

# Partial Classification Forest

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Abstract

## I. Introduction

Some datasets do not allow a classifier to generate a decision surface good enough to be able to predict unseen observations well. Well, in this case, refers to a context dependent threshold for any quality measurement of a classifier, for example the accuracy or an information loss metric.

But for some of those problems, it may still be valuable to predict only on partitions of the feature space, in which the dataset is ‘clean’ enough, meaning a classifier can be found within the subset of the dataset laying inside one of those partitions which equals or exceeds the threshold.

This paper proposes a Monte Carlo based ensemble method called Partial Classification Forest (PCF), which builds an ensemble of trees having a structure similar to k-d trees to partition the feature space of the dataset in order to find ‘clean’ partitions. In the following a tree generated by the PCF is spelled Tree with a capital T, rather than tree, which is used to denote the tree data structure.

It should be noted here, that this paper is rather a Proof of Concept of a very early version of the PCF and has several shortcomings in research and empirical tests, due to a lack of time and no complete, fast implementation. I will discuss these shortcomings further in Chapter VI, but the most important ones I will list right away:

- Tests with a more sophisticated  $\gamma$  (cmp. II-A)

- The PCF’s performance on high dimensional data
- A Tree’s growing behaviour
- Benchmarks

In Section II I will lay out the structure and the operations of a Tree generated by the PCF before, in Section III, describing how PCF utilizes Tree instances. After that I will continue displaying test results using PCF. In Section V I will discuss further optimizations and possible additional features before finishing with a conclusion.

## II. The Tree structure

A Tree generated by the PCF is a binary search tree structure similar to k-d trees. Its purpose is to randomly generate disjoint partitions of a feature space.

A Tree has two types of nodes, non-leaf nodes, here denoted as Nodes and leaf nodes denoted as Leafs. It provides two operations: (i) FIT, initializing the Tree and (ii) PREDICT, returning a label for an observation.

The Node structure contains three properties: (i) a split value; (ii) a left and (iii) a right successor, both references to either another Node or a Leaf.

A Leaf on the other hand, is the structure representing a partition of the feature space, having the following properties: (i) active, a boolean value deciding whether the partition’s quality, determined during the FIT operation, is equal or better than the defined threshold or not; (ii) optionally a predictor which is used to classify observations during the PREDICT operation. Only if a Leaf’s active property is true, a predictor must be provided. A Leaf also has

two vectors with arbitrary length as properties: (iii) a vector containing the observations of the dataset used in FIT, which are laying inside the partition and (iv) their inherent labels.

During the FIT operation a Tree contains a third type of node, Nil. Nil is used to initialize Trees and the left and right successor of a Node. These nodes are transformed during FIT to either a Node or a Leaf, so after the FIT operation a Tree does not contain Nil nodes anymore. A Nil node does not have any properties.

#### A. The FIT operation

The FIT operation constructs a Tree, based on a dataset split in observations ( $X$ ) and their labels ( $y$ ). Algorithm 1 shows how FIT recursively builds a Tree, which is at the beginning a pointer to a Nil node.

The most important parameter passed to FIT is  $\gamma$ .  $\gamma$  is a function returning (i) a predictor and (ii) the loss of it. Otherwise  $\gamma$  is treated as a black box by the PCF, so what the predictor is and how its loss is calculated are not relevant to the PCF, as long as the predictor is callable and returns an element from the label set when called (Algorithm 2, line 9). The loss returned by  $\gamma$  gets compared to the quality threshold  $\tau_l$ . Is the loss  $\leq \tau_l$  the predictor is good enough and  $\Theta$  is transformed to an active Leaf (Algorithm 1, lines 2, 3).

There are two other thresholds besides  $\tau_l$ ,  $\tau_{|X|}$ ,  $\tau_h$ . Both regulate the behaviour of a Tree's growth.  $\tau_{|X|}$  defines a minimum amount of observations a Leaf must contain. One can easily imagine, without  $\tau_{|X|}$  or  $\tau_{|X|} = 0$  a Tree would never stop growing, since FIT would continue to split empty partitions, trying to find a smaller partition which would be predictable, even though no predictor could be generated without observations to train it on.

$\tau_h$  further regulates the maximum path length of a Tree. It is necessary besides  $\tau_{|X|}$ , because of the following scenario: be  $\tau_{|X|} = 2$  and there are two equal observations in the dataset, but both having a different label than the other one. Now  $\gamma$ , passed  $X$  containing only those two

identical observations, returns a predictor with a loss  $> \tau_l$ . Since  $|X|$  is still not smaller than  $\tau_{|X|}$  FIT would continue trying to separate the two inseparable observations. To prevent such a szenario  $\tau_h$  tells FIT to stop before the Tree's height, the amount of edges of the longest path, would exceed  $\tau_h$ . The path length of the Tree's root to  $\Theta$  is passed as a parameter  $h$  to FIT.

Now, if neither  $\tau_l$  is exceeded nor  $\tau_{|X|}$  or  $\tau_h$  is violated, FIT performs a split and transforms  $\Theta$  to a Node (Algorithm 1, lines 7ff). The dimension the split is performed on is chosen in a cyclic manner, a practise also applied to k-d trees (Algorithm 1, line 7). [1] But rather than choosing the splitting value at the median of the observations in the dimension, which is done in order to construct balanced k-d trees, the splitting value is random.[1]

In order to chose a proper splitting value  $\beta_X$  is passed as another parameter to FIT.  $\beta_X$  represents the boundries for every dimension of the feature space based on  $X$ . For each dimension  $\beta_X$  contains a tuple with the minimum and maximum value in the dimension of all observations in  $X$ .

$\beta_X[\text{dimension}]$  is passed to a pseudo-random number generator generating a random value so that  $\text{lower}(\beta_X[\text{dimension}]) \leq \text{random number} \leq \text{upper}(\beta_X[\text{dimension}])$  (Algorithm 1, line 8).

Afterwards  $X$ ,  $y$ ,  $\beta_X$  are splitted into two new disjoint partitions and FIT is recursively applied to the two new partitions (Algorithm 1, lines 10ff).

Since  $\tau_h$  is defined, the maximum amount of nodes a Tree can have is  $2^{\tau_h+1} - 1$ , if the Tree would be perfectly balanced. For each node FIT is called, so building a Tree has a worst case time complexity of  $\mathcal{O}((2^{\tau_h+1} - 1) * \mathcal{O}(\text{FIT}))$ .  $\mathcal{O}(\text{FIT})$  is determined by the size of  $X$ , since  $X$  has to be splitted and by  $\mathcal{O}(\gamma)$ . That said, a single FIT operation would have a worst case time complexity of  $\mathcal{O}(|X| + \mathcal{O}(\gamma))$ , which would mean the time complexity of the whole fitting process would be  $\mathcal{O}((2^{\tau_h+1} - 1) * (|X| + \mathcal{O}(\gamma)))$ .



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Algorithm 1 : FIT( $\Theta, X, y, h, \beta_X, \gamma, \tau_l, \tau_{|X|}, \tau_h$ )

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A Tree's FIT operation.

Inputs:

- $\Theta$  — a pointer to a Nil node; initially pointing to the root node of an empty Tree,
- $X$  — input data,
- $y$  — labels of  $X$ ,
- $h$  — height of the Tree; initially  $h = 0$ ,
- $\beta_X$  — lower and upper boundaries of every dimension of  $X$ ,
- $\gamma$  — function returning a predictor and its loss,
- $\tau_l$  — loss threshold,
- $\tau_{|X|}$  — threshold for the size of  $X$ ,
- $\tau_h$  — height limit of the Tree

Output: void

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- 1: predictor, loss  $\leftarrow \gamma(X, y)$
  - 2: if loss  $\leq \tau_l$  then
  - 3:    $\Theta \leftarrow \text{LEAF}(\text{true}, \text{predictor}, X, y)$
  - 4: else if  $h > \tau_h$  or  $|X| < \tau_{|X|}$  or loss  $> \tau_l$  then
  - 5:    $\Theta \leftarrow \text{LEAF}(\text{false}, \text{predictor}, X, y)$
  - 6: else
  - 7:   dimension  $\leftarrow h \bmod |X[0]|$
  - 8:   split  $\leftarrow \text{RANDOM}(\beta_X[\text{dimension}])$
  - 9:    $\Theta \leftarrow \text{NODE}(\text{split}, \text{NIL}, \text{NIL})$
  - 10:   split  $X, y$  and  $\beta_X$  into  $X', X'', y', y'', \beta'_X, \beta''_X$
  - 11:   FIT( $\Theta.\text{left}, X', y', h + 1, \beta'_X, \dots$ )
  - 12:   FIT( $\Theta.\text{right}, X'', y'', h + 1, \beta''_X, \dots$ )
  - 13: end if
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Both FIT and PREDICT can be implemented as multithreaded operations as long as  $\gamma$  is threadsafe, since the Tree instances are independent of each other and the shared parameters  $X, y$  (FIT) and  $x$  (PREDICT) are read only, making synchronization unnecessary.

FIT first computes  $\beta_X$  which has a time complexity of  $\mathcal{O}(|X[0]| * |X|)$ ,  $|X[0]|$  denoting the amount of dimensions the feature space has.

After that the Tree's FIT operation is called  $N$  times, which means the PCF's FIT operation has a worst case time complexity of

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Algorithm 2 : PREDICT( $\Theta, x, h$ )

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A Tree's PREDICT operation.

Inputs:

- $\Theta$  — a Tree node; initially pointing to the root of the Tree,
- $x$  — an observation,
- $h$  — height of the Tree; initially  $h = 0$

Output: the predicted label or  $\Lambda$

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- 1: if TYPE( $\Theta$ ) is Node then
  - 2:   dimension  $\leftarrow h \bmod |x|$
  - 3:   if  $x[\text{dimension}] \leq \Theta.\text{split}$  then
  - 4:     PREDICT( $\Theta.\text{left}, x, h + 1$ )
  - 5:   else
  - 6:     PREDICT( $\Theta.\text{right}, x, h + 1$ )
  - 7:   end if
  - 8: else if  $\Theta.\text{active}$  then
  - 9:   return  $\Theta.\text{predictor}(x)$
  - 10: else
  - 11:   return  $\Lambda$
  - 12: end if
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$\mathcal{O}(N * (2^{\tau_h+1} - 1) * (|X| + \mathcal{O}(\gamma)) + |X[0]| * |X|)$  (cmp. II-A).

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Algorithm 3 : FIT( $\Pi, X, y, \gamma, \tau_l, \tau_{|X|}, \tau_h$ )

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The PCF's FIT operation.

Inputs:

- $\Pi$  — a PCF instance,
- $X$  — input data,
- $y$  — labels of  $X$ ,
- $\gamma$  — function returning a predictor and its loss,
- $\tau_l$  — loss threshold,
- $\tau_{|X|}$  — threshold for the size of  $X$ ,
- $\tau_h$  — height limit of the Tree

Output: void

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- 1: compute  $\beta_X$
  - 2: for all  $\Theta \in \Pi.\text{trees}$  do
  - 3:   FIT( $\Theta, X, y, 0, \beta_X, \dots$ )
  - 4: end for
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The PCF's PREDICT operation first initializes an array with  $N$  elements (Algorithm 4, line 1). Each Tree instance fills one element of the

array with its prediction. After that the PCF's PREDICT operation takes the label predicted most and returns it as its prediction for the observation  $x$  (Algorithm 4, lines 5, 6).

The worst case time complexity of the PCF's PREDICT operation is  $\mathcal{O}(N * (\tau_h + \mathcal{O}(\text{predictor})) + N)$ , since a Tree's PREDICT operation is executed  $N$  times, plus the most predicted label must be determined, which is  $\mathcal{O}(N)$ .

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Algorithm 4 : PREDICT( $\Pi, x$ )

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The PCF's PREDICT operation.

Inputs:

$\Pi$     –    a PCF instance,  
 $x$     –    an observation,

Output: the predicted label or  $\Lambda$

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1: predictions  $\leftarrow [\Lambda; N]$ 
2: for  $i = 1$  to  $N$  do
3:   predictions[ $i$ ] = PREDICT( $\Pi.\text{trees}[i]$ ,  $x$ ,
    0)
4: end for
5: determine  $l_{max}$ , the label predicted most
6: return  $l_{max}$ 

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#### IV. Tests

#### V. Further optimizations and additional features

#### VI. Conclusion

#### References

- [1] Brown, R. A. Building a balanced  $k$ -d tree in  $o(kn \log n)$  time. Journal of Computer Graphics Techniques (JCGT) 4, 1 (March 2015), 50–68.
- [2] Cormen, T. H., Leiserson, C. E., Rivest, R. L., and Stein, C. Introduction to Algorithms, Third Edition, 3rd ed. The MIT Press, 2009.

