_state = seed)
5      else:
6          X_train = X
7          Y_train = y
8          root = learn_rec(X_train, Y_train, [],
impurity_measure)
9      if pruning:
10         root = pruning_function(root, X_prun, Y_prun)
11     return root

```

### 3 Implementation of the impurity measures (Benjamin)

Both, entropy as well as the Gini-impurity, need for its calculation only the probabilities of different possible values. The latter is computed by the function “probability\_list” which takes a list of numbers and first counts the occurrence for all the different values. This is done through a for-loop over all elements in the input-list.

```

1  for i in range(count_values):
2      if (values[i] not in value_list):
3          value_list.append(values[i])
4          prob_list.append(1)
5          different_val +=1
6      else:
7          for j in range(different_val):
8              if(values[i]==value_list[j]): prob_list[j]+=1

```

If the value of the viewed element didn’t occur before, it gets appended to the value\_list while prob\_list saves the value 1 at the same index. Else the algorithm searches the index in the value\_list with the right value and increments the value in the prob\_list at the right index. To get the probabilities, every numbers of occurrence is divided by the total length of the input-list.

```

1  for k in range(different_val):
2      prob_list[k] = float(prob_list[k]) / count_values

```

In order to get the entropy and Gini-impurity, one can first compute this probability\_list and implement the summation with a simple for-loop.

```

1  def entropy(values):
2      prob_list = probability_list(values)
3      different_values = len(prob_list)
4      sum = 0

```

```

5         for i in range(different_values):
6             sum -= math.log(prob_list[i], 2) * prob_list[i]
7         return sum
8
9     def gini(values):
10         prob_list = probabiliy_list(values)
11         different_values = len(prob_list)
12         sum = 0
13         for i in range(different_values):
14             sum += prob_list[i]*(1-prob_list[i])
15         return sum

```

## 4 Implementation of the decrease in impurity through a split (Benjamin)

The value of decrease in impurity is computed by the function “information\_gain”, which takes a feature-vector  $x$ , a label-vector  $y$  and a string describing the impurity-measure as input and is implemented in three steps. First, the mean value of all values in  $x$  is calculated. This is done by a simple for-loop over all values in  $x$ .

```

1     mean = 0
2     count = 0
3     for i in range(len(x)):
4         mean += x[i]
5         count +=1
6     mean /= count

```

Secondly, the split of the label-vector is realised in two lists “leq\_mean” and “g\_mean” containing all values from  $x$  less equal or greater than the mean. Additionally, the percentual occurrence of the values in “leq\_mean” is computed in “prob\_leq\_mean”.

```

1     leq_mean = []
2     g_mean = []
3     count_leq_mean = 0
4     for i in range(len(x)):
5         if(x[i] <= mean):
6             count_leq_mean +=1
7             leq_mean.append(y[i])
8         else:
9             g_mean.append(y[i])
10    prob_leq_mean = count_leq_mean / count

```

Lastly the Impurities before and after the split are calculated using the already implemented functions “entropy” or “gini”.

```

1     f(impurity_measure=="entropy"):
2         impurity = entropy(y)
3         conditional_impurity = prob_leq_mean * entropy(
leq_mean)+ (1-prob_leq_mean) * entropy(g_mean)

```

```

4     elif(impurity_measure=="gini"):
5         impurity = gini(y)
6         conditional_impurity = prob_leq_mean * gini(leq_mean)
7         + (1-prob_leq_mean) * gini(g_mean)
8     else:
9         print("Not known impurity measure")

```

Using that, the function returns the value of decrease in impurity through the split, which is the information gain in the case of entropy.

```

1     return impurity - conditional_impurity

```

## 5 Implementation of predict (Benjamin)

This function gets as parameter a vector  $x$  and returns the predicted label. Generally, it is implemented as recursion starting with the root and then continuing with the child for which the value of  $x$  at the looked at feature fits. If the (sub-)tree is a leaf, that is when the (sub-)tree's children-list is empty, then the function returns the label memorized in `node.data`.

```

1     if (not node.children):
2         return node.data

```

Otherwise, a for loop goes through the children-list, which is sorted by the value of split. In the case of a binary split, there is only one split point, so in `children[0][1]` and `children[1][1]` there is both times memorized the mean-value. Furthermore, if the value of the feature-vector at the splitting feature, which memorized as index in `node.data` for not-leaf nodes, is less than the mean value, then we recursively call predict for the child-node memorized in `children[i][0]`.

```

1     if(x[node.data] < node.children[i][1]):
2         return predict(node.children[i][0], x)

```

If we reached the end of the children-list, it means the vector at the looked upon feature has a higher value than every splitting point and we recursively call predict for the last child-node.

```

1     elif(i==count_values-1):
2         return predict(node.children[i][0], x)

```

Remark: We implemented this function before we knew we only had to compute binary trees, so this could have been implemented easier by just looking at the left and right child.

## 6 Implementation of pruning (Benjamin)

The pruning-function takes as input a node representing a (sub-)tree and the pruning data `X_prun` and `y_prun`. It returns a node representing the post-

pruned input-tree. First, the pruning function searches the leafs of the decision tree. Therefore, it recursively calls itself until the length of the children-list is not zero.

```

1     if (len(node.children)!=0):
2         for i in range(len(node.children)):
3             #call recursion on each child
4             pruning_function((node.children[i])[0],
X_prun, y_prun)
5     else:
6         #reached leaf
7         return node

```

Afterwards, it goes back up the tree and checks for every node whether the accuracy of the (sub-)tree on the pruning-data is lower than the accuracy when predicting the majority label. In this case, we change the node into a leaf predicting the majority label.

```

1     if (acc(node, X_prun, y_prun) < majority_label_acc(node.
majority_label, y_prun)):
2         node.data = node.majority_label
3         node.children = []
4     return node

```

The accuracies are implemented through a simple for-loop counting the right predictions.

```

1     def majority_label_acc(majority_label, y_prun):
2         x = 0
3         for i in range(len(y_prun)):
4             if(y_prun[i]==majority_label):
5                 x += 1
6         return float(x) / len(y_prun)
7
8     def acc(node, X, Y):
9         x = 0
10        for i in range(len(Y)):
11            if(Y[i]==predict(node, X[i])):
12                x += 1
13        return float(x) / len(Y)

```

## 7 Evaluation (Benjamin, Sophie)

First, we split the Data into Training-, Evaluation and Test-data.

```

1     seed = 111
2     X_train, X_val_test, Y_train, Y_val_test =
model_selection.train_test_split(X, y, test_size= 0.3,
shuffle=True, random_state = seed)
3     seed = 112
4     X_val, X_test, Y_val, Y_test = model_selection.
train_test_split(X_val_test, Y_val_test, test_size= 0.5,
shuffle=True, random_state = seed)

```

We chose a ratio of 0.7, 0.15, 0.15 to have sufficient training-data whilst getting representative accuracies. Afterwards we implemented four trees, one for each impurity measure with and without pruning and computed the accuracy on the training- as well as validation-data. This example shows one the procedure for the tree using entropy and pruning.

```
1 tree_ent_prun = learn(X_train, Y_train, impurity_measure=  
    "entropy", pruning=True)  
2 val_acc_ent_prun = acc(tree_ent_prun, X_val, Y_val)  
3 train_acc_ent_prun = acc(tree_ent_prun, X_train, Y_train)
```

We got the following results:

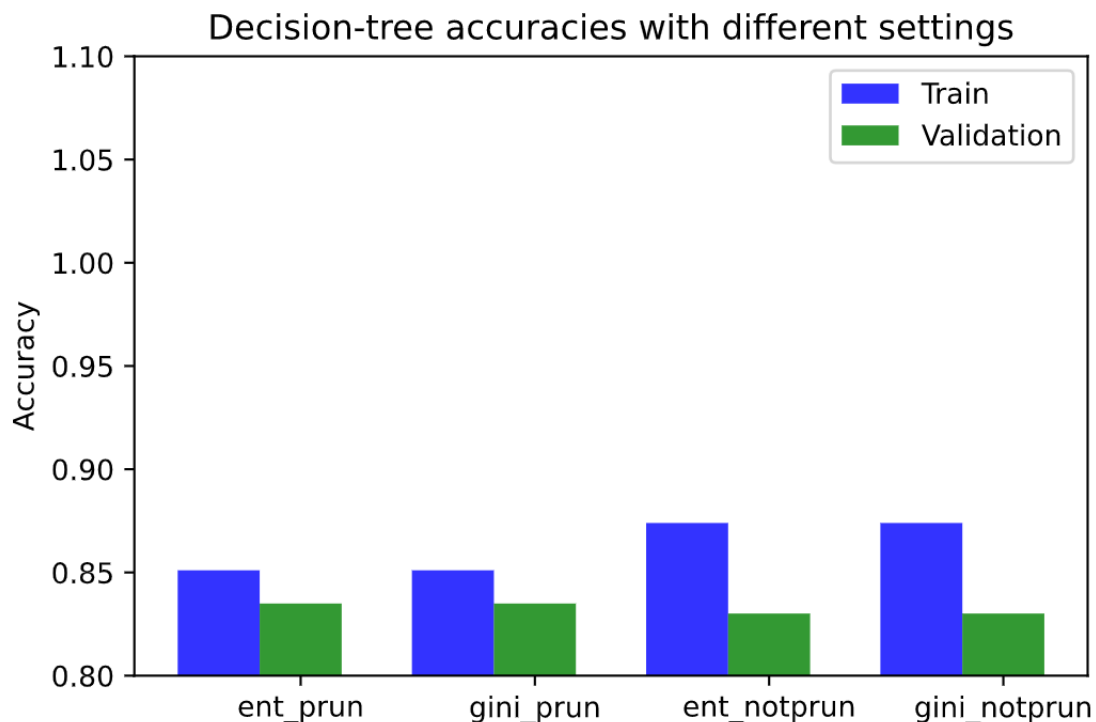


Abbildung 1: Results

The accuracies of the trees don't change for the different impurity measures. Moreover not using pruning, the training-accuracy is higher than the validation-accuracy, which was expected as it is susceptible to overfitting. As the validation-accuracy is higher using pruning, we decided to use the tree trained by using entropy and pruning.

## 8 Testing (Benjamin, Sophie)

We now tested the chosen tree to get a representative estimate of the accuracy.

```
1 test_acc = acc(tree_ent_prun, X_test, Y_test)
```

The test-accuracy was with 0.9515 lower than the according training- and validation-accuracy. Nevertheless, the difference was less than 1.6% and a slightly lower test-accuracy is to be expected as we used the other accuracies to choose this tree.

## 9 Comparison to existing implementations (Sophie)

The time measurements show, that the given implementation from the sklearn library is a lot faster. In learning and testing it is 10 times faster than our implementation. This is as expected for many reasons. The sklearn decision tree doesn't use pruning per default, which saves a lot of time. The library probably uses optimized data structures to build the tree as well as optimized algorithms. In the design of our decision tree we didn't consider time a critical factor and therefore didn't pay attention to time-saving code and functions.