

Complete decision tree induction functionality in scikit-learn

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Preface

I would like to thank everybody who kept me busy the last year, especially my promoter and my assistants. I would also like to thank the jury for reading the text. My sincere gratitude also goes to my wive and the rest of my family.

Ir. Sven Van Hove

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Abstract

The abstract environment contains a more extensive overview of the work. But it should be limited to one page. [EHWP16]

Introduction

1.1 Context

Decision tree induction is one of the most well-known tools in the machine learning community. Most of the theoretical groundwork was laid in the last three decades of the previous century. Researchers Leo Breiman and Ross Quinlan have been particularly influential in this space. Some well know algorithms include the Concept Learning System [HMS66], ID3 [Qui79, Qui83, Qui86] by Quinlan and Classificatiom And Regression Trees (CART) [BFSO84] by Breiman.

Contemporary AI researchers focus most of their attentions on neural networks and in particular deep learning — the recent hype around DeepMind's AlphaGo [SSS⁺17] victories comes to mind — but decision tree research is not dead. Researchers still continue to propose new or improved algorithms and analyses.

Theory is one thing, but the algorithms need to be implemented as computer programs to actually be useful. Sci-kit learn [PVG+11] is a very popular machine learning library written in Python. As such, it also contains implementations of various decision tree induction algorithms. Before sci-kit learn became popular, a Java-based library called Weka [EHWP16] (or "Waikato Environment for Knowledge Analysis" in full) was often used instead. Even today, the implementations of decision tree algorithms in Weka are still in many respects superior to those in scikit-learn. Other libraries that implement similar algorithms exist (e.g., Apache Spark [ZXW+16]), but those are beyond the scope of this text.

1.2 Goal

The goal of this thesis is to alleviate the discrepancies between sci-kit learn and Weka concerning decision tree induction. Mind that decision tree induction tools can never be truly "complete" as stated in the title because the field is immensely broad and still continues to grow. Nevertheless, an effort can be made to improve feature parity between these two popular tools.

One such discrepancy was found when comparing the performance of Weka and sci-kit learn on an activity dataset [KWM11]. The difference between classification accuracies in this case was considerable at about 25% in favor of Weka.

1.3 Motivation

Some would perhaps question the relevance of such *outdated* techniques anno 2018. This feeling is misguided. The advantages of decision tree induction algorithms are still hard to compete with, even for more modern algorithms [PVG⁺11, Mur98, KZP07]:

- 1. Comprehensible: makes intuitive sense even for the uninitiated.
- 2. Transparent, as opposed to for example artificial neural networks
- 3. Easy to visualize tree (if number of nodes remains small)
- 4. Non-parametric, makes very few assumptions about data
- 5. No data normalization required
- 6. Handles both categorical and numeric data
- 7. Handles missing data elegantly
- 8. Handles multiclass, multilabel and multioutput problems natively
- 9. Fast training
- 10. Fast inference

Of course decision tree induction algorithms are not perfect:

- 1. Unstable: small modifications in training data can result in a completely different tree
- 2. Learning optimal trees is an NP-Complete problem [HR76], so heuristics are used to find approximations
- 3. Prone to overfitting if not actively countered by adding early stopping criteria or an extra pruning step
- 4. Prone to bias when one class appears more much frequently in the training set than others.

1.4 Thesis structure

The structure of the remainder of this text is as follows. First, an overview of the literature study concerning decision tree induction will be presented. In particular the link between an implementation and its underlying algorithm will be clarified, including the effects of that choice on the capabilities of the tool.

Literature review

The relevant literature for this thesis mostly consists of papers concerning decision tree induction. These go back many decades, but fortunately there are some review and survey papers that provide a convenient overview [Mur98, RM05, KZP07]. On top of the academic literature, the source code and accompanying documentation of scikit-learn and Weka has also been a rich source of information.

2.1 Prerequisites

The reader ought to be familiar with basic machine learning concepts such as supervised learning, classification, regression, bias-variance trade-offs, model validation and ensemble learning. Furthermore, elementary knowledge of decision tree induction is expected. The most important basic concepts will be discussed briefly. Topics that are particularly important for the next chapters will be elaborated on.

2.2 Scope

A wide variety of decision tree induction algorithms exists. Here, only the *top down* induction of decision trees (TDIDT) family is considered. It is the most common approach and it is particularly relevant to the software tools under scrutiny.

Furthermore, only classification trees are considered. With little effort, most TDIDT classification algorithms can be converted to regression algorithms. Yet, these are far less popular and better alternatives such as XGBoost [CG16] exist.

Ensemble methods are also out of scope. Recent decision tree algorithms rarely work with a single tree, but rather with an ensemble of trees. Random forests [Bre01] is a very popular example of bootstrap aggregating or *bagging*. Regardless, the scope of this thesis concerns the fundamentals of decision trees, and not their derivatives. Implementation improvements suggested in this thesis could still potentially benefit related ensemble methods.

The algorithms in scope are all offline learning methods invented before the big data era. This implies that computation is done locally and that all data has to fit in memory. As such, online learning methods or distributed algorithms are out of scope.

Finally, only univariate tests are in scope. The test performed in each internal node must only evaluate one attribute of the observation. For categorical attributes, this typically implies checking whether or not the input is equal to a fixed category. For numeric (and thus ordered) attributes, the input value is compared against a fixed threshold using the less than or equal and greater than operators. Consequently, the input space is partitioned recursively using axis-aligned hyperplanes. This scope limitation precludes well-known but seldom used extensions such as oblique trees.

2.3 Terminology

Throughout the relevant literature, there is a lack of ubiquitous vocabulary shared by all researchers. Decision trees are used in various scientific fields, each with its own jargon. Specifically, there is a big divide between researchers that approach the problem from a machine learning perspective compared to those who come from a statistics background. To avoid confusion, some basic terms are reviewed. A decision tree consists of (internal) nodes which are connected to other nodes via a one-to-many parent-child relation on one hand, and leaves which have no children on the other hand. The root node is the only node without parent. In a binary tree, all internal nodes have two children.

Induction algorithms typically receive a training set as input data to construct a decision tree while a test set is used afterwards for model validation. These sets are tables of data where each row represents an observation. All observation are fully described by a common set of attributes. Some attributes are categorical, others may be numeric. Because decision tree induction is a part of supervised learning, one or more class labels are also associated with each observation. If the total number of distinct class values equals two, the task is called binary classification. Otherwise it is called multiclass classification. Multilabel classification occurs when one observation can be tagged with a variable number of class values at once. Multioutput classification on the other hand occurs when multiple distinct classes, each with their own set of values, have to be derived from the same set of attributes. This can be accomplished trivially by creating multiple trees, each handling one class. Regardless, combining them in one tree might offer performance benefits. In a way, multilabel classification is a special case of multioutput classification. Decision trees are one of the few machine learning algorithms that can handle all these modes of operation natively.

During training, first one root node is created and all observations in the training set are stored in this node. When a node is *split* using some test function, that test partitions the observations in subsets and then creates a child node for each subset. This process is repeated recursively until some stopping citerium is reached. The *purity* of a node is defined as the percentage of observations in that node that belong to the majority class. A *pure node* is a node with 100% purity.

2.4 A generic TDIDT algorithm

A typical TDIDT algorithm for classification consists of two phases: a grow phase and an optional prune phase. The grow phase requires three functions with fixed signatures: a test generation function, a splitting function and a stopping function. Historically, researchers presented their TDIDT algorithms with fixed functions. Because of the common interface it is now common to choose these functions \grave{a} la carte. One could try to evaluate the performance of each function separately, but choosing the best of each function does not guarantee a global optimum. Holistic tests must be performed to ensure the best configuration is chosen.

2.4.1 Univariate test generation

Based on the observations in a node, tests can be devised that spit those observations in a number of subsets. The goal of this step is to generate a finite number of tests $\tau_i \in \mathcal{T}$ based on one given attribute. Recall the tests based on multiple attributes exist but are out of scope. In the next step, one specific test is chosen from this set of possible tests.

Generating tests for categorical attributes is trivial. For binary trees the value of the attribute is compared against one specific category. If it matches, it belongs to the first subset, else to the second. This results in as many tests as there are possible categories for the attribute. For non-binary trees, one test suffices that maps each distinct category to a specific subset.

In the case of numeric, ordered attributes, threshold are introduced to partition the observations based on that ordering. That way, an infinite number of tests can be generated, which is of course undesireable. However, at least for the training data, not all tests will result in a different partitioning. A clever choice of thresholds should bring the number of subsets back to a manageable level.

2.4.2 Splitting

Classic TDIDT algorithms work by recursively splitting nodes based on some optimal test $\tau \in \mathcal{T}$, the set of all possible tests. A heuristic called the splitting criterion is required to determine this τ . A few such criteria have stood the test of time.

Purity

The perfect test τ^* creates a partition $S_{\tau^*} = \{S_1, \ldots, S_k\}$ wherein each subset is pure, so optimizing for weighted average partition purity is a sensible first criterion.

$$p(S_{\tau}) = \sum_{i} \frac{|S_i|}{|S|} p(S_i)$$
(2.1)

Here, $S = S_1 \cup \ldots \cup S_k$ and p(S) is the set purity as described above.

Entropy and information gain

In practice purity does not appear to work very well. That is why researchers came up with an alternative based on Shannon's information theory [Sha48]. Quinlan used such metrics in many of his prominent algorithms such as ID3 and C4.5 [Qui86, Qui93], but it was already invented earlier for the Concept Learning System (CLS) [HMS66]. Define entropy (or missing information) of a variable V with possible values v_i and associated probabilities p_i as follows:

$$s(V) = -\sum_{i} p_i \log_2(p_i) \tag{2.2}$$

The same concept can be applied to the class variable. Define the class entropy $s_C(S)$:

$$s_C(S) = -\sum_c p(c) \log_2(p(c))$$
 (2.3)

where p(c) is the probability that a random observation in S belongs to class c. This value can be defined for any node, independent of any specific partition.

For a given test τ , a similar definition can be given for each subset S_i of the induced partition on S:

$$s_C(S_i) = -\sum_c p_i(c) \log_2(p_i(c))$$
 (2.4)

For the entropy of the whole partition S_{τ} , again use the weighted average entropy of its subsets:

$$s_C(\mathcal{S}_\tau) = \sum_i \frac{|S_i|}{|S|} s_C(S_i) \tag{2.5}$$

Finally, calculate the information gain $h_{IG}(\tau, S)$ of the split that resulted from test τ :

$$h_{IG}(\tau, S) = s_C(S) - s_C(S_\tau) \tag{2.6}$$

where S_{τ} is the partition resulting from test τ .

Gain ratio

The information gain criterion is biased towards tests with many possible outcomes. This could be a problem in non-binary trees. The gain ratio alleviates this problem. First define split information $SI(\tau, S)$ — the maximum possible information gain — as follows:

$$SI(\tau, S) = -\sum_{i} \frac{|S_i|}{|S|} \log_2 \frac{|S_i|}{|S|}$$
 (2.7)

Finally, define the gain ratio:

$$h_{GR}(\tau, S) = \frac{h_{IG}(\tau, S)}{SI(\tau, S)}$$
(2.8)

In binary trees, this heuristic typically causes a less balanced tree compared to the information gain criterion [Qui93].

Gini

Distance metrics such as the Gini impurity index can be used instead of heuristics based on information theory [BFSO84]. The definitions follow the same pattern as those of the information gain:

$$g(S) = \sum_{c} p(c)(1 - p(c))$$
 (2.9)

$$g(S_i) = \sum_{c} p_i(c)(1 - p_i(c))$$
 (2.10)

$$g(\mathcal{S}_{\tau}) = \sum_{i} \frac{|S_i|}{|S|} g(S_i)$$
(2.11)

$$h_G(\tau, S) = g(S) - g(S_\tau) \tag{2.12}$$

2.4.3 Stopping

Breiman argues in his CART book [BFSO84] that choosing good stopping criteria is far more important than choosing good splitting criteria. If early stopping was not applied or no pruning (see below) was performed afterwards, trees would grow excessively large on real world data sets. This is a classic case of overfitting. It negatively impacts many factors that make decision trees attractive in the first place, such as their comprehensibility and their fast training and inference. It is also detrimental to the performance of the model on unseen data since the model fails to generalize properly.

Some simple stopping criteria are based on the depth of the tree, the purity of a node or the number of observations belonging to a node.

More complex stopping criteria are based on the Minimum Discription Length (MDL) of a tree [Ris78] or on statistical techniques such as a Ξ^2 -test. Quilan proposed to use the latter in his ID3 algorithm but decided not to include it in the successor (C4.5) [Qui86, Qui93].

2.4.4 Pruning

A better alternative to early stopping criteria is to let the tree grow freely, and to prune it afterwards in a bottom-up fashion. Typically, the current performance of an internal node including its children is compared to what the performance would be if this node would be converted to a leaf. If it would preform better as a leaf, the children are pruned away. Many different pruning algorithms exists. What follows is a non-exhaustive list of common pruning approaches.

Reduced Error Pruning (REP)

Reduced Error Pruning is one of the most straightforward and statistically sound methods of pruning a tree [Qui87, EK01a, EK01b]. Instead of using the whole training set to grow the tree, some randomly chosen observations are withheld in a seperate validation set. This way, an unbiased estimate of the performance of (a part of) tree can be calculated.

The disadvantage of this method is that less data is available for growing the tree, potentially negatively impacting this process. This downside is mitigated if training data is available in abundance.

Error Based Pruning (EBP)

Error Based Pruning is a technique used in C4.5 [Qui93]. It does not require a separate validation set, so the full training set can be used to grow the tree. The downside of this is that this method is less statistically sound. An upper bound is calculated based on the observed error and that upper bound used instead in the performance calculation.

Cost Complexity Pruning (CCP)

Cost Complexity Pruning, used in the CART algorithm [BFSO84], takes another approach akin to regularization in classic optimization problems. It considers both the total measured error and a cost factor proportional to the size of the tree. If the error increases, but it is compensated for by a much smaller tree, the operation as a whole can still be considered positive.

Pessimistic Pruning

[Man97] [Qui87] [Qui93]

Others

Many other pruning algorithms exist [BA97, Elo99, Min89, EMSK97]. Some use the MDL concept [MRA⁺95, QR89]. Others use a small neural network and backpropagation to find good pruning candidates [KC01].

Alternative: Rule-based Pruning

An outlier in this list is rule-based pruning. Decistion trees can be converted to a series of if-then statements where the condition is a conjunctive clause. These statements can be further simplified to if-then-else statements and then optimized, which can be seen as an alternative form of pruning. The resulting model is no long a tree, but it can still approximate the underlying concept that the tree used to represent.

2.5 Conclusion

TDIDT algorithms are made up of different components, for each of which a number of alternatives are available. This makes them a very flexible tool with uses in a variety of settings. Popular algorithms such as C4.5 and CART are opiniated in the sense that they each propose one specific configuration of components. Fortunately, that does not stop algorithm implementers from offering more choice to their users, as shown in the next chapter. Note also that there is no single precise definition of ID3, C4.5 or CART. New insights were aquired over time and added to the solution, but the algorithm name rarely changed.

Software for decision tree induction

Intro

- 3.1 Capabilities
- 3.2 Conclusion

Methodology

Intro

4.1 Conclusion

Results and discussion

Intro

- 5.1 Pruning
- 5.2 Categorical attributes
- 5.3 Conclusion

Conclusion

Intro

- 6.1 Contributions
- 6.2 Retrospective
- 6.3 Future work

Appendices

Appendix A

The First Appendix

Appendices hold useful data which is not essential to understand the work done in the master's thesis. An example is a (program) source. An appendix can also have sections as well as figures and references.

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