Fair and Optimal Decision Trees: A Dynamic Programming Approach

Abstract

Interpretable and fair machine learning models are required for many applications, such as credit assessment and in criminal justice. Decision trees offer inherent interpretability, especially when they are small. Optimal decision trees are of particular interest because they offer the best performance possible for a given size. However, state-of-the-art algorithms for fair and optimal decision trees have scalability issues, often requiring several hours to find such trees even for small datasets. In contrast to these state-of-the-art methods that use mixed integer programming, we propose a method that exploits the tree structure using dynamic programming. A key component in our method is a new pruning mechanism that reduces the search space by comparing partial solutions based on upper and lower bounds on their final fairness values. As a result, our model can find fair and optimal trees several orders of magnitude faster than previous methods, also for larger datasets that were previously beyond reach. Moreover, we show that with this substantial improvement our method can find the full Pareto front in the trade-off between accuracy and fairness.

1 Introduction

As machine learning (ML) is used in more domains that involve discrimination-sensitive decisions, demand for *fair*, *interpretable* and *accurate* models increases. ML, for example, is used in criminal justice [8], credit assessment [28], and housing appointments [7], each of which requires fair decisions. Simply removing the discrimination-sensitive attribute from the dataset does not necessarily result in less discrimination [33]. Instead, fairness is obtained by adding a constraint on the classifier: often either an individual or a group fairness constraint (see [10, 20] for a comparison of the two). The focus in this paper is on *group fairness*. With group fairness, a binary classifier is considered fair if the probability of receiving the positive outcome is the same for all protected groups. This fairness metric is formulated as a global constraint over the full dataset.

To increase confidence and prevent errors in these sensitive-decision domains, it also important that the ML models are interpretable [39]. Decision trees offer inherent interpretability and do not carry some of the risks of post-hoc attempts at explaining black-box models [36]. This is especially the case when decision trees are small [34]. On average, *optimal* decision trees, i.e., trees that for a given size provably maximize an objective such as accuracy, provide better out-of-sample performance than trees generated by heuristics when comparing trees of equal size [9, 15]. Therefore small optimal decision trees are ideal when searching for accurate and interpretable models.

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There are several approaches for finding fair and optimal decision trees [2, 3, 25, 40], all of which use mixed integer programming (MIP). However, none of these models scale well. If fairness is not considered, small optimal decision trees can be found efficiently by using dynamic programming (DP) methods that exploit the separability present in the tree structure [4, 5, 15, 23, 31]. Demirović and Stuckey [14] can even find the Pareto front of optimal decision trees for biobjective nonlinear metrics. However, a global fairness constraint cannot trivially be added to these algorithms, because the fairness constraint breaks the separability assumption by introducing dependency between subproblems.

Motivated by the success of DP methods for conventional optimal trees, we developed a novel DP method that incorporates a global fairness constraint. We solve the subproblem dependency by computing upper and lower bounds on the final fairness values of partial solutions, thus enabling early pruning of the search space. As a result of this and other improvements in the merging and comparing of subproblems, our method is several orders of magnitude faster than the state-of-the-art MIP methods.

This has several benefits. First, our algorithm can find fair and optimal decision trees for datasets that were previously too large to consider, often even within seconds. Second, it can find deeper and more accurate trees, while still satisfying the fairness constraint. Last, our algorithm can find the full Pareto front, helping data analysts to understand the trade-off between fairness and accuracy for a specific dataset.

As is commonly done in optimal decision tree search, we assume that all features have been binarized in advance [12]. Similarly, it is assumed that the label of every instance in the dataset is binary: either the positive/preferred outcome or the negative outcome. This is a common assumption in fair classification [16] and was also assumed in previous fair and optimal decision tree methods [25, 40].

The paper setup is as follows: we first discuss related work and preliminaries, and then we present our method. Finally, we evaluate the performance of the new method on twelve datasets from different application domains and we compare its performance to a leading MIP method from the literature.

2   Related work

This section discusses, in order, the related work of 1) algorithms for finding optimal decision trees, 2) MIP models for finding fair optimal decision trees, and finally 3) other (non-optimal) methods for finding fair decision trees.

Algorithms for Optimal Decision Trees.   The search for optimal decision trees is an NP-Hard problem [24]. Therefore decision tree search was traditionally done by means of heuristics such as the much-used CART algorithm [11]. Because of increasing computation power, searching for optimal decision trees has become feasible and Bertsimas and Dunn [9] showed that optimal decision trees can offer 1-5% better out-of-sample accuracy scores than previous heuristics. Their MIP model was consecutively improved by several others [41, 42].

During the same period, custom search and bound-based approaches, using techniques such as caching and branching, for finding (binary) classification trees were developed. These methods often outperformed the MIP models in terms of runtime and scalability, sometimes reducing runtimes of several hours to several seconds, by utilizing dynamic programming in a depth-first [4, 5, 15] or breadth-first approach [23, 31].

Of particular interest for this paper is the method presented in [14] which extends on these methods and finds decision trees for biobjective optimisation, minimizing the misclassification score for both classes in binary classification at the same time and returning the Pareto front of nondominated solutions. This method is further discussed in the preliminaries.

Fairness in Optimal Decision Trees.   Currently, the only methods for finding fair and optimal decision trees are MIP-based. Verwer and Zhang [40] developed a MIP model with a soft group fairness constraint in the objective. They report how their method can find a fair depth-three tree for a

dataset of 1000 instances in 15 minutes, when the model is given an initial solution from CART. But for the fair tree case they do not report being able to prove optimality within this time limit.

Aghaei et al. [2] present a MIP model that co-optimizes fairness and error for both classification and regression. Apart from group fairness, they also consider a notion of individual fairness separately. However, they are criticized by Ranzato et al. [35] that their model often provides accuracy not much better than a constant classifier. In [3] and [25], they continue their work and present a MIP model based on a max-flow formulation that results in a considerable improvement of the runtime. However, these models still have scalability issues because of the large number of binary variables.

Other methods for Fair Decision Trees. For completeness but deviating from the focus of this paper, the following section mentions a number of related fair decision tree methods that do not guarantee a globally optimal solution. These can be divided into pre-, in-, and postprocessing methods.

Kamiran and Calders [26] present a preprocessing method that changes the labels in the dataset according to a group fairness metric. The updated dataset can then be used to train any regular classifier.

Some of the in-processing methods consider fairness in the classifier itself. Others use a regular classifier, combined with a meta- or master problem that takes care of the fairness constraint. Pedreschi et al. [33], for example, develop a method for finding fair association rules that can then be used to compile decision trees. Agarwal et al. [1] iteratively solve a number of cost-sensitive classification problems, with the costs determined by the number of violations of the fairness constraint. Grari et al. [21] propose a gradient tree boosting approach with an adversarial fairness constraint. Detassis et al. [17] use a MIP master problem that changes the labels for the next training iteration to satisfy the fairness constraint. There is, however, no guarantee of convergence. Valdivia et al. [38] developed an evolutionary algorithm that returns a Pareto front of decision trees with the multiobjective of both fairness and accuracy.

Kamiran et al. [27] test a CART-based approach with a split criterion that is a mix of information gain in the label and the sensitive feature. They did not observe a significant improvement when using this approach. As an alternative, they suggest a postprocessing relabeling method that uses a greedy knapsack algorithm to decide which leaf nodes to relabel to improve fairness at a minimum cost of accuracy.

Summary of related work. In summary, the current state-of-the-art has scalable custom methods for finding optimal conventional decision trees, but adding fairness constraints to these methods is non-trivial. MIP-based solutions exist for this problem, but they do not scale well. The alternative is to use heuristic methods, which result in suboptimal solutions.

3 Preliminaries

This section discusses the preliminaries of our method: an introduction of notation, a formal introduction of group fairness, a formal problem definition, and a dynamic programming formulation for optimal decision tree search that does not consider fairness.

Notation. A dataset $\mathcal{D}$ contains instances $(\mathbf{x}, a, y)$, with $\mathbf{x}$ the feature vector, a the sensitive attribute and y the label value. The number of instances in the dataset $\mathcal{D}$ is $N = |\mathcal{D}|$. Let y_i be the label of instance i and a_i the value of the protected feature. Let $\mathcal{F}$ be the feature set and $f \in \mathcal{F}$ a feature in the feature set. We use f_i to denote the value of this feature for instance i. Since we are only considering binary features and binary classification, we introduce the shorter notation f_i and $\bar{f}_i$ to mean respectively that instance i satisfies or does not satisfy feature f. The same idea applies for y and $\bar{y}$, and a and $\bar{a}$. We use subscript notation to refer to a subgroup of the dataset: for example, $\mathcal{D}_{\bar{f}}$ refers to all instances in dataset $\mathcal{D}$ that do not satisfy feature f; N_a is the number of instances in

the dataset that belong to group a, and $N_{y,\bar{a}}$ is the number of instances in the dataset with positive outcome that does not belong to group a.

Group Fairness Formalized. The specific group fairness definition considered in this paper is *demographic parity* (although our method can easily be extended to support other notions of group fairness, such as equality of opportunity). Demographic parity requires that the performance (the expected percentage of positive outcomes) per group is the same [20]. More formally, if $\hat{y}$ is the predicted outcome, 1 is the positive outcome, and a represents a (binary) sensitive/protected group, then demographic parity holds iff:

$$P(\hat{y} = 1 \mid a = 1) = P(\hat{y} = 1 \mid a = 0) \tag{1}$$

In binary classification, this constraint can be measured by considering the relative average performance of the two groups. Let $\hat{p}$ be the percentage of instances that receive the positive outcome, so $\hat{p}_a = N_{\hat{y},a}/N_a$. The difference in the performance between two groups is what we will call the imbalance $I = \hat{p}_a - \hat{p}_{\bar{a}}$. When the imbalance is limited to some value δ, we say that it satisfies demographic parity up to bias δ:

$$|I| = |\hat{p}_a - \hat{p}_{\bar{a}}| \le \delta \tag{2}$$

According to the definition of demographic parity, if a leaf node n receives label 0, the partial imbalance I_n in that node is also 0. Otherwise, the partial imbalance in a leaf node n can be expressed as follows:

$$I_n = \frac{N_{n,a}}{N_a} - \frac{N_{n,\bar{a}}}{N_{\bar{a}}} \tag{3}$$

The partial imbalance of a branching node is the sum of the imbalance of both sub-nodes.

Problem Definition. The task of learning an optimal fair decision tree is to find the feature value tests in the branch nodes and the leaf node assignments for a tree of a given maximum size that minimize the number of misclassifications in a training data set while observing a fairness constraint. The percentage of correctly classified instances is called accuracy. This can be formalized as follows. The task is to find a decision tree classifier $h : \{0, 1\}^{|\mathcal{F}|} \to \{0, 1\}$ of depth d that minimizes the misclassification score for a given dataset $\mathcal{D}$ while observing a fairness constraint up to bias δ:

$$\min_h \sum_{(\mathbf{x},a,y) \in \mathcal{D}} |h(\mathbf{x}) - y|$$

$$\text{s.t.} \left| \sum_{(\mathbf{x},a,y) \in \mathcal{D}_a} \frac{h(\mathbf{x})}{N_a} - \sum_{(\mathbf{x},a,y) \in \mathcal{D}_{\bar{a}}} \frac{h(\mathbf{x})}{N_{\bar{a}}} \right| \le \delta \tag{4}$$

Alternatively, when searching for the full Pareto front, the parameter δ is not necessary, and the problem can be written as a multiobjective problem:

$$\min_h \left\{ \sum_{(\mathbf{x},a,y) \in \mathcal{D}} |h(\mathbf{x}) - y|, \left| \sum_{(\mathbf{x},a,y) \in \mathcal{D}_a} \frac{h(\mathbf{x})}{N_a} - \sum_{(\mathbf{x},a,y) \in \mathcal{D}_{\bar{a}}} \frac{h(\mathbf{x})}{N_{\bar{a}}} \right| \right\} \tag{5}$$

The Pareto front for Eq. 5 consists of all nondominated solutions (M, I) with M the misclassification score and I the imbalance. A solution is considered dominated if there is another solution that performs better or similar on all objectives. Therefore, we can define the dominance relation $(\succ)$ for two solutions (M_1, I_1) and (M_2, I_2), and the function nondom as follows:

$$(M_1, I_1) \succ (M_2, I_2) \text{ iff } M_1 \le M_2 \wedge |I_1| \le |I_2| \wedge (M_1, I_1) \ne (M_2, I_2) \tag{6}$$

$$\text{nondom}(S) = \{s_1 \in S \mid \neg \exists \, s_2 \in S, s_2 \succ s_1\} \tag{7}$$

Dynamic Programming Formulation for Decision Trees. The problem of finding accurate decision trees has been formulated as a DP problem before in [15]. Of interest to this paper is the

formulation for biobjective optimization in [14]. They minimize the misclassification score of two classes at the same time and return the Pareto front of nondominated solutions:

$$T_{\text{BI}}(\mathcal{D}, d) = \begin{cases} \{(|\mathcal{D}_y|, 0), (0, |\mathcal{D}_{\bar{y}}|)\} & d = 0 \\ \text{nondom}\left(\cup_{f \in \mathcal{F}} \text{merge}\left(T_{\text{BI}}(\mathcal{D}_f, d-1), T_{\text{BI}}(\mathcal{D}_{\bar{f}}, d-1)\right)\right) & \text{else} \end{cases} \quad (8)$$

At the leaf node ($d = 0$), the function returns two solutions consisting of two values: the misclassification score per class. In all other cases, the function should return the merge of two solution sets and retain only the nondominated solutions resulting from that merge. The function merge combines two sets of partial solutions (from the left and right subtree) by element-wise addition:

$$\text{merge}(S_1, S_2) = \{(a_1 + a_2, b_1 + b_2) \mid (a_1, b_1) \in S_1, (a_2, b_2) \in S_2\} \quad (9)$$

Their method assumes that the objective is *additive* and *monotonic*. The additive property holds if two partial solutions (a, b) and (a', b') can be combined by addition: $(a + a', b + b')$. This property holds also for the misclassification and imbalance objective. The monotonic property holds for a biobjective function $f(a, b)$ if for any two possible different inputs a, b and a', b', the following dominance relation holds:

$$a \leq a' \wedge b \leq b' \rightarrow f(a, b) \succ f(a', b') \quad (10)$$

However, the dominance relation for fair and accurate trees expressed in Eq. 6 is not monotonic due to the absolute value function for the imbalance.

4 Method Description

In this section, we present our method DPF. Similar to Eq. 8, we define a recursive function T_{F} that returns a set of all nondominated (partial) solutions, resulting in a full Pareto front of fair and optimal decision trees. In order to guarantee that the method returns the full Pareto front we must do an exhaustive search of all combinations of subtrees, pruning only those solutions for which we can prove that they will never be part of an optimal (nondominated) solution.

The following 1) first explains how solutions for subproblems (partial solutions) can be compared and filtered, 2) how the function T_{F} is defined, 3) how the method can be changed to find a single best tree, and finally, 4) what further improvements we have implemented in our method.

Upper and lower bounds for fairness. The key idea in our method is to compute upper and lower bounds for the imbalance value of partial solutions to enable comparison. When upper and lower bounds $\bar{I}_R$ and $\underline{I}_R$ are known for the imbalance in the rest (R) of the tree, then the final imbalance value of a tree that contains node n will be in the range: $[\underline{I}_R + I_n, \bar{I}_R + I_n]$. The lower bound ($\underline{I}$) and the upper bound ($\bar{I}$) for the final *absolute* imbalance value can now be computed:

$$\underline{I}(I_n, [\underline{I}_R, \bar{I}_R]) = \begin{cases} 0 & \text{if } 0 \in [\underline{I}_R + I_n, \bar{I}_R + I_n] \\ \min\left(|\underline{I}_R + I_n|, |\bar{I}_R + I_n|\right) & \text{else} \end{cases} \quad (11)$$

$$\bar{I}(I_n, [\underline{I}_R, \bar{I}_R]) = \max\left(|\underline{I}_R + I_n|, |\bar{I}_R + I_n|\right) \quad (12)$$

Trivial bounds on the value I_R can be derived from inspecting the dataset (e.g., assigning one label to all instances of one group and the other label to all other instances, or vice versa). When partial solutions for the rest of the tree are known, the bounds can be derived by taking the partial solutions with the most extreme value for I_R.

With these upper and lower bounds on the final absolute imbalance value, we can redefine the dominance relation of Eq. 6:

$$\begin{aligned} (M_1, I_1) \succ (M_2, I_2)) \text{ iff } & (M_1, I_1) \neq (M_2, I_2) \\ & \wedge M_1 \leq M_2 \\ & \wedge \left(\bar{I}(I_1, [\underline{I}_R, \bar{I}_R]) \leq \underline{I}(I_2, [\underline{I}_R, \bar{I}_R]) \vee I_1 = I_2\right) \end{aligned} \quad (13)$$

A solution is dominated by another solution if 1) the other solution has a lower misclassification score; and 2) either an upper bound on the absolute imbalance lower than the lower bound on the absolute imbalance of this solution, or precisely the same imbalance score.

Finding the Pareto front of optimal fair trees. Now we can define a new function for finding fair trees $T_F(\mathcal{D}, d, [\underline{I}_R, \bar{I}_R])$ that returns all nondominated subtrees. Its building blocks are the functions merge and nondom based on our new dominance comparison as described in Eq. 13. (For ease of notation, $[\underline{I}_R, \bar{I}_R]$ will often be left out.)

In a leaf node n, the function should return the following (M, I) solutions, with imbalance values based on Eq. 3 and using the dominance relation as defined in Eq. 13:

$$\text{leaf}(\mathcal{D}, [\underline{I}_R, \bar{I}_R]) = \text{nondom}\left(\left\{(|\mathcal{D}_y|, 0), \left(|\mathcal{D}_{\bar{y}}|, \frac{N_{n,a}}{N_a} - \frac{N_{n,\bar{a}}}{N_{\bar{a}}}\right)\right\}\right) \quad (14)$$

When branching, the data set is split on some feature f, resulting in two subproblems. Each subproblem is solved independently by passing the best known bounds for the other subproblem. Let $U(\mathcal{D}_{\bar{f}}, [\underline{I}_R, \bar{I}_R], d-1)$ denote the function that returns these best known bounds. These bounds can be trivial, based on dataset inspection, or based on previous (cached) solutions. The solutions from the two subproblems are then merged as follows:

$$\begin{aligned}
\text{branch}(\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]) = \text{nondom}\,(\cup_{f \in \mathcal{F}}\, \text{merge}\,(\\
T_F(\mathcal{D}_{\bar{f}}, d-1, \quad U(\mathcal{D}_f, [\underline{I}_R, \bar{I}_R], d-1)),\\
T_F(\mathcal{D}_f, d-1, \quad U(\mathcal{D}_{\bar{f}}, [\underline{I}_R, \bar{I}_R], d-1))))
\end{aligned} \quad (15)$$

All combined this results in the DP formulation that returns the full Pareto-front that optimizes Eq. 5:

$$T_{\text{F}}(\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]) = \begin{cases} \text{leaf}(\mathcal{D}, [\underline{I}_R, \bar{I}_R]) & d = 0 \\ \text{branch}(\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]) & \text{else} \end{cases} \quad (16)$$

Finding a single best tree. It is also possible with this same DP formulation to search for a single optimal and fair tree up to bias δ, as defined in Eq. 4. This can be achieved by pruning all (partial) solutions and retaining only those that are *possibly* fair. Then, in the root node of the search, the tree with minimum misclassification score is selected from the list of solutions. A solution is possibly fair if the lower bound on the absolute imbalance value is less than or equal to δ. For this, we introduce the function prune, which should filter a set of solutions S before it is passed to the nondom function.

$$\text{prune}(S, [\underline{I}_R, \bar{I}_R]) = \left\{(M_n, I_n) \mid (M_n, I_n) \in S,\ \underline{I}(I_n, [\underline{I}_R, \bar{I}_R]) \leq \delta\right\} \quad (17)$$

Other algorithmic improvements. To reduce the runtime cost of the merge function, before merging, the two sets of partial solutions are first pruned based on bounds on the misclassification score and discrimination score derived from the other set. Then the two sets are sorted by misclassification score to enable early termination of the merge if it can be proven that all consecutive combinations of partial solutions will exceed the misclassification upper bound. In the root node of the search, the solutions are sorted on the imbalance value to enable faster elimination of non-fair solutions.

The runtime of the nondom function can be reduced by observing that partial solutions with a lower bound imbalance of 0 do not need to be compared with other solutions because they will always be nondominated. For the sake of brevity, all these have been left out of the method description. See our code repository for full implementation details.[1]

Similar to [14, 15] our method uses caching, upper and lower bounds on the misclassification score, and a special depth-two solver to increase the runtime performance. Appendix A presents the pseudocode of DPF. Note that our algorithm may easily be modified to also optimize tree sparsity. Appendix B shows how.

5 Experimental Results

The following section analyzes the performance of DPF. We focus on the analysis of the runtime and scalability of DPF: 1) How does DPF compare to the state-of-the-art in terms of runtime? 2) What

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[1]

impact do our pruning method and merge improvements have on the runtime? 3) What is the runtime cost to find the full Pareto front? 4) What impact do parameters such as number of features, instances, etc. have on the runtime of DPF?

For an out-of-sample analysis, see Appendix C.

Experiment setup. We evaluate DPF on all datasets mentioned in the survey by Le Quy et al. [30]. The data preprocessing and binarization is also done as described in that paper. Categorical variables are encoded through one-hot encoding, except for variables with twenty or more categories, which were removed (this applies to the KDD census dataset). The binarized datasets are included in our code repository (if the license permits redistribution). The Diabetes and Law-school dataset are left out of the evaluation because the best tree of depth three is (almost) identical to the trivial solution (assign the majority label to all instances). The references for the original datasets can be found here [6, 13, 18, 19, 29, 32, 37].

The algorithm receives the whole dataset as input and is asked to find a tree with an unfairness limit of $\delta = 1\%$. In a later experiment, we also examine the impact of this parameter on the runtime. The values shown are the average of five runs and a time limit of one hour is set for every run. All experiments are run on a 2.6Ghz Intel i7 CPU with 8GB RAM using only one thread.

The FairOCT model To our knowledge, the only methods available for finding fair and optimal decision trees are MIP models [2, 25, 40]. Verwer and Zhang [40] limit their model to solving problems with less than 1000 instances. In our initial comparison of [2] and [25], the latter outperformed the first by a large margin. Therefore, we compare our method with their FairOCT model. They model the decision tree as a flow graph where all instances must flow from the source (connected to the root node of the tree) to one of the sink nodes (one for each class), to which all nodes are connected. In our experiments, the FairOCT model is solved with Gurobi 9.0 using the default parameters.

Runtime comparison. Table 1 shows the runtime comparison between FairOCT and our method DPF. DPF can find optimal trees for $d = 2$ within one second for all datasets, about four to five orders of magnitude faster than FairOCT. FairOCT is able to find $d = 2$ trees only for the smallest datasets within one hour. It struggles with an increasing number of data instances because every data instance is represented by a binary variable. Also for $d = 3$, DPF can find optimal trees within one second for several datasets and all problems are solved within a minute. For $d = 4$, the exponential nature of the problem becomes clearly visible. Datasets with both large number of features and instances are no longer solvable within one hour. However, DPF still succeeds in finding the best tree for a number of datasets, among which Adult, with over 45000 instances.

Table 1: Runtime comparison of DPF and FairOCT. Runtime is in seconds. FairOCT resulted in an out of memory error for KDD census income, so no runtime can be reported.

Dataset	a	$|\mathcal{D}|$	$|\mathcal{F}|$	FairOCT $d = 2$	DPF $d = 2$	$d = 3$	$d = 4$
Adult	Gender	45222	17	> 1h	< 1	< 1	1012
Bank Marketing	Married	45211	46	> 1h	< 1	2	> 1h
Communities & Crime	Race	1994	97	> 1h	< 1	4	1261
COMPAS recid.	Race	6172	9	> 1h	< 1	< 1	9
COMPAS viol. recid.	Race	4020	9	1594	< 1	< 1	< 1
Dutch census	Gender	60420	58	> 1h	< 1	5	> 1h
German credit	Gender	1000	69	> 1h	< 1	2	741
KDD census income	Race	284556	117	-	< 1	32	> 1h
OULAD	Gender	21562	45	> 1h	< 1	3	> 1h
Ricci	Race	118	4	< 1	< 1	< 1	< 1
Student-Mathematics	Gender	395	55	284	< 1	< 1	24
Student-Portuguese	Gender	649	55	866	< 1	< 1	32

The impact of our novel pruning method and the merge improvements   Table 2 shows the runtime results when DPF is run 1) without our novel pruning method (not applying the prune and nondom methods of Eqs. 7 and 17; 2) without our improvements to the merge method; 3) without our improvement to the nondom method, and 4) to generate a full Pareto front.

For several datasets, our pruning method decreases the runtime by one or more orders of magnitude. This is specifically the case for depth four, where the subproblem dependency is more significant than for depth three. Our method can find optimal trees for depth four for eight out of twelve datasets within one hour, but without the pruning method, this is possible for only four datasets.

Our improvements to the merge function in some cases significantly reduce the runtime. This is specifically the case for datasets that have a high initial bias (the bias of an optimal tree if fairness would not be considered): Adult, COMPAS recid. and Dutch census have an initial bias of 18%, 16%, and 14% respectively, and for these datasets the merge improvements reduce the runtime by an order of magnitude. The other data sets in Table 2 have an initial bias of 1-6% and the merge improvements result in only a small reduction of the runtime. The improvements to the nondom function are also essential for the performance of our pruning mechanism.

Table 2: DPF runtimes without the pruning and dominance checks (¬P), without the merge improvements (¬M), without the nondom improvements (¬D), for finding the full Pareto-front (Pr), and for comparison the default runtime (Df). Datasets for which DPF has a default runtime below one second or over one hour, are left out.

| | | $d = 3$ | | | | | | $d = 4$ | | | |
Dataset	¬P	¬M	¬D	Pr	Df	Dataset	¬P	¬M	¬D	Pr	Df
Bank	5	3	319	4	2	Adult	> 1h	> 1h	> 1h	> 1h	1012
Com.&Cr.	13	5	19	7	4	Com.&Cr.	> 1h	1329	2473	1939	1261
Dutch	> 1h	101	1814	510	5	COMP. rcd.	> 1h	394	1658	1659	9
German	8	2	22	2	2	German	> 1h	770	1971	738	741
KDD	144	35	1254	50	32	Stud. Math	144	24	29	412	24
OULAD	5	4	1448	3	3	Stud. Port.	1834	32	38	259	32

Generating the full Pareto-front   Because of the runtime improvements of DPF, it is also able to find a *full*, fair and optimal decision tree Pareto front. In contrast, Valdivia et al. [38] find a partial and non-optimal Pareto front and the FairOCT model is used to find an optimal, but still partial, Pareto front. Table 2 shows the runtime performance of DPF when tasked to find the full Pareto front. In comparison to FairOCT, DPF is several orders of magnitude faster. For COMPAS recid., for example, DPF found the Pareto front for depth two consisting of 29 solutions in 0.12 ± 0.04 seconds. The FairOCT method generates the partial Pareto front by repeatedly solving the problem with a different maximum bias, resulting in a runtime several times higher than the values reported in Table 1.

See Figure 1 for a number of full Pareto fronts generated by DPF for depth 2-4. These plots confirm the trade-off between accuracy and fairness as reported in previous works.

Figure 1: The full Pareto front of training accuracy and discrimination for four datasets for trees of depth 2, 3 and 4, generated by DPF.

Figure 2: The impact of the number of features, instances, minimum leaf node size and maximum allowed bias in the fairness constraint, δ, on the runtime of DPF with $d = 3$. The shaded area shows the values within one standard deviation. Note the logarithmic scale of the y-axis.

Method evaluation. To provide more insight into our method we evaluate the impact of the number of features, the number of dataset instances, the minimum leaf node size and the maximum allowed bias in the fairness constraint on the runtime of DPF for three datasets when searching for trees of depth three. Figure 2 shows the results. It can be observed from these experiments that the number of features is the most important factor, resulting in an exponential increase, which is in line with previous studies [31]. One might expect that the method would scale linearly with increasing dataset size, but the experiments show almost no increase in runtime after a certain size. The reason for this is that the runtime of the method is mostly dependent on the number of unique partial solutions that need to be merged by Eq. 9. This number of unique solutions is strongly dependent on the number of features and is only weakly related to the number of instances.

DPF can easily be changed to limit the minimum leaf node size, by returning an empty set in Eq. 14 when the number of instances does not exceed the required leaf node size. Increasing the minimum leaf node size decreases the amount of unique partial solutions, and consequently we see a significant decrease in the runtime when the minimum leaf node size is increased. Adding such a minimum leaf node size often does not have a significant impact on the accuracy score.

Relaxing the fairness constraint to values larger than $\delta = 1\%$ causes the runtime to increase for the Dutch census dataset. This can be explained because a more relaxed constraint means a larger search space and less strong fairness bounds. However, when the constraint is even more relaxed, the problem is closer to a normal optimal tree search, and the accuracy bounds become stronger, again resulting in a shorter runtime. For the Bank marketing and OULAD datasets, the default discrimination when no fairness constraint is in place, is lower: 1.2% and 3.0%, versus Dutch census: 14.1%. This explains why these datasets show almost no difference in runtime when changing the value δ.

6 Conclusion

We present a first specialized algorithm for finding fair optimal decision trees that can also find the full Pareto front in terms of accuracy and fairness. Dynamic programming is used to exploit the separability present in the tree structure. We solve the subproblem dependency introduced by the global fairness constraint by computing upper and lower bounds on the final fairness value and thus enable comparison of partial solutions. The results show that our method can find fair optimal trees several orders of magnitude faster than the state-of-the-art. As a result, our method succeeds in finding fair and optimal decision trees even for large datasets that were previously beyond reach.

An extension of our method would be to support other group-based notions of fairness, such as equality of opportunity. To better deal with unbalanced datasets, another extension of the presented research would be to replace the accuracy objective with an objective that can cope better with unbalanced datasets, for example, with the biobjective method presented in [14], or by using balanced accuracy. Finally, based on the evaluation results, new heuristic methods could be investigated, for example, by considering not all possibly-fair partial solutions, but only a representative subset.

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Algorithm 1: Tree search of depth d with fairness on a dataset $\mathcal{D}$ for a feature set $\mathcal{F}$.

$T_{\mathrm{F}}(\mathcal{D}, d, \mathrm{UB}, [\underline{I}_R, \bar{I}_R])$

    if $d = 0$ **then**

        $S \leftarrow \left\{ (|\mathcal{D}_y|, 0), \left(|\mathcal{D}_{\bar{y}}|, \frac{N_{n,a}}{N_a} - \frac{N_{n,\bar{a}}}{N_{\bar{a}}} \right) \right\}$

        return filter$(S, \mathrm{UB}, [\underline{I}_R, \bar{I}_R])$

    $\langle F, \mathrm{lb}, \mathrm{stat} \rangle \leftarrow \mathrm{cache}[\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]]$

    if $\mathrm{lb} \geq \mathrm{UB}$ **then return** $\emptyset$

    if $\mathrm{stat} = \mathrm{optimal}$ **then return** filter$(F, \mathrm{UB}, [\underline{I}_R, \bar{I}_R])$

    $\mathrm{PF} \leftarrow \emptyset$

    for $f \in \mathcal{F}$ **do**

        $\mathrm{LB}_R \leftarrow \mathrm{LB}(\mathcal{D}_f, d-1, [\underline{I}_R, \bar{I}_R])$

        $\mathrm{PF}_L \leftarrow T_{\mathrm{F}}(\mathcal{D}_{\bar{f}}, d-1, \mathrm{UB} - \mathrm{LB}_R, U(\mathcal{D}_f, [\underline{I}_R, \bar{I}_R], d-1))$

        if $\mathrm{PF}_L = \emptyset$ **then continue**

        $\mathrm{LB}_L \leftarrow \mathrm{LB}(\mathcal{D}_{\bar{f}}, d-1, [\underline{I}_R, \bar{I}_R])$

        $\mathrm{PF}_R \leftarrow T_{\mathrm{F}}(\mathcal{D}_f, d-1, \mathrm{UB} - \mathrm{LB}_L, U(\mathcal{D}_f, [\underline{I}_R, \bar{I}_R], d-1))$

        if $\mathrm{PF}_R = \emptyset$ **then continue**

        $F \leftarrow \mathrm{prune}\left(\mathrm{merge}(\mathrm{PF}_L, \mathrm{PF}_R, [\underline{I}_R, \bar{I}_R]), \mathrm{UB}, [\underline{I}_R, \bar{I}_R]\right)$

        $\mathrm{PF} \leftarrow \mathrm{PF} \cup F$

    $\mathrm{PF} \leftarrow \mathrm{nondom}(\mathrm{PF})$

    if $\mathrm{PF} = \emptyset$ **then**

        $\mathrm{cache}[\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]] \leftarrow \langle \emptyset, \mathrm{UB}, \mathrm{lower\ bound} \rangle$

        return $\emptyset$

    $\mathrm{cache}[\mathcal{D}, d, [\underline{I}_R, \bar{I}_R]] \leftarrow \langle \mathrm{PF}, \underline{M}(\mathrm{PF}), \mathrm{optimal} \rangle$

    return PF

A Pseudocode for DPF

This section gives the pseudocode for DPF. Note that in contrast to the main text this pseudocode does consider upper and lower bounds and cache. Therefore, the function T_{F} receives the current best known upper bound on the misclassification score as an extra parameter.

The function prune must be redefined to take into account this upper bound on the misclassification score.

$$\mathrm{prune}(S, \mathrm{UB}, [\underline{I}_R, \bar{I}_R]) = \{ (M_n, I_n) \mid (M_n, I_n) \in S,$$
$$\underline{I}(I_n, [\underline{I}_R, \bar{I}_R]) \leq \delta, M_n < \mathrm{UB} \} \tag{18}$$

For shorter notation, define:

$$\mathrm{filter}(S, \mathrm{UB}, [\underline{I}_R, \bar{I}_R]) = \mathrm{nondom}\left(\mathrm{prune}\left(S, \mathrm{UB}, [\underline{I}_R, \bar{I}_R] \right), [\underline{I}_R, \bar{I}_R] \right) \tag{19}$$

Also we define the functions $\underline{M}$ to return the best misclassification score within a set of solutions:

$$\underline{M}(S) = \min \{ M \mid (M, I) \in S \} \tag{20}$$

Furthermore, let LB return the best known lower bound for a subproblem, or zero if no such lower bound is known.

With these changes, see Algorithm 1 for the resulting pseudocode of DPF. For a full Pareto front, the algorithm must be called with a fairness cut-off value of $\delta = 1$.

B Non-complete trees

For the sake of brevity, the main text of this paper only shows how to search for complete trees, that is, trees of depth d with $2^d - 1$ branch nodes and 2^d leaf nodes. Our method also allows to search for smaller trees, as was done similarly in [15].

Figure 3: Out-of-sample accuracy and discrimination for three datasets, for maximum bias $\delta = 1\%, 5\%, 100\%$. The figure shows the distribution of 10 runs.

The DP formulation in Eq. 15 can be extended as follows to also allow for incomplete/sparse trees. Here n signifies the number of nodes in the tree.

$$\text{nondom} \left(\text{prune} \left(\cup_{f \in \mathcal{F}, i \in [0, n-1]} \text{merge} \left(\right. \right. \right.$$
$$T_F(\mathcal{D}_{\bar{f}}, d-1, n-i-1, \quad U(\mathcal{D}_f, [\underline{I}_R, \bar{I}_R], d-1, i)), \tag{21}$$
$$\left. \left. \left. T_F(\mathcal{D}_f, d-1, i, \quad U(\mathcal{D}_{\bar{f}}, [\underline{I}_R, \bar{I}_R], d-1, n-i-1)) \right) \right) \right)$$

With this change in place, the solver can search for incomplete trees. This also allows to add a parameter α to prevent overfitting. The misclassification score now becomes $M + \alpha n$. The parameter α describes how much the misclassification score should at least decrease in order to justify adding another node to the tree. The addition of this parameter to the algorithm is trivial.

C Test Evaluation

Experiment setup. The evaluation in the main text is focused on analyzing the runtime of our DPF method. This section further analyzes the out-of-sample performance of DPF and compares it with the heuristic proposed in [27]. They propose several methods, but conclude that the best results are obtained by extending CART with a splitting criterion that is a mix of information gain in the class label and information gain in the sensitive attribute (IGC+IGS). After generating the tree, they heuristically change the label of some leaves in such a way that discrimination is minimized with the least loss of accuracy. We will call this method after their first author: Kamiran.

We evaluate the out-of-sample performance by running DPF for $d = 2, 3, 4$ and Kamiran for $d = 3, 4$ for different maximum allowed bias $\delta = 1\%, 5\%, 100\%$ on three datasets that are commonly evaluated

in the literature. Note that Kamiran with $\delta = 100\%$ is the same as plain CART. We run each case 10
times on random stratified train-test splits of 75% vs 25%. Figure 3 shows the resulting distribution
of test accuracy and test discrimination. The sign of the discrimination score tells which of the two
groups is discriminated against.

Evaluation. From the results in Figure 3 it can be observed that the optimal decision trees generated
by DPF in general have better out-of-sample accuracy than Kamiran for the same or smaller depth.
In several cases DPF $d = 2$ even outperforms Kamiran, even though it uses four times less nodes.
However, in general the variance in test accuracy is high.

The variance in test discrimination is also high, and both methods often exceed the imposed dis-
crimination limit in the test evaluation. This problem is less visible with the Adult dataset, probably
because of its larger number of data instances.

For both the COMPAS-recid. and German credit dataset almost maximum accuracy can already be
obtained with a decision tree of depth two, so the addition of extra nodes does not help much. For the
Adult dataset, however, deeper trees can provide better accuracy.

Tuned for number of nodes. In our next analysis, we prevent overfitting during training for both
methods by using 10 random validation splits of 25% of the training data to find what number of
branching nodes is best. The tree size with the best average accuracy in the validation set, while on
average respecting the discrimination constraint is selected as best. The full training dataset is then
used to generate a tree of that size. Table 3 shows the results when both algorithms are used to find
trees of a maximum bias of 5%.

Discussion. DPF searches for optimal decision trees, which means it will always find the tree with
maximum accuracy for the training dataset, and thus always outperform heuristics on performance
in the training dataset. The results in Table 3 show that when DPF is tuned for selecting the right
number of nodes, this also generalizes to better performance than Kamiran in the test set. DPF has
higher accuracy score for almost all datasets, while still respecting the discrimination constraint on
average. For some datasets the difference in accuracy is as high as 11.6% (Student-Mathematics)
or 9.2% (Dutch census). Only for the COMPAS-viol.-recid. dataset is the average accuracy of DPF
worse by 0.1% than for Kamiran, but this difference is not statistically significant.

Table 3: Out-of-sample average accuracy and discrimination $\pm$ the standard deviation (%) for
solutions with a maximum training bias of $\delta = 5\%$ and a maximum depth of $d = 3$. Best performing
accuracy score per dataset is marked bold, if significant (p-value $\leq 5\%$), except for Ricci, because
DPF's discrimination score is too high for fair comparison.

Dataset	DPF		Kamiran	
	Accuracy	Discrimination	Accuracy	Discrimination
Adult	**80.4** $\pm$ 0.3	-5.1 $\pm$ 0.4	78.0 $\pm$ 1.5	-1.9 $\pm$ 1.1
Bank marketing	**89.7** $\pm$ 0.1	1.7 $\pm$ 0.6	89.0 $\pm$ 0.4	0.5 $\pm$ 0.3
Communities & Crime	**94.6** $\pm$ 0.3	-3.2 $\pm$ 1.0	93.9 $\pm$ 0.3	-1.3 $\pm$ 2.3
COMPAS-recid.	**59.1** $\pm$ 2.4	-4.6 $\pm$ 3.1	55.6 $\pm$ 1.9	-1.1 $\pm$ 2.1
COMPAS-viol.-recid.	83.7 $\pm$ 0.2	-0.3 $\pm$ 0.5	83.8 $\pm$ 0.1	-0.3 $\pm$ 0.8
Dutch census	**77.4** $\pm$ 0.3	-5.0 $\pm$ 1.0	68.2 $\pm$ 10.9	-0.6 $\pm$ 0.6
German credit	70.3 $\pm$ 1.7	-2.1 $\pm$ 3.0	70.3 $\pm$ 1.0	-0.9 $\pm$ 3.4
KDD census income	**94.3** $\pm$ 0.0	-1.0 $\pm$ 0.1	93.9 $\pm$ 0.0	0.0 $\pm$ 0.1
OULAD	**68.7** $\pm$ 0.3	-2.1 $\pm$ 1.0	68.0 $\pm$ 0.1	0.1 $\pm$ 0.2
Ricci	66.0 $\pm$ 4.4	-13.4 $\pm$ 12.3	53.3 $\pm$ 0.0	0.0 $\pm$ 0.0
Student-Mathematics	**85.5** $\pm$ 6.6	-2.8 $\pm$ 8.3	73.9 $\pm$ 11.8	-2.5 $\pm$ 4.2
Student-Portuguese	90.5 $\pm$ 2.8	0.8 $\pm$ 5.0	89.4 $\pm$ 4.2	2.3 $\pm$ 3.1

Both methods have difficulties with finding fair trees for Ricci and Student-Mathematics. This can be explained by the small size of these datasets. Ricci has only 4 features and 118 instances. This makes it hard to find splits that are fair and accurate both in the training and test set.

The results also confirm the findings from Figure 3 that the variance in the discrimination value is often high, specifically for the small datasets. This means that for those instances it is difficult to generalize and overfitting in terms of discrimination is still happening. It is an open question how this can be reduced.