

A fast algorithm to compute a curve of confidence upper bounds for the False Discovery Proportion using a reference family with a forest structure

Guillermo Durand ¹ Laboratoire de Mathématiques d'Orsay, Université Paris-Saclay

Date published: 2025-02-11 Last modified: 2025-02-11

Abstract

This paper presents a new algorithm (and an additional trick) that allows to compute fastly an entire curve of post hoc bounds for the False Discovery Proportion when the underlying bound V_{\Re}^* construction is based on a reference family \Re with a forest structure à la Durand et al. (2020). By an entire curve, we mean the values $V_{\Re}^*(S_1), \dots, V_{\Re}^*(S_m)$ computed on a path of increasing selection sets $S_1 \subseteq \dots \subseteq S_m$, $|S_t| = t$. The new algorithm leverages the fact that going from S_t to S_{t+1} is done by adding only one hypothesis.

Keywords: multiple testing, algorithmic, post hoc inference, false discovery proportion, confidence bound

Contents

2	1	Introduction	Z
3	2	Notation and reference family methodology	3
4		2.1 Multiple testing notation	3
5		2.2 Post hoc bounds with reference families	3
6		2.3 Regions with a forest structure	4
7	3	New algorithms	7
8		3.1 Pruning the forest	7
9		3.2 Fast algorithm to compute a curve of confidence bounds on a path of selection sets	9
10		3.3 Illustration on a detailed example	12
11		3.4 Proof of Theorem 3.1	17
12		3.4.1 Derivation of (18)	17
13		3.4.2 Proof of (16) and (17)	17
14	4	Implementation	23
15	5	Numerical experiments	24
16	6	Conclusion	27
17	7	Acknowledgements	27
18	Re	eferences	27
19	Se	ssion information	29

 $^{^{1}} Corresponding\ author:\ guiller mo.dur and @universite-paris-saclay. fr$

1 Introduction

Multiple testing theory is often used for exploratory analysis, like Genome-Wide Association Studies, where multiple features are tested to find promising ones. Classical multiple testing theory like Family-Wise Error Rate (FWER) control or False Discovery Rate (FDR) control (Benjamini and Hochberg, 1995) can be used, but a more recent trend consists in the computation of post hoc bounds, also named post selection bounds or confidence envelopes, for the number of false positives, or, equivalently, for the False Discovery Proportion (FDP). This approach is notably advocated for in the context of exploratory research by (Goeman and Solari, 2011, Section 1).

Mathematically speaking, a confidence upper bound (we prefer to say upper bound instead of envelope for obvious reasons) is a function $\hat{V}: \mathscr{P}(\mathbb{N}_m^*) \to \mathbb{N}_m$, where $\mathbb{N}_m = \{0, ..., m\}$, $\mathbb{N}_m^* = \{1, ..., m\}$ and m is the number of hypotheses, such that

$$\forall \alpha \in]0, 1[, \mathbb{P}\left(\forall S \subseteq \mathbb{N}_m^*, |S \cap \mathcal{H}_0| \le \hat{V}(S)\right) \ge 1 - \alpha. \tag{1}$$

Here, α is a target error rate and \mathcal{H}_0 is the set of hypotheses indices that are true null hypotheses. Note that the construction of \hat{V} depends on α and on the random data X and the dependence is omitted to lighten notation and because there is no ambiguity. The meaning of Equation (1) is that \hat{V} provides an upper bound of the number of null hypotheses in S for any selection set $S \subseteq \mathbb{N}_m^*$, which allows the user to perform post hoc selection on their data without breaching the statistical guarantee. Also note that by dividing by $|S| \vee 1$ in Equation (1) we also get a confidence bound for the FDP:

$$\forall \alpha \in]0, 1[, \mathbb{P}\left(\forall S \subseteq \mathbb{N}_m^*, \text{FDP}(S) \le \frac{\hat{V}(S)}{|S| \vee 1}\right) \ge 1 - \alpha.$$
 (2)

So post hoc bounds provide ways to construct FDP-controlling sets instead of FDR-controlling sets, which is much more desirable given the nature of the FDR as an expected value. See for example (Bogdan et al., 2015, Figure 4) for a credible example where the FDR is controlled but the FDP has a highly undesirable behavior (either 0 because no discoveries at all are made, either higher than the target level).

The first confidence bounds are found in Genovese and Wasserman (2006) and Meinshausen (2006), although, in the latter, only for selection sets of the form $\{i \in \mathbb{N}_m : P_i \leq t\}$ where P_i is the p-value associated to the null hypothesis $H_{0,i}$. In Goeman and Solari (2011) the authors re-wrote the generic construction of Genovese and Wasserman (2006) in terms of closed testing Marcus et al. (1976), proposed several practical constructions and sparked a new interest in multiple testing procedures based on confidence envelopes. This work was followed by a prolific series of works like Meijer et al. (2015) or Vesely et al. (2023). In Blanchard et al. (2020), the authors introduce the new point of view of references families (see Section 2.2) to construct post hoc bounds, and show the links between this meta-technique and the closed testing one, along with new bounds.

Following the reference family trail, in Durand et al. (2020) the authors introduce new reference families with a special set-theoretic constraint that allows an efficient computation of the bound denoted by $V_{\mathfrak{R}}^*$ on a single selection set S. The problem is that one often wants to compute $V_{\mathfrak{R}}^*$ on a whole path of selection sets $(S_t)_{t\in\mathbb{N}_m^*}$, for example the hypotheses attached to the t smallest p-values. Whereas the algorithm provided in the aforementioned work (Durand et al., 2020, Algorithm 1), which is reproduced here, see Algorithm 1, is fast for a single evaluation, it is slow and inefficient to repeatedly call it to compute each $V_{\mathfrak{R}}^*(S_t)$. If the S_t 's are nested, and growing by one, that is $S_1 \subseteq \cdots \subseteq S_m$ and $|S_t| = t$, there is a way to efficiently compute $(V_{\mathfrak{R}}^*(S_t))_{t\in\mathbb{N}_m}$ by leveraging the nested structure

This is the main contribution of the present paper: a new and fast algorithm computing the curve $\left(V_{\mathfrak{R}}^{*}(S_{t})\right)_{t\in\mathbb{N}_{m}}$ for a nested path of selection sets, that is presented in Section 3.2. An additional

algorithm that can speed up computations both for the single-evaluation algorithm and the new curve-evaluation algorithm is also presented, in Section 3.1. A detailed example illustrating how the new algorithms work is provided in Section 3.3, and the proof that the fast algorithm indeed computes correctly the curve is in Section 3.4. In Section 2.1, all necessary notation and vocabulary is re-introduced, most of it being the same as in Durand et al. (2020). In Section 4 we discuss the current implementations of all the presented algorithms in the R (R Core Team, 2024) package sanssouci (Neuvial et al., 2024). Finally, a few numerical experiments are presented in Section 5 to demonstrate the computation time gain.

2 Notation and reference family methodology

2.1 Multiple testing notation

As is usual in multiple testing theory, we consider a probability space $(\Omega, \mathcal{A}, \mathbb{P})$, a model \mathscr{P} on a measurable space $(\mathcal{X}, \mathfrak{X})$, and data that is represented by a random variable $X : (\Omega, \mathcal{A}) \to (\mathcal{X}, \mathfrak{X})$ with $X \sim P \in \mathscr{P}$, that is, the law of X is comprised in the model \mathscr{P} .

Then we consider $m \geq 1$ null hypotheses $H_{0,1}, \ldots, H_{0,m}$ which formally are submodels, that is subsets of \mathscr{P} . The associated alternative hypotheses $H_{1,1}, \ldots, H_{1,m}$ are submodels such that $H_{0,i} \cap H_{1,i} = \emptyset$ for all $i \in \mathbb{N}_m^*$. We denote by $\mathscr{H}_0 = \mathscr{H}_0(P)$ (the dependence in P will be dropped when there is no ambiguity) the set of all null hypotheses that are true, that is $\mathscr{H}_0(P) = \{i \in \mathbb{N}_m^* : P \in H_{0,i}\}$. In other words, $H_{0,i}$ is true if and only if $i \in \mathscr{H}_0$. For testing each $H_{0,i}, i \in \mathbb{N}_m^*$, we have at hand a p-value $p_i = p_i(X)$ (the dependence in X will be dropped when there is no ambiguity) which is a random variable with the following property: if $i \in \mathscr{H}_0$, then the law of p_i is super-uniform, which is sometimes denoted $\mathscr{L}(p_i) \succeq \mathscr{U}([0,1])$. This means that in such case, the cumulative distribution function (cdf) of p_i is always smaller than or equal to the cdf of a random variable $U \sim \mathscr{U}([0,1])$:

$$\forall x \in \mathbb{R}, \mathbb{P}(p_i \le x) \le \mathbb{P}(U \le x) = 0 \lor (x \land 1). \tag{3}$$

For every subset of hypotheses $S \subseteq \mathbb{N}_m^*$, let $V(S) = |S \cap \mathcal{H}_0|$. If we think of S as a selection set of hypotheses deemed significant, V(S) is then the number of false positives (FP) in S. V(S) is our main object of interest and the quantity that we wish to over-estimate with confidence upper bounds (see Equation (1) or the more formal Equation (4) below).

Finally let us consider the following toy example, that will be re-used in the remainder of the paper.

Example 2.1 (Gaussian one-sided). In this case we assume that $X = (X_1, ..., X_m)$ is a Gaussian vector and the null hypotheses refer to the nullity of the means in contrast to their positivity. That is, formally, $(\mathcal{X}, \mathfrak{X}) = (\mathbb{R}^m, \mathcal{B}(\mathbb{R}^m))$, $\mathcal{P} = {\mathcal{N}(\boldsymbol{\mu}, \Sigma) : \forall j \in \mathbb{N}_m^*, \mu_j \geq 0, \Sigma \text{ positive semidefinite}}$, for each $i \in \mathbb{N}_m^*$, $H_{0,i} = {\mathcal{N}(\boldsymbol{\mu}, \Sigma) \in \mathcal{P} : \mu_i = 0}$ and $H_{1,i} = {\mathcal{N}(\boldsymbol{\mu}, \Sigma) \in \mathcal{P} : \mu_i > 0}$. Then we can construct *p*-values by letting $p_i(X) = \bar{\Phi}(X_i) = 1 - \Phi(X_i)$, where Φ denotes the cdf of $\mathcal{N}(0, 1)$ and $\bar{\Phi}$ the associated survival function.

2.2 Post hoc bounds with reference families

With the formalism introduced in last section, a confidence upper bound is a functional $\hat{V}: \mathcal{X} \times]0,1[\to (\mathcal{P}(\mathbb{N}_m^*) \to \mathbb{N}_m)]$ such that,

$$\forall P \in \mathcal{P}, \forall X \sim P, \forall \alpha \in]0, 1[, \mathbb{P}\left(\forall S \subseteq \mathbb{N}_m^*, V(S) \le \hat{V}(X, \alpha)(S)\right) \ge 1 - \alpha. \tag{4}$$

In the remainder, the dependence in (X, α) will be dropped when there is no ambiguity and $\hat{V}(X, \alpha)$ will simply be written \hat{V} .

As said in the Introduction, many constructions, ultimately theoretically equivalent but differing by the practical steps involved, exist, and in this paper we focus on the meta-construction of Blanchard et al. (2020) based on reference families. A reference family is a family $\Re = \Re(X,\alpha) = (R_k,\zeta_k)_{k\in\mathscr{K}}$ with $|\mathscr{K}| \leq 2^m$, $R_k \subseteq \mathbb{N}_m^*$, $\zeta_k \in \{0,\dots,|R_k|\}$ where everything (that is, \mathscr{K} and all the R_k and ζ_k) depends on (X,α) but the dependency is not explicitly written. The R_k are all distinct. We also define the following error criterion for a reference family, named Joint Error Rate (JER):

$$JER(\Re) = \mathbb{P}\left(\exists k \in \mathcal{K}, |R_k \cap \mathcal{H}_0| > \zeta_k\right) = \mathbb{P}\left(\exists k \in \mathcal{K}, V(R_k) > \zeta_k\right). \tag{5}$$

In the following, we are only interested in reference families that control the JER at level α :

$$\forall P \in \mathcal{P}, \forall X \sim P, \forall \alpha \in]0, 1[, 1 - \text{JER}(\Re(X, \alpha)) = \mathbb{P}(\forall k \in \mathcal{K}, V(R_k) \le \zeta_k) \ge 1 - \alpha. \tag{6}$$

Note that Equation (6) is really similar to Equation (4) except that the uniform guarantee, instead of being over all $S \subseteq \mathbb{N}_m^*$, is only over all the $R_k \subseteq \mathbb{N}_m^*$, $k \in \mathcal{K}$, with \mathcal{K} having cardinality potentially much smaller than 2^m . A global confidence bound is then derived from a JER-controlling reference family by interpolation. Let

$$\mathscr{A}(\mathfrak{R}) = \{ A \subseteq \mathbb{N}_m^* : \forall k \in \mathscr{K}, |R_k \cap A| \le \zeta_k \}. \tag{7}$$

What says the JER control is that $\mathcal{H}_0 \in \mathcal{A}(\mathfrak{R})$. We leverage this information with the following confidence bound construction:

$$V_{\mathfrak{R}}^{*}(S) = \max_{A \in \mathscr{A}(\mathfrak{R})} |S \cap A| \tag{8}$$

which optimally uses the information provided by the JER control of the reference family, as proven by Proposition 2.1 of Blanchard et al. (2020). Because of the $\max_{A \in \mathscr{A}(\Re)}$, the computation of $V_{\Re}^*(S)$ is generally intractable (see Proposition 2.2 of Blanchard et al. (2020)), but for specific structures of reference families, a polynomial computation can be derived. This is the topic of Durand et al. (2020) and of next section.

118 2.3 Regions with a forest structure

The core concept of this section is to assume that the regions R_k 's of the reference family are what we called in Durand et al. (2020) a forest structure, that is two regions are either disjoint or nested:

$$\forall k, k' \in \mathcal{K}, R_k \cap R_{k'} \in \{R_k, R_{k'}, \emptyset\}. \tag{9}$$

Representing the R_k 's with a directed graph, where there is an oriented edge $R_k \leftarrow R_{k'}$ if and only if $R_k \subset R_{k'}$ and there is no $R_{k''}$ such that $R_k \subseteq R_{k''} \subseteq R_{k'}$ gives a forest, hence the name. See Example 2.2 and its representation in Figure 1.

We also need to introduce the notion of depth with the following function:

$$\phi : \begin{cases} \mathcal{K} \to \mathbb{N}^* \\ k \mapsto 1 + |\{k' \in \mathcal{K} : R_k \subseteq R_{k'}\}|. \end{cases}$$
 (10)

Example 2.2. Let m = 25, $R_1 = \{1, ..., 20\}$, $R_2 = \{1, 2\}$, $R_3 = \{3, ..., 10\}$, $R_4 = \{11, ..., 20\}$, $R_5 = \{5, ..., 10\}$, $R_6 = \{11, ..., 16\}$, $R_7 = \{17, ..., 20\}$, $R_8 = \{21, 22\}$, $R_9 = \{22\}$. This is the same example as Example 2 of Durand et al. (2020) and it is graphically depicted in Figure 1. The sets R_1 , R_8 are of depth 1; the sets R_2 , R_3 , R_4 , R_9 are of depth 2; the sets R_5 , R_6 , R_7 are of depth 3.

Another tool of Durand et al. (2020) that will be used is its Lemma 2, that is the identification of \Re with a set $\mathscr{C} \subset \left\{ (i,j) \in \left(\mathbb{N}_N^*\right)^2 : i \leq j \right\}$ such that for $(i,j), (i',j') \in \mathscr{C}, \{i,\dots,j\} \cap \{i',\dots,j'\} \in \mathscr{C}$

```
%%| filename: ../figure-tikz/fig-tikz-01
\begin{tikzpicture} [scale=1]
 \tikzstyle{quadri}=[circle,draw,text=black, thick]
 \tikzstyle{estun}=[->,>=latex,very thick]
 \node[quadri] (R1) at (0,3) {$R_1$};
 \node[quadri] (R2) at (-2,1) {$R_2$};
 \node[quadri] (R3) at (0,1) {$R_3$};
 \node[quadri] (R4) at (2,1) {$R_4$};
 \node[quadri] (R5) at (0,-1) {$R_5$};
 \node[quadri] (R6) at (1.5,-1) {$R_6$};
 \node[quadri] (R7) at (2.5,-1) {$R_7$};
 \node[quadri] (R8) at (4,3) {$R_8$};
 \node[quadri] (R9) at (4,1) {$R_9$};
 \draw[estun] (R1)--(R2);
 \draw[estun] (R1)--(R3);
 \draw[estun] (R1) -- (R4);
 \draw[estun] (R3)--(R5);
 \draw[estun] (R4)--(R6);
 \draw[estun] (R4)--(R7);
 \draw[estun] (R8)--(R9);
\end{tikzpicture}
```

Figure 1: The regions of Example 2.2.

 $\{\emptyset, \{i, ..., j\}, \{i', ..., j'\}\}$. With this identification, each $R_k = R_{(i,j)}$ can be written as $P_{i:j} = \bigcup_{i \le n \le j} P_n$ where $(P_n)_{1 \le n \le N}$ is a partition of \mathbb{N}_m^* . The P_n 's were called atoms in Durand et al. (2020) because they have the thinnest granularity in the structure, but to continue the analogy with graphs, forests and trees, they can also be called leafs. See Example 2.3 for a concrete example.

Example 2.3 (Continuation of Example 2.2). For the reference family given in Example 2.2, a partition of atoms is given by $P_1 = R_2$, $P_2 = R_3 \setminus R_5$, $P_3 = R_5$, $P_4 = R_6$, $P_5 = R_7$, $P_6 = R_8 \setminus R_9$, $P_7 = R_9$, $P_8 = \mathbb{N}_m^* \setminus \{R_1 \cup R_8\}$. Then $R_1 = P_{1:5}$, $R_3 = P_{2:3}$, $R_4 = P_{4:5}$ and $R_8 = P_{6:7}$. Note that not all atoms are regions of the family. Those new labels are graphically depicted in Figure 2. The nodes that correspond to atoms that are not in the family are depicted with a dashed circle, and all atoms are depicted in gray. This is the same example as Example 3 of Durand et al. (2020).

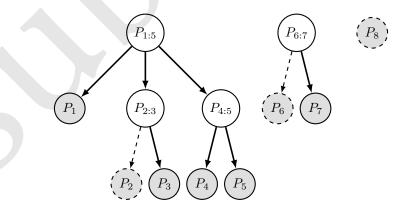


Figure 2: The regions of Example 2.2 but with the labels of Example 2.3.

When all leaves are regions of the family, it is said that the family is complete. If this is not the case, the family can easily be completed by adding the missing leaves (and using their cardinality as

associated ζ) without changing the value $V_{\mathfrak{R}}^*$. See Definition 2, Lemma 6 and Algorithm 2 of Durand et al. (2020) for the details.

Durand et al. (2020) also proved in their Theorem 1 that:

$$V_{\mathfrak{R}}^{\star}(S) = \min_{Q \subseteq \mathcal{X}} \left(\sum_{k' \in Q} \zeta_{k'} \wedge |S \cap R_{k'}| + \left| S \setminus \bigcup_{k' \in Q} R_{k'} \right| \right)$$
(11)

and, even better, in their Corollary 1 (iii) that:

$$V_{\mathfrak{R}}^{\star}(S) = \min_{Q \in \mathfrak{P}} \sum_{k' \in O} \zeta_{k'} \wedge |S \cap R_{k'}|, \tag{12}$$

provided that the family is complete. Here, $\mathfrak{P} \subseteq \mathcal{P}(\mathcal{K})$ is the set of subsets of \mathcal{K} that realize a partition, that is, the set of $Q \subseteq \mathcal{K}$ such that the R_k , $k \in Q$, form a partition of \mathbb{N}_m^* . So the minimum in Equation (12) is over way less elements than in Equation (11).

Finally, that paper provides a polynomial algorithm to $V_{\mathfrak{R}}^*(S)$ for a single $S \subseteq \mathbb{N}_m^*$, which we reproduce here in Algorithm 1 . The family is assumed complete, otherwise the first step would be to complete it. In the original paper, \mathscr{K}^h used to designate the elements of \mathscr{K} at depth h plus the atoms at depth $\leq h$. Actually one can realize that the last assumption is not needed for this algorithm to perform exactly the same, with the added benefit of not repeating computations at the atoms that don't have the maximal depth. The only change is that sometimes $Succ_k$ can be empty, in which case we simply let $newVec_k = \zeta_k \wedge |S \cap R_k|$. Thus, here in this paper, we define \mathscr{K}^h as only the elements of \mathscr{K} at depth h (the previous intricate definition may still be necessary for the proof of Theorem 1 of Durand et al. (2020)): $\mathscr{K}^h = \{(i,j) \in \mathscr{K} : \phi(i,j) = h\}, \quad h \geq 1$. This is the only deviation from the notation of Durand et al. (2020). Finally note that in the ongoing analogy with graph theory, the elements of \mathscr{K}^1 are the roots of the different trees making up the forest.

Algorithm 1 Computation of a given $V_{\Re}^*(S)$

```
1: procedure VSTAR(S, \Re = (R_k, \zeta_k)_{k \in \mathcal{X}} with \Re complete)
 2:
            H \leftarrow \max_{k \in \mathcal{K}} \phi(k)
                                                                                                                                       ⊳ maximum depth
            for h = H - 1, ..., 1 do
 3:
                  \mathcal{K}^h \leftarrow \{k \in \mathcal{K} : \phi(k) = h\}
 4:
                  newVec \leftarrow (0)_{k \in \mathcal{K}^h}
 5:
                  for k \in \mathcal{K}^h do
 6:
                        Succ_k \leftarrow \{k' \in \mathcal{K}^{h+1} : R_{k'} \subseteq R_k\}
 7:
                        if Succ_k = \emptyset then
 8:
                              newVec_k \leftarrow \zeta_k \wedge |S \cap R_k|
 9:
10:
                              newVec_k \leftarrow \min\left(\zeta_k \wedge |S \cap R_k|, \sum_{k' \in Succ_k} Vec_{k'}\right)
11:
                         end if
12:
                  end for
13:
                  Vec \leftarrow newVec
14:
            end for
15:
            return \sum_{k \in \mathcal{K}^1} Vec_k
16:
17: end procedure
```

Tip

In the practical implementation of this algorithm (and of the following Algorithm 2), Vec and *newVec* are always of size N(the number of leaves) instead of the cardinality of \mathcal{X}^h . And the sum $\textstyle \sum_{k' \in Succ_k} Vec_{k'} \text{ is really easy to compute: if } R_k = R_{(i_0,i_p-1)} = \bigcup_{j=1}^p R_{(i_{j-1},i_j-1)} = \bigcup_{i_0 \leq n \leq i_p-1} P_n \in \mathbb{R}_{(i_0,i_p-1)}$ \mathcal{K}^h for some $p \geq 2$, a strictly increasing sequence (i_0, \dots, i_p) and $R_{(i_{j-1}, i_{j}-1)} \in \mathcal{K}^{h+1}$ for all $1 \le j \le p$, then we simply sum *Vec* over the indices from i_0 to $i_p - 1$. After that, the computed quantity is set in newVec at index i_0 . So actually computing $Succ_k$ is not needed and not done.

161 162

169

The computation time of the algorithm is in $O(m|\mathcal{K}|)$, which is fast for a single evaluation, but calling it repeatedly on a path of selection sets $(S_t)_{t \in \mathbb{N}_m^*}$ has complexity $O(m^2 |\mathcal{X}|)$ which is not desirable and makes computations difficult in practice, hence the need for a new, faster algorithm.

Remark 2.1. The specific computation of the R_k 's and the ζ_k 's such that Equation (6) holds is outside the scope of the present paper, but different constructions can be found in Blanchard et al. (2020), Durand et al. (2020) or Blain et al. (2022) for example.

New algorithms

Pruning the forest

We remark the simple fact that if, for example, $(1, 1), (2, 2), (1, 2) \in \mathcal{K}$, and $\zeta_{(1,2)} \ge \zeta_{(1,1)} + \zeta_{(2,2)}$, then $R_{(1,2)}$ never contributes to the computation of any $V_{\Re}^{\star}(S)$ and it could just be removed from \Re . We now formalize and prove this pruning scheme.

Definition 3.1 (Pruning). We define by $\mathcal{K}^{\mathfrak{pr}}$ (\mathcal{K} pruned) the set of elements of \mathcal{K} such that we removed all (i, i') such that there exists $p \ge 2$ and integers i_1, \dots, i_{p-1} such that, when setting $i_0 = i$ and $i_p = i' + 1$, the sequence (i_0, \dots, i_p) is strictly increasing, $(i_{j-1}, i_j - 1) \in \mathcal{K}$ for all $1 \le j \le p$ and finally $\zeta_{(i,i')} = \zeta_{(i_0,i_n-1)} \ge \sum_{j=1}^p \zeta_{(i_{j-1},i_j-1)}$.

An important note is that for a removed $(i, i') \in \mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}}$, we can always choose the indices i_1, \dots, i_{p-1} such that actually $(i_i, i_{i+1} - 1) \in \mathcal{K}^{\mathfrak{pr}}$ and not only \mathcal{K} , because if $(i_i, i_{i+1} - 1) \in \mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}}$ it can itself be fragmented, and this decreasing recursion eventually ends (the later possible being at the atoms of the forest structure). Also note that removing elements from $\mathcal K$ does not alter the fact that we 180 have at hand a forest structure, that is, the reference family defined by $\Re^{\mathfrak{pr}} = (R_k, \zeta_k)_{k \in \mathscr{K}^{\mathfrak{pr}}}$ has a forest structure. Because pruning a forest structure does not touch the atoms, note finally that if ${\mathcal X}$ is complete then so is $\mathcal{K}^{\mathfrak{pr}}$.

The following proposition states that pruning the forest does not alter the bound.

Proposition 3.1. For any $S \subseteq \mathbb{N}_m^*$, $V_{\mathfrak{R}}^*(S) = V_{\mathfrak{R}^{\mathfrak{pr}}}^*(S)$.

Proof. Recall Equation (11) and, because $\Re^{\mathfrak{pr}}$ also has a forest structure,

$$V_{\mathfrak{R}^{\mathfrak{p}_{\mathfrak{r}}}}^{\star}(S) = \min_{Q \subseteq \mathscr{X}^{\mathfrak{p}_{\mathfrak{r}}}} \left(\sum_{k' \in Q} \zeta_{k'} \wedge |S \cap R_{k'}| + \left| S \setminus \bigcup_{k' \in Q} R_{k'} \right| \right), \tag{13}$$

so we immediately get that $V_{\Re}^*(S) \leq V_{\Re^{\mathfrak{pr}}}^*(S)$.

Let any $Q \subseteq \mathcal{H}$. We split Q in A elements of $\mathcal{H} \setminus \mathcal{H}^{\mathfrak{pr}}$, denoted $(i_{0,a}, i_{p_a,a}-1), 1 \leq a \leq A$ for some $p_a \geq 2$, and B elements of $\mathcal{K}^{\mathfrak{pr}}$, simply denoted k_b , $1 \le b \le B$. By the definition of $\mathcal{K}^{\mathfrak{pr}}$ and the previous

remarks, for any $1 \leq a \leq A$, there exist integers $i_{1,a}, \ldots, i_{p_a-1,a}$ such that $i_{0,a} < i_{1,a} < \cdots < i_{p_a-1,a} < i_{p_a,a}$, $(i_{j-1,a},i_{j,a}-1) \in \mathscr{K}^{\mathfrak{pr}}$ for all $1 \leq j \leq p_a$, and $\zeta_{(i_{0,a},i_{p_a,a}-1)} \geq \sum_{j=1}^{p_a} \zeta_{(i_{j-1,a},i_{j,a}-1)}$. Now let

$$Q^{\mathfrak{pr}} = \{k_b : 1 \le b \le B\} \cup \{(i_{j-1,a}, i_{j,a} - 1) : 1 \le a \le A, 1 \le j \le p_a\}. \tag{14}$$

We have that $Q^{\mathfrak{pr}} \subseteq \mathscr{K}^{\mathfrak{pr}}$ and $\bigcup_{k \in Q} R_k = \bigcup_{k \in Q^{\mathfrak{pr}}} R_k$. Then,

$$\begin{split} \sum_{k \in Q} \zeta_k \wedge |S \cap R_k| + \left| S \setminus \bigcup_{k \in Q} R_k \right| &= \sum_{b=1}^B \zeta_{k_b} \wedge |S \cap R_{k_b}| \\ &+ \sum_{a=1}^A \zeta_{(i_{0,a}, i_{p_a,a} - 1)} \wedge |S \cap R_{(i_{0,a}, i_{p_a,a} - 1)}| \\ &+ \left| S \setminus \bigcup_{k \in Q} R_k \right|, \end{split}$$

but for all $1 \le a \le A$,

$$\begin{split} \zeta_{(i_{0,a},i_{p_{a},a}-1)} &\geq \sum_{j=1}^{p_{a}} \zeta_{(i_{j-1,a},i_{j,a}-1)} \\ &\geq \sum_{i=1}^{p_{a}} \zeta_{(i_{j-1,a},i_{j,a}-1)} \wedge |S \cap R_{(i_{j-1,a},i_{j,a}-1)}|, \end{split}$$

so the term $\sum_{a=1}^A \zeta_{(i_{0,a}i_{p_a,a}-1)} \wedge |S \cap R_{(i_{0,a}i_{p_a,a}-1)}|$ is greater than or equal to

$$\sum_{a=1}^{A} \left(\sum_{j=1}^{p_a} \zeta_{(i_{j-1,a},i_{j,a}-1)} \wedge |S \cap R_{(i_{j-1,a},i_{j,a}-1)}| \right) \wedge |S \cap R_{(i_{0,a},i_{p_a,a}-1)}|,$$

which is simply equal to

$$\sum_{a=1}^{A} \sum_{i=1}^{p_a} \zeta_{(i_{j-1,a},i_{j,a}-1)} \wedge |S \cap R_{(i_{j-1,a},i_{j,a}-1)}|.$$

Furthermore $\left| S \setminus \bigcup_{k \in Q} R_k \right| = \left| S \setminus \bigcup_{k \in Q^{\text{pr}}} R_k \right|$ so finally:

$$\sum_{k \in Q} \zeta_k \wedge |S \cap R_k| + \left| S \setminus \bigcup_{k \in Q} R_k \right| \ge \sum_{k \in Q^{\mathfrak{pr}}} \zeta_k \wedge |S \cap R_k| + \left| S \setminus \bigcup_{k \in Q^{\mathfrak{pr}}} R_k \right|$$

$$\ge V_{\mathfrak{R}^{\mathfrak{pr}}}^*(S).$$
(15)

Note that Equation (15) is true even if there are some $b \in \{1, \dots, B\}, a \in \{1, \dots, A\}, j \in \{1, \dots, p_a\}$ such that $k_b = (i_{j-1,a}, i_{j,a} - 1)$. We minimize over all Q to get that $V_{\mathfrak{R}}^*(S) \geq V_{\mathfrak{R}}^*\mathfrak{pr}(S)$. \square

This gives a practical way to speed up computations by first pruning the family before computing any $V_{\Re}^*(S)$, because $\mathscr{K}^{\mathfrak{pr}}$ is smaller than \mathscr{K} , and by the above Proposition there is no theoretical loss in doing so.

Furthermore, pruning can be done really simply by following Algorithm 1 for $S = \mathbb{N}_m^*$, and pruning when appropriate. This gives the following Algorithm 2, assuming, for simplicity, that the family is complete. The computation time of the algorithm is the same as Algorithm 1, that is $O(m|\mathcal{K}|)$. Note that the only differences between Algorithm 2 and Algorithm 1 are the pruning step and ζ_k replacing $\zeta_k \wedge |S \cap R_k|$, because $\zeta_k \leq |R_k|$ and here $S = \mathbb{N}_m^*$, so $\zeta_k \wedge |\mathbb{N}_m^* \cap R_k| = \zeta_k$. Also note that the algorithm returns $V_{\mathfrak{R}}^*(\mathbb{N}_m^*)$ as a by-product. The following proposition states that Algorithm 2 indeed produces the pruned region as in Definition 3.1.

Algorithm 2 Pruning of R

219

220

221

```
1: procedure Pruning(\Re = (R_k, \zeta_k)_{k \in \mathcal{K}} with \Re complete)
 2:
            H \leftarrow \max_{k \in \mathcal{K}} \phi(k)
                                                                                                                                          ⊳ maximum depth
 3:
             for h = H - 1, ..., 1 do
 4:
                   \mathcal{K}^h \leftarrow \{k \in \mathcal{K} : \phi(k) = h\}
  5:
                   newVec \leftarrow (0)_{k \in \mathcal{K}^h}
 6:
                   for k \in \mathcal{K}^h do
 7:
                         Succ_k \leftarrow \{k' \in \mathcal{K}^{h+1} : R_{k'} \subseteq R_k\}
 8:
                         if Succ_k = \emptyset then
 9:
                               newVec_k \leftarrow \zeta_k
10:
                         else
11:
                               if \zeta_k \geq \sum_{k' \in Succ_k} Vec_{k'} then
12:
                                      \mathscr{L} \leftarrow \mathscr{L} \setminus \{k\}
                                                                                                          \triangleright pruning of the region indexed by k
13:
14:
                               newVec_k \leftarrow \min\left(\zeta_k, \sum_{k' \in Succ_k} Vec_{k'}\right)
15:
                         end if
16:
                   end for
17:
                   Vec \leftarrow newVec
18:
             end for
19:
            return (\mathcal{L}, \sum_{k \in \mathcal{K}^1} Vec_k)
20:
21: end procedure
```

Proposition 3.2. The final \mathcal{L} returned by Algorithm 2 is equal to $\mathcal{K}^{\mathfrak{pr}}$: $\mathcal{L} = \mathcal{K}^{\mathfrak{pr}}$

```
Proof. First, \mathcal{K} \setminus \mathcal{L} \subseteq \mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}} is trivial: a k such that \zeta_k \geq \sum_{k' \in Succ_k} Vec_{k'} obviously satisfies the
     condition of Definition 3.1 to be pruned.
211
     Now let (i,i') \in \mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}} an element that is pruned by Definition 3.1, so there exists p \geq 2
     and integers i_1, \dots, i_{p-1} such that, when setting i_0 = i and i_p = i' + 1, the sequence (i_0, \dots, i_p) is
     strictly increasing, (i_{j-1}, i_j - 1) \in \mathcal{K} for all 1 \le j \le p and finally \zeta_{(i,i')} = \zeta_{(i_0,i_p-1)} \ge \sum_{j=1}^p \zeta_{(i_{j-1},i_j-1)}.
214
     Then by the proof of Theorem 1 of Durand et al. (2020) but applied to S = R_{(i,i')} we have that
     \sum_{j=1}^{p} \zeta_{(i_{j-1},i_{j}-1)} \ge \sum_{k' \in Succ_{(i,i')}} Vec_{k'} \text{ (see the unnumbered line just above Equation (A4) in that paper)}
```

In the end, $\mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}} = \mathcal{K} \setminus \mathcal{L}$ so $\mathcal{K}^{\mathfrak{pr}} = \mathcal{L}$. \square

and so $\zeta_{(i,i')} \geq \sum_{k' \in Succ_{(i,i')}} Vec_{k'}$ hence (i,i') is pruned by Algorithm 2 and $\mathcal{K} \setminus \mathcal{K}^{\mathfrak{pr}} \subseteq \mathcal{K} \setminus \mathcal{L}$.

Fast algorithm to compute a curve of confidence bounds on a path of selection sets

Let $(i_1, ..., i_m)$ a permutation of \mathbb{N}_m^* , eventually random, and, for all $t \in \mathbb{N}_m^*$, let $S_t = \{i_1, ..., i_t\}$ and $S_0 = \emptyset$. For example, (i_1, \dots, i_m) can be the permutation ordering the p-values in increasing order and in that case S_t becomes the set of indices of the t smallest p-values. Assume that we want to compute all $V_{\mathfrak{R}}^{*}(S_{t})$ for all $t \in \{0, ..., m\}$, this is what we call the curve of confidence bounds indexed 224 by (i_1, \dots, i_m) . Applying Algorithm 1 to compute $V_{\mathfrak{R}}^*(S_t)$ for a given t has complexity $O(t|\mathcal{X}|)$, so using it to sequentially compute the full curve has complexity $O\left(\sum_{t=0}^{m} t |\mathcal{X}|\right) = O\left(m^2 |\mathcal{X}|\right)$. In this section, 226 we present a new algorithm that computes the curve with a $O(m|\mathcal{K}|)$ complexity. The algorithm will need that \Re is complete, so if that is not the case we first need to complete \Re following the Algorithm 2 of Durand et al. (2020), which has a $O(m|\mathcal{K}|)$ complexity. In the remainder of this section

we assume that \Re is complete.

We first recall and introduce some notation. Recall that ϕ is the depth function inside of \Re , that $\Re \subseteq \mathscr{P}(\mathscr{K})$ is the set of subsets of \mathscr{K} that realize a partition, recall the important result stated by Equation (12), and that $\mathscr{K}^h = \{k \in \mathscr{K} : \phi(k) = h\}$ for all $1 \le h \le H$ where $H = \max_{k \in \mathscr{K}} \phi(k)$. For any $t \in \mathbb{N}_m^*$ and $1 \le h \le H$, we denote by $k^{(t,h)}$ the element of \mathscr{K}^h such that $i_t \in R_{k^{(t,h)}}$ if it exists, and we denote by $h_{\max}(t)$ the highest h such that $k^{(t,h)}$ exists.

Example 3.1 (Continuation of Example 2.2 and Example 2.3). Assume that the reference family of Example 2.2 has been labeled as in Example 2.3 and completed. Let $(i_1, ..., i_{25})$ such that $i_1 = 7$, $i_2 = 1$ and $i_3 = 24$. Then for t = 1, $k^{(t,1)} = (1,5)$, $k^{(t,2)} = (2,3)$, $k^{(t,3)} = (3,3)$ and $h_{\max}(t) = H = 3$. For t = 2, $k^{(t,1)} = (1,5)$, $k^{(t,2)} = (1,1)$, $k^{(t,3)}$ does not exist and $h_{\max}(t) = 2$. For t = 3, $k^{(t,1)} = (8,8)$, $k^{(t,2)}$ does not exist and $h_{\max}(t) = 1$.

Now we can finally present the new algorithm and the proof that it computes the curve $(V_{\Re}^*(S_t))_{t\in\mathbb{N}_m}$. We present two versions of the algorithm (strictly equivalent): one very formal (Algorithm 3), to introduce additional notation used in the proof of Theorem 3.1, and, later, a simpler version that is the one actually implemented (Algorithm 4). Recall that a detailed illustration of the steps of the algorithms will be provided in Section 3.3.

The core idea of the algorithm is that, as we increase t and add new hypotheses in S_t , we inflate a counter η_k^t for each region R_k , by 1 if $i_t \in R_k$ (line 12), by 0 if not (lines 23 and 27), but only until the counter reaches ζ_k (line 13). After this point, the hypotheses in R_k don't contribute to $V_{\mathfrak{R}}^*(S_t)$, we keep track of those hypotheses with \mathcal{K}_t^- (line 6), so as soon as $\eta_{k(t,h)}^t = \zeta_k$ we update \mathcal{K}_t^- by adding $k^{(t,h)}$ (line 18) to it and we update \mathcal{S}_t^t accordingly (line 17).

We will see in the following Theorem 3.1 how this algorithm allows to compute $V_{\mathfrak{R}}^*(S_t)$. We first need a final notation. Let

$$\mathcal{K}_t = \{k \in \mathcal{K} : \exists k' \in \mathcal{P}^t : R_{k'} \subseteq R_k\}.$$

The elements of \mathcal{K}_t index the regions of the forest that "are above" the regions of the current partition-realizing \mathcal{P}^t . In particular, we always have, for any $t \in \mathbb{N}_m$, $\mathcal{K}^1 \subseteq \mathcal{K}_t$ and $\mathcal{P}^t \subseteq \mathcal{K}_t$. We can also remark that the sequence $(\mathcal{K}_t)_{0 \le t \le m}$ is non-increasing for the inclusion relation, and that $\mathcal{K}_0 = \mathcal{K}$.

Theorem 3.1 (Fast curve computation). Let any $t \in \mathbb{N}_m$. Then, $\mathscr{P}^t \in \mathfrak{P}$, and for all $k \in \mathscr{K}_t$, we have

$$V_{\mathfrak{R}}^*(S_t \cap R_k) = \eta_k^t \tag{16}$$

258 and

$$V_{\mathfrak{R}}^{\star}(S_t \cap R_k) = \sum_{\substack{k' \in \mathscr{D}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}|. \tag{17}$$

Furthermore,

$$V_{\mathfrak{R}}^{\star}(S_t) = \sum_{k \in \mathscr{P}^t} \zeta_k \wedge |S_t \cap R_k| = \sum_{k \in \mathscr{K}^1} \eta_k^t.$$
 (18)

The proof of this Theorem is postponed to Section 3.4. The first equality of Equation (18) states that the minimum in (12) is realized on the partition \mathcal{P}^t , and the last equality of the same Equation is the basis of the following light corollary.

Corollary 3.1 (Easy computation). For $t \in \{0, ..., m-1\}$, $V_{\Re}^*(S_{t+1}) = V_{\Re}^*(S_t)$ if $i_{t+1} \in \bigcup_{k \in \mathscr{K}_t^-} R_k$, and $V_{\Re}^*(S_{t+1}) = V_{\Re}^*(S_t) + 1$ if not.

Algorithm 3 Formal computation of $(V_{\mathfrak{R}}^*(S_t))_{0 \le t \le m}$

```
1: procedure Curve(\Re = (R_k, \zeta_k)_{k \in \mathcal{X}} with \Re complete, path (S_t)_{1 \le t \le m} with S_t = \{i_1, \dots, i_t\})
             \mathcal{P}^0 \leftarrow \{(i,i) : 1 \le i \le n\}
                                                                                                                               > the set of all atoms indices
 2:
             \mathcal{K}_0^- \leftarrow \{k \in \mathcal{K} : \zeta_k = 0\}
 3:
             \eta_k^0 \leftarrow 0 for all k \in \mathcal{K}
  4:
             \mathbf{for}\ t = 1, \dots, m\ \mathbf{do}
  5:
                   if i_t \in \bigcup_{k \in \mathcal{K}_{t-1}^-} R_k then
  6:
                          \mathscr{P}^t \leftarrow \mathscr{P}^{t-1}
 7:
                          \begin{aligned} \mathcal{K}_t^- &\leftarrow \mathcal{K}_{t-1}^- \\ \eta_k^t &\leftarrow \eta_k^{t-1} \text{ for all } k \in \mathcal{K} \end{aligned}
 8:
 9:
10:
                          for h = 1, ..., h_{\max}(t) do
11:
                                \eta_{k^{(t,h)}}^t \leftarrow \eta_{k^{(t,h)}}^{t-1} + 1
12:
                                 if \eta_{k^{(t,h)}}^t < \zeta_k then
13:
14:
15:
                                       h_t^f \leftarrow h
                                                                                                                                                            ⊳ final depth
16:
                                       \mathcal{P}^t \leftarrow \left(\mathcal{P}^{t-1} \setminus \{k \in \mathcal{P}^{t-1} : R_k \subseteq R_{k^{(t,h_t^f)}}\}\right) \cup \{k^{(t,h_t^f)}\}
17:
                                       \mathcal{K}_t^- \leftarrow \mathcal{K}_{t-1}^- \cup \{k^{(t,h_t^f)}\}
18:
                                        Break the loop
19:
                                 end if
20:
                           end for
21:
                          if the loop has been broken then
22:
                                 \eta_k^t \leftarrow \eta_k^{t-1} for all k \in \mathcal{K} not visited during the loop, that is all k \notin \{k^{(t,h)}, 1 \leq h \leq t \leq t \}
23:
      h_t^J
                          else
24:
                                 \mathscr{P}^t \leftarrow \mathscr{P}^{t-1}
25:
                                 26:
27:
      h_{\max}(t)
                           end if
28:
                    end if
29:
30:
             return \mathcal{P}^t, \eta_k^t for all t = 1, ..., m and k \in \mathcal{K}
32: end procedure
```

```
Proof. From (18), V_{\mathfrak{R}}^{*}(S_{t+1}) = \sum_{k \in \mathcal{X}^{1}} \eta_{k}^{t+1} and V_{\mathfrak{R}}^{*}(S_{t}) = \sum_{k \in \mathcal{X}^{1}} \eta_{k}^{t}. If i_{t+1} \in \bigcup_{k \in \mathcal{K}_{t}^{-}} R_{k}, \eta_{k}^{t+1} = \eta_{k}^{t} for all k \in \mathcal{K}^{1}. If not, \eta_{k}^{t+1} = \eta_{k}^{t} for all k \in \mathcal{K}^{1}, k \neq k^{(t+1,1)}, whereas for k = k^{(t+1,1)}, \eta_{k}^{t+1} = \eta_{k}^{t} + 1. \square
```

We note that, from Theorem 3.1 and Corollary 3.1, if one is only interested in the computation of the curve $(V^*_{\Re}(S_t))_{1 \leq t \leq m}$, tracking \mathscr{D}^t is actually useless, what is important is to track and update \mathscr{K}^-_t correctly. Hence the simpler, alternative Algorithm 4 . Note that Algorithm 4 is less formal than Algorithm 3: as in Algorithm 1 and Algorithm 2, it recycles notation (mimicking the actual code implementation) so the t subscript or superscript is dropped from the \mathscr{K}^-_t and the η^t_k . In Algorithm 4, the notation V_t is actually equal to $V^*_{\Re}(S_t)$ by Corollary 3.1.

It is easy to see that each step t has a complexity in $O(|\mathcal{K}|)$ hence the total complexity is in $O(m|\mathcal{K}|)$. This is because, if the regions are carefully stocked in memory, especially if their bounds (in terms

Algorithm 4 Practical computation of $(V_{\mathfrak{R}}^*(S_t))_{0 \le t \le m}$

```
1: procedure Curve(\Re = (R_k, \zeta_k)_{k \in \mathcal{X}} with \Re complete, path (S_t)_{1 \le t \le m} with S_t = \{i_1, \dots, i_t\})
 2:
             \mathcal{K}^- \leftarrow \{k \in \mathcal{K} : \zeta_k = 0\}
 3:
            \eta_k \leftarrow 0 \text{ for all } k \in \mathcal{K}
  4:
            for t = 1, ..., m do
  5:
                   if i_t \in \bigcup_{k \in \mathcal{K}^-} R_k then
 6:
                         V_t \leftarrow V_{t-1}
 7:
                   else
 8:
                         for h = 1, ..., h_{\max}(t) do
 9:
                               find k^{(t,h)} \in \mathcal{K}^h such that i_t \in R_{k^{(t,h)}}
10:
                               \eta_{k(t,h)} \leftarrow \eta_{k(t,h)} + 1
                               if \eta_{k(t,h)} < \zeta_k then
12:
                                      pass
13:
14:
                                      \mathcal{K}^- \leftarrow \mathcal{K}^- \cup \{k^{(t,h)}\}\
15:
                                      break the loop
16:
                               end if
17:
                         end for
18:
                         V_t \leftarrow V_{t-1} + 1
19:
20:
                   end if
             end for
21:
             return (V_t)_{1 \le t \le m}
23: end procedure
```

of hypothesis index) are stocked, then finding $k^{(t,h)}$ has a complexity in $O(|\mathcal{K}^h|)$ and checking if $i_t \in \bigcup_{k \in \mathcal{K}_{t-1}^-} R_k$ has a complexity in $O(|\mathcal{K}|)$.

3.3 Illustration on a detailed example

```
We still continue Example 2.2 and Example 2.3. Recall that m=25, P_{1:5}=R_1=\{1,\ldots,20\}, P_1=\{1,\ldots,20\}
          R_2 = \{1,2\}, \ P_{2:3} = R_3 = \{3,\dots,10\}, \ P_{4:5} = R_4 = \{11,\dots,20\}, \ P_2 = \{3,4\}, \ P_3 = R_5 = \{5,\dots,10\}, \ P_{4:5} = \{1,2\}, \ P_{4:5} = \{1,2
           P_8 = \{23, 24, 25\}.
           Now assume that we have the following values for the \zeta_k's: \zeta_{(1,5)} = 6, \zeta_{(1,1)} = 2, \zeta_{(2,3)} = 1, \zeta_{(3,3)} = 4,
           \zeta_{(4,5)} = 4, \zeta_{(4,4)} = 2, \zeta_{(5,5)} = 3, \zeta_{(6,7)} = 2, \zeta_{(7,7)} = 0. Because P_2, P_6 and P_8 come from the completion
           operation (see Section 2.3), we also have \zeta_{(2,2)} = |P_2| = 2, \zeta_{(6,6)} = |P_6| = 1 and \zeta_{(8,8)} = |P_8| = 3. Theses
           values are summarized in Figure 3.
          We want to compute the curve (V_{\mathfrak{R}}^{*}(S_{t}))_{1 \leq t \leq 9} with S_{t} = \{i_{1}, \dots, i_{t}\} and i_{1} = 11, i_{2} = 17, i_{3} = 12, i_{4} = 13, \dots, i_{t}\}
286
           i_5 = 18, i_6 = 3, i_7 = 19, i_8 = 22 and i_9 = 5.
           First, we apply Algorithm 2 to the family. This results in pruning P_{6:7} (and only this region), because
           2 = \zeta_{(6,7)} \ge \zeta_{(6,6)} + \zeta_{(7,7)} = 1 + 0. This gives Figure 4.
           Now we initialize Algorithm 3, that is we let t = 0. Because \zeta_{(7,7)} = 0, (7,7) is added to \mathcal{X}_t^-:
           \mathcal{K}_0^- = \{(7,7)\}. Furthermore, all \eta_k^t are set to 0. The initial state of Algorithm 3 is shown in Figure 5,
           with (7,7) being in red to show that it will not contribute to the computations.
          We move on to t = 1, with i_1 = 11. i_1 \in P_4 \subseteq P_{4:5} \subseteq P_{1:5}. The appropriate \eta_k^t are increased by one,
          and by (18) we have V_{\Re}^*(S_1) = \eta_{(1,5)}^1 + \eta_{(6,6)}^1 + \eta_{(7,7)}^1 + \eta_{(8,8)}^1 = 1 + 0 + 0 + 0 = 1. The state of the step
```

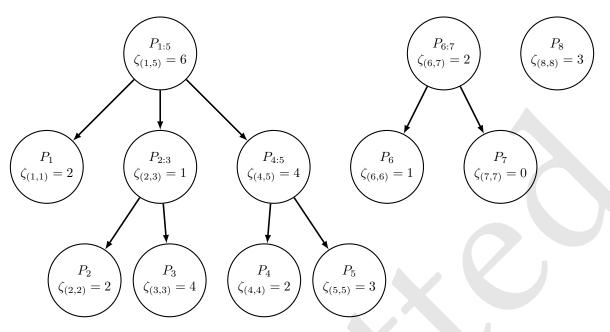


Figure 3: The regions of Example 2.2 with the ζ_k values.

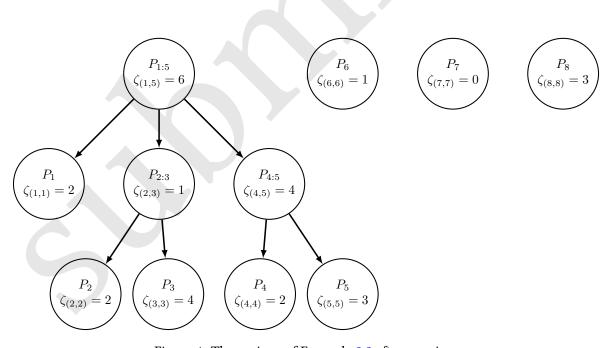


Figure 4: The regions of Example 2.2 after pruning.

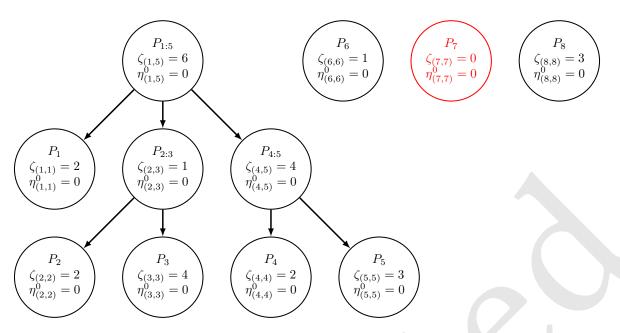


Figure 5: The regions of Example 2.2 at t = 0 in Algorithm 3.

is summarized in Figure 6.

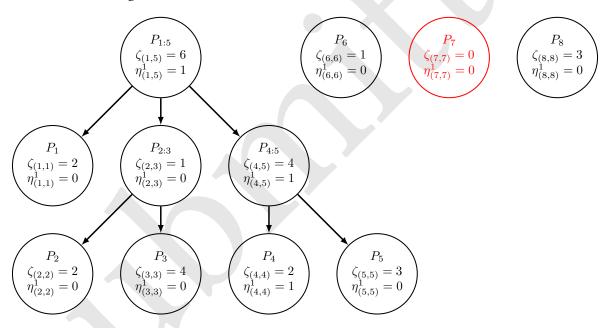


Figure 6: The regions of Example 2.2 at t = 1 in Algorithm 3.

We move on to t=2, with $i_2=17$. $i_1 \in P_5 \subseteq P_{4:5} \subseteq P_{1:5}$. The appropriate η_k^t are increased by one, and by (18) we have $V_{\Re}^*(S_2)=2$. The state of the step is summarized in Figure 7.

We move on to t=3, with $i_3=12$. $i_3\in P_4\subseteq P_{4:5}\subseteq P_{1:5}$. The appropriate η_k^t are increased by one, and we notice that $\eta_{(4,4)}^3=2=\zeta_{(4,4)}$. So P_4 will stop contributing, we add it to $\mathcal{K}_t^-:\mathcal{K}_3^-=\{(4,4),(7,7)\}$.

By (18), we have $V_{\mathfrak{R}}^{(3,3)}(S_3)=3$. The state of the step is summarized in Figure 8, with P_4 now also in red.

We move on to t=4, with $i_4=13$. $i_4\in P_4\in\bigcup_{k\in\mathscr{K}_3^-}R_k$. No η_k^t is increased (see line 9 of Algorithm 3), and by (18), we have $V_{\Re}^*(S_4)=3$.

We move on to t=5, with $i_5=18$. $i_5\in P_5\subseteq P_{4:5}\subseteq P_{1:5}$. We first increase $\eta^t_{(1,5)}$: $\eta^5_{(1,5)}=4<\zeta_{(1,5)}$,

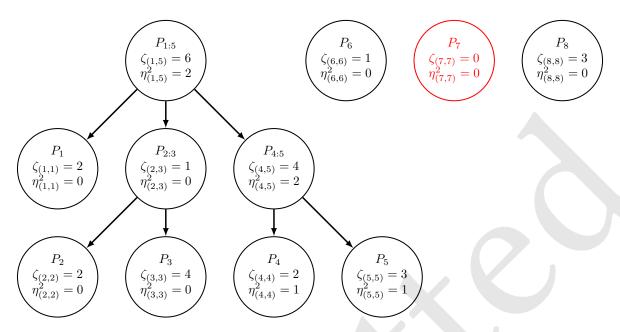


Figure 7: The regions of Example 2.2 at t = 2 in Algorithm 3.

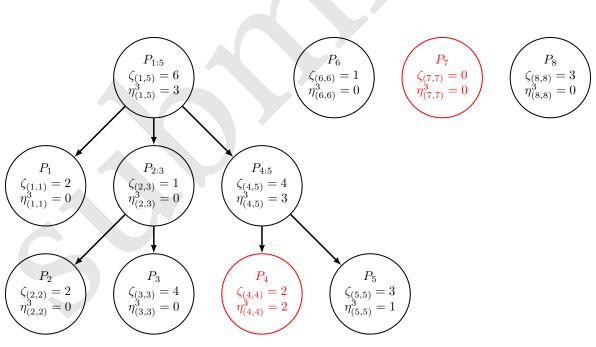


Figure 8: The regions of Example 2.2 at t=3 in Algorithm 3 .

then $\eta^t_{(4,5)}$: $\eta^5_{(4,5)} = 4$, and we stop there because $\eta^5_{(4,5)} = 4 = \zeta_{(4,5)}$. $P_{4:5}$ will stop contributing, we add it to \mathcal{K}_t^- : $\mathcal{K}_5^- = \{(4,5), (4,4), (7,7)\}$. Note that $\eta^t_{(5,5)}$ is not updated because we stopped the loop before, see line 23 of Algorithm 3 . By (18), we have $V^*_{\mathfrak{R}}(S_5) = 4$. The state of the step is summarized in Figure 9, with $P_{4:5}$ now also in red.

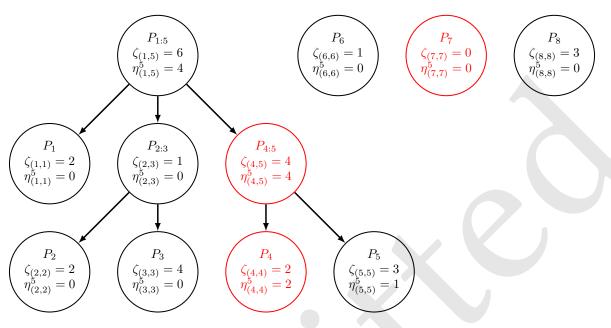


Figure 9: The regions of Example 2.2 at t = 5 in Algorithm 3.

We move on to t=6, with $i_6=3$. $i_6\in P_3\subseteq P_{2:3}\subseteq P_{1:5}$. We first increase $\eta^t_{(1,5)}\colon \eta^6_{(1,5)}=5<\zeta_{(1,5)},$ then $\eta^t_{(2,3)}\colon \eta^6_{(2,3)}=1$, and we stop there because $\eta^6_{(2,3)}=1=\zeta_{(2,3)}$. $P_{2:3}$ will stop contributing, we add it to $\mathcal{K}^+_t\colon \mathcal{K}^-_6=\{(2,3),(4,5),(4,4),(7,7)\}$. Note that $\eta^t_{(3,3)}$ is not updated because we stopped the loop before, see line 23 of Algorithm 3 . By (18), we have $V^*_{\mathfrak{R}}(S_6)=5$. The state of the step is summarized in Figure 10, with $P_{2:3}$ now also in red.

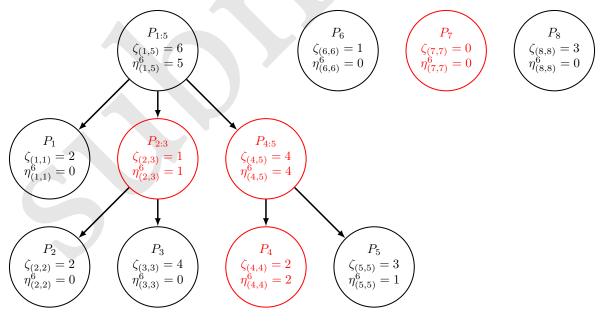


Figure 10: The regions of Example 2.2 at t = 6 in Algorithm 3.

We move on to the remaining steps. $i_7=19\in P_{4:5},\,i_8=22\in P_7$ and $i_9=5\in P_{2:3}$ are all

in $\bigcup_{k \in \mathcal{K}_6^-} R_k$ so no η_k^t is increased at their step (see line 9 of Algorithm 3), and by (18), we have $V_{\mathfrak{R}}^*(S_7) = V_{\mathfrak{R}}^*(S_8) = V_{\mathfrak{R}}^*(S_9) = 5$.

3.4 Proof of Theorem 3.1

3.4.1 Derivation of (18)

We first derive (18) from (16) and (17). First note that for all $Q \in \mathfrak{P}$,

$$Q = \bigcup_{k \in \mathcal{K}^1} \{ k' \in Q : R_{k'} \subseteq R_k \}$$

$$\tag{19}$$

and the union is disjoint. From (12), let $Q^* \in \mathfrak{P}$ such that $V_{\mathfrak{R}}^*(S_t) = \sum_{k' \in Q^*} \zeta_{k'} \wedge |S_t \cap R_{k'}|$. Then by (19),

$$V_{\mathfrak{R}}^{*}(S_{t}) = \sum_{k' \in \mathcal{Q}^{*}} \zeta_{k'} \wedge |S_{t} \cap R_{k'}|$$

$$= \sum_{k \in \mathcal{H}^{1}} \sum_{\substack{k' \in \mathcal{Q}^{*} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t} \cap R_{k'}|$$

$$= \sum_{k \in \mathcal{H}^{1}} \sum_{\substack{k' \in \mathcal{Q}^{*} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t} \cap (R_{k} \cap R_{k'})|$$

$$= \sum_{k \in \mathcal{H}^{1}} \sum_{\substack{k' \in \mathcal{Q}^{*} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |(S_{t} \cap R_{k}) \cap R_{k'}|$$

$$\geq \sum_{k \in \mathcal{H}^{1}} V_{\mathfrak{R}}^{*}(S_{t} \cap R_{k}), \tag{20}$$

where the equality in (20) comes from the fact that if $R_{k'} \nsubseteq R_k$, then $R_{k'} \cap R_k = \emptyset$, that is, $R_k \subseteq R_{k'}$ is impossible because $k \in \mathcal{K}^1$. Furthermore, (21) holds again by (12).

Because $\mathscr{K}^1 \subseteq \mathscr{K}_t$, by (17), $V_{\mathfrak{R}}^*(S_t \cap R_k) = \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}|$ for all $k \in \mathscr{K}^1$. Then,

$$\sum_{k \in \mathcal{X}^1} V_{\mathfrak{R}}^*(S_t \cap R_k) = \sum_{k \in \mathcal{X}^1} \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}|$$

$$= \sum_{k \in \mathcal{P}^t} \zeta_k \wedge |S_t \cap R_k| \text{ by (19)}$$

$$\geq V_{\mathfrak{R}}^*(S_t) \text{ by (12)}.$$

So we proved that $V_{\mathfrak{R}}^{*}(S_{t}) = \sum_{k \in \mathscr{T}^{t}} \zeta_{k} \wedge |S_{t} \cap R_{k}| = \sum_{k \in \mathscr{K}^{1}} V_{\mathfrak{R}}^{*}(S_{t} \cap R_{k})$ and finally $V_{\mathfrak{R}}^{*}(S_{t}) = \sum_{k \in \mathscr{K}^{1}} V_{\mathfrak{R}}^{*}(S_{t} \cap R_{k}) = \sum_{k \in \mathscr{K}^{1}} \eta_{k}^{t}$ by (16), again because $\mathscr{K}^{1} \subseteq \mathscr{K}_{t}$. Every equality in (18) is proven.

26 3.4.2 Proof of (16) and (17)

We show the remainder of the statements by a strong recursion over t. We have $\mathscr{P}^0 \in \mathfrak{P}$ by definition, and given that $S_0 = \emptyset$ and $\eta_k^0 = 0$ for all $k \in \mathscr{K}$ (recall that $\mathscr{K}_0 = \mathscr{K}$), everything is equal to 0 in (16) and (17).

So we let $t \in \{0, ..., m-1\}$, and assume that $\mathscr{P}^{t'} \in \mathfrak{P}$ and that (16) and (17) hold for all $t' \leq t$.

In all the following, \bar{k} is the element of \mathscr{P}^t such that $i_{t+1} \in R_{\bar{k}}$. We will distinguish two cases: if

 $i_{t+1} \in \bigcup_{k \in \mathcal{K}_t^-} R_k$ or not. First we show an inequality that will be used in both cases. We have, for all $k \in \mathcal{K}_t$,

$$V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_k) \leq \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|. \tag{22}$$

334 Indeed, by (12),

$$V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_k) \leq \sum_{k' \in \mathscr{D}^t} \zeta_{k'} \wedge |S_{t+1} \cap R_k \cap R_{k'}|.$$

For any $k' \in \mathcal{P}^t$, we have either $R_{k'} \cap R_k = \emptyset$, in which case $|S_{t+1} \cap R_k \cap R_{k'}| = 0$, either $R_{k'} \subseteq R_k$, in which case $|S_{t+1} \cap R_k \cap R_{k'}| = |S_{t+1} \cap R_{k'}|$, but $R_k \subseteq R_{k'}$ is impossible. Indeed, by definition of \mathcal{K}_t , there exists $\tilde{k} \in \mathcal{P}^t$ such that $R_{\tilde{k}} \subseteq R_k$, so $R_k \subseteq R_{k'}$ would entail $R_{\tilde{k}} \subseteq R_{k'}$ which is impossible since $k', \tilde{k} \in \mathcal{P}^t \in \mathfrak{P}$ and so $R_{\tilde{k}}$ and $R_{k'}$ are part of a partition of \mathbb{N}_m^* . This gives (22).

339 **3.4.2.1** First case: $i_{t+1} \in \bigcup_{k \in \mathcal{K}_t} R_k$

In this case, $\mathscr{P}^{t+1} = \mathscr{P}^t \in \mathfrak{P}$ and $\mathscr{K}_{t+1} = \mathscr{K}_t$. For any $k \in \mathscr{K}_{t+1}$ such that $i_{t+1} \notin R_k$ (or, equivalently, such that $S_{t+1} \cap R_k = S_t \cap R_k$),

$$\sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| = \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}|$$

$$= V_{\Re}^* (S_t \cap R_k) \text{ by (17)}$$

$$= \eta_k^t \text{ by (16)}$$

$$= \eta_k^{t+1}$$

because $\eta_k^t = \eta_k^{t+1}$ for all $k \in \mathcal{K}$. Furthermore $S_{t+1} \cap R_k = S_t \cap R_k$ so $V_{\mathfrak{R}}^*(S_{t+1} \cap R_k) = V_{\mathfrak{R}}^*(S_t \cap R_k)$. So everything is proved for such a k.

Now we let $k \in \mathcal{K}_{t+1}$ such that $i_{t+1} \in R_k$ or, equivalently, such that $R_{\bar{k}} \subseteq R_k$. We first need to show that $\zeta_{\bar{k}} \le |S_t \cap R_{\bar{k}}|$, and for that we need to distinguish two subcases: if \bar{k} has been added to \mathcal{P}^t during a previous step of the algorithm, of if not.

$_{347}$ $\,$ 3.4.2.1.1 $\,$ First subcase: $ar{k}$ has never been added during the process of line 17

Then $\bar{k} \in \mathscr{P}^0$ and $R_{\bar{k}}$ is an atom, so $i_{t+1} \in \bigcup_{k' \in \mathscr{K}_t^-} R_{k'}$ implies that $R_{\bar{k}} \subseteq \bigcup_{k' \in \mathscr{K}_t^-} R_{k'}$ (because of the forest structure). Let k'_{\max} such that

$$R_{k'_{max}} = \max\{R_{k'} : k' \in \mathcal{K}_t^-, R_{\bar{k}} \subseteq R_{k'}\}$$

(this a maximum for the inclusion relation, and it is well defined thanks to the forest structure). By reductio ad absurdum we show that $k'_{\max} = \bar{k}$. If that wasn't the case, by the joint construction of \mathscr{P}^t and \mathscr{K}^-_t during the algorithm we would have $k'_{\max} \in \mathscr{P}^t$ and a contradiction with the fact that $\mathscr{P}^t \in \mathfrak{P}$: we can't have both $\bar{k} \in \mathscr{P}^t$ and $k'_{\max} \in \mathscr{P}^t$ if they are distinct. So $k'_{\max} = \bar{k}$, so $\bar{k} \in \mathscr{K}^-_t$, but it cannot have been added to \mathscr{K}^-_t during a previous step of the algorithm, otherwise it would have been added to \mathscr{P}^t , too. Hence $\bar{k} \in \mathscr{K}^-_0$ which means that $\zeta_{\bar{k}} = 0$ and $\zeta_{\bar{k}} = 0 \leq |S_t \cap R_{\bar{k}}|$.

3.4.2.1.2 Second subcase: \bar{k} has been added to \mathcal{P}^t at a previous step

Let $t' \leq t$ be this step. This means that $\bar{k} = k^{(t',h^f_{t'})}$ and that at that step $\eta^{t'}_{\bar{k}} = \zeta_{\bar{k}}$. Indeed, the if condition in line 13 failed so $\eta^{t'}_{\bar{k}} \geq \zeta_{\bar{k}}$, but for all t'' < t' we had $\eta^{t''}_{\bar{k}} \leq \zeta_{\bar{k}}$ which implies equality. Also $\bar{k} \in \mathscr{P}^{t'}$ so $\bar{k} \in \mathscr{H}_{t'}$ so we can write

$$\zeta_{\bar{k}} = \eta_{\bar{k}}^{t'}$$

$$= V_{\mathfrak{R}}^{*}(S_{t'} \cap R_{\bar{k}}) \text{ by (16)}$$

$$\leq |S_{t'} \cap R_{\bar{k}}|$$

$$\leq |S_{t} \cap R_{\bar{k}}|.$$

This concludes the two subcases dichotomy: $\zeta_{\bar{k}} \leq |S_t \cap R_{\bar{k}}|$ and we can go back to our $k \in \mathcal{K}_{t+1}$ such that $i_{t+1} \in R_k$ and $R_{\bar{k}} \subseteq R_k$.

We write the following chain:

$$\begin{split} V^*_{\mathfrak{R}}(S_{t+1} \cap R_k) &\leq \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \text{ by (22) and } \mathscr{K}_{t+1} \subseteq \mathscr{K}_t \\ &= \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_{t+1} \cap R_{\bar{k}}| \\ &= \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge (|S_t \cap R_{\bar{k}}| + 1) \\ &= \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_t \cap R_{\bar{k}}| \text{ because } \zeta_{\bar{k}} \leq |S_t \cap R_{\bar{k}}| \\ &= \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\ &= V^*_{\mathfrak{R}}(S_t \cap R_k) \text{ by (17)} \\ &= \eta_k^t \text{ by (16)} \\ &= \eta_k^{t+1}. \end{split}$$

But on the other hand, $S_t \subseteq S_{t+1}$ and so (12) also gives $V_{\mathfrak{R}}^*(S_t \cap R_k) \leq V_{\mathfrak{R}}^*(S_{t+1} \cap R_k)$ and so in the end we have the desired outcome:

$$V_{\mathfrak{R}}^{*}(S_{t+1} \cap R_{k}) = \eta_{k}^{t+1} = \sum_{\substack{k' \in \mathscr{D}^{t+1} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|,$$

which concludes this first case.

3.4.2.2 Second case: $i_{t+1} \notin \bigcup_{k \in \mathcal{K}_t^-} R_k$

We first prove that $\mathcal{P}^{t+1} \in \mathfrak{P}$ whether it came form the adjustment in line 17 or not. If it didn't, it stayed equal to $\mathcal{P}^t \in \mathfrak{P}$. If it did, we have

$$\mathscr{P}^{t+1} = \left(\mathscr{P}^t \setminus \{k \in \mathscr{P}^t, R_k \subseteq R_{k^{(t+1,h_{t+1}^f)}}\} \right) \cup \{k^{(t+1,h_{t+1}^f)}\}. \tag{23}$$

To prove that $\mathscr{P}^{t+1} \in \mathfrak{P}$ in that case, it suffices to prove there are no $k' \in \mathscr{P}^t$ such that $R_{k^{(t+1,h_{t+1}^f)}} \subseteq R_{k'}$.

If it was the case, because of the strict inclusion, we would have $k' \notin \mathscr{P}^0$, so k' would have been added to $\mathscr{P}^{t'}$ at a previous step $t' \leq t$ of the algorithm, but in that case it would also have been added to $\mathscr{K}_{t'} \subseteq \mathscr{K}_{t}^-$. So in the end we would have

$$i_{t+1} \in R_{k^{(t+1,h_{t+1}^f)}} \subseteq R_{k'} \subseteq \bigcup_{k \in \mathscr{K}_{-}^-} R_k$$

which is a contradiction and so $\mathscr{P}^{t+1} \in \mathfrak{P}$.

Like in the first case, considering a $k \in \mathcal{K}_{t+1} \subseteq \mathcal{K}_t$ such that $i_{t+1} \notin R_k$ is not problematic, because in that case k is not visited at all by the algorithm at step $t+1: \eta_k^{t+1} = \eta_k^t, \{k' \in \mathcal{P}^{t+1}: R_{k'} \subseteq R_k\} = \{k' \in \mathcal{P}^t: R_{k'} \subseteq R_k\}$, and for all $k' \in \mathcal{K}$ such that $R_{k'} \subseteq R_k, S_{t+1} \cap R_{k'} = S_t \cap R_{k'}$. Hence, from

$$V_{\mathfrak{R}}^{\star}(S_t \cap R_k) = \eta_k^t = \sum_{\substack{k' \in \mathscr{P}^t \\ R_{k'} \subset R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|,$$

we directly have

$$V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_k) = \eta_k^{t+1} = \sum_{\substack{k' \in \mathscr{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|.$$

So we now focus on the $k \in \mathcal{K}_{t+1}$ such that $i_{t+1} \in R_k$. Note that for such k,

$$\eta_k^{t+1} = \eta_k^t + 1 = V_{\Re}^*(S_t \cap R_k) + 1 = \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + 1$$

by construction, by (16) and by (17). Indeed, such a k is equal to a $k^{(t+1,h)}$ with $h \le h_{max}(t+1)$, and even $h \le h_{t+1}^f$ if the latter exists.

Also, similarly to the first case, for all $k \in \mathcal{K}_{t+1}$ such that $i_{t+1} \in R_k$ (recall that this is equivalent to $R_{\bar{k}} \subseteq R_k$), we can write:

$$V_{\Re}^{*}(S_{t+1} \cap R_{k}) \leq \sum_{\substack{k' \in \mathscr{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \text{ by (22) and } \mathscr{K}_{t+1} \subseteq \mathscr{K}_{t}$$

$$= \sum_{\substack{k' \in \mathscr{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_{t+1} \cap R_{\bar{k}}|$$

$$= \sum_{\substack{k' \in \mathscr{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t} \cap R_{k'}| + \zeta_{\bar{k}} \wedge (|S_{t} \cap R_{\bar{k}}| + 1)$$

$$\leq \sum_{\substack{k' \in \mathscr{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t} \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_{t} \cap R_{\bar{k}}| + 1$$

$$= \sum_{\substack{k' \in \mathscr{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t} \cap R_{k'}| + 1$$

$$= V_{\Re}^{*}(S_{t} \cap R_{k}) + 1 \text{ by (17)}. \tag{24}$$

Note that by the joint construction of \mathcal{K}_t^- and \mathcal{P}^t on lines 17 and 18, the fact that $i_{t+1} \notin \bigcup_{k \in \mathcal{K}_t^-} R_k$ implies that \bar{k} is the index of an atom, so actually $h_{\max}(t+1) = \phi(\bar{k})$, $\bar{k} = k^{(t+1,\phi(\bar{k}))}$ and the R_k ,

 $k \in \mathcal{K}_t$, such that $R_{\bar{k}} \subseteq R_k$ are nested and are exactly indexed by the $k^{(t+1,h)}$, $1 \le h \le \phi(\bar{k})$. We now prove that for all of them, $V_{\mathfrak{R}}^*(S_{t+1} \cap R_k) \ge V_{\mathfrak{R}}^*(S_t \cap R_k) + 1$, which will be true in particular for the ones that are in \mathcal{K}_{t+1} , given that $\mathcal{K}_{t+1} \subseteq \mathcal{K}_t$. We do that by constructing some sets A_h with good properties with a descending recursion on h, starting from $\phi(\bar{k})$. We only give the first two steps of the construction, because every other step is exactly the same as the second one, which contains the recursive arguments. We go back to the real definition of $V_{\mathfrak{R}}^*$ to do so, for any $S \subseteq \mathbb{N}_m$:

$$V_{\mathfrak{R}}^{*}(S) = \max_{\substack{A \subseteq \mathbb{N}_{m} \\ \forall k' \in \mathcal{K}, |A \cap R_{k'}| \le \zeta_{k'}}} |A \cap S| = \max_{\substack{A \subseteq S \\ \forall k' \in \mathcal{K}, |A \cap R_{k'}| \le \zeta_{k'}}} |A|. \tag{25}$$

By (25), we have that $V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}) = |A_{\phi(\bar{k})}|$ for a given $A_{\phi(\bar{k})} \subseteq S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}$ and such that $|A_{\phi(\bar{k})} \cap R_{k'}| \leq \zeta_{k'}$ for all $k' \in \mathcal{K}$. Now for the second set, we construct $A_{\phi(\bar{k})-1}$. Note that $V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}) = |B|$ for some $B \subseteq S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}$ and such that $|B \cap R_{k'}| \leq \zeta_{k'}$ for all $k' \in \mathcal{K}$. By reductio ad absurdum, if there are strictly less than $V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}) - V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}) = |B| - |A_{\phi(\bar{k})}|$ elements in $S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}} \setminus S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}$, then $|B| + |S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}| - |S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}| > |A_{\phi(\bar{k})}| = V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k}))}})$. Given that $B \cup (S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}) \subseteq S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}$, this entails $|B \cap S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}| = |B| + |S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}| - |B \cup (S_t \cap R_{k^{(t+1,\phi(\bar{k}))}})| > V^*_{\mathfrak{R}}(S_t \cap R_{k^{(t+1,\phi(\bar{k}))}})$ which contradicts the maximality of $A_{\phi(\bar{k})}$ in (25).

So we construct $A_{\phi(\bar{k})-1}$ by taking the disjoint union of $A_{\phi(\bar{k})}$ and $V_{\mathfrak{R}}^*(S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}) - V_{\mathfrak{R}}^*(S_t \cap R_{k^{(t+1,\phi(\bar{k}))}})$ elements of $S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}} \setminus S_t \cap R_{k^{(t+1,\phi(\bar{k}))}}$. We now establish the properties of $A_{\phi(\bar{k})-1}$. First, $A_{\phi(\bar{k})-1} \subseteq S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}}$, and $|A_{\phi(\bar{k})-1}| = V_{\mathfrak{R}}^*(S_t \cap R_{k^{(t+1,\phi(\bar{k})-1)}})$. For all $k' \in \mathscr{K}$ such that $R_{k^{(t+1,\phi(\bar{k})-1)}} \cap R_{k'} = \emptyset$, we have $|A_{\phi(\bar{k})-1} \cap R_{k'}| = 0 \le \zeta_k'$. Furthermore,

$$\begin{split} |A_{\phi(\bar{k})-1} \cap R_{k^{(t+1,\phi(\bar{k}))}}| &= |A_{\phi(\bar{k})} \cap R_{k^{(t+1,\phi(\bar{k}))}}| \\ &\leq \zeta_{k^{(t+1,\phi(\bar{k}))}} \end{split}$$

by construction of $A_{\phi(\bar{k})}$. Finally, for all k' such that $R_{k^{(t+1,\phi(\bar{k})-1)}}\subseteq R_{k'}$, $|A_{\phi(\bar{k})-1}\cap R_{k'}|=|A_{\phi(\bar{k})-1}|=V_{\mathfrak{R}}^*(S_t\cap R_{k^{(t+1,\phi(\bar{k})-1)}})=|B|$ with the previously defined B, in particular $|B\cap R_{k'}|\leq \zeta_{k'}$, but given that $B\subseteq S_t\cap R_{k^{(t+1,\phi(\bar{k})-1)}}, |B\cap R_{k'}|=|B|$. Wrapping all those equalities, it comes that $|A_{\phi(\bar{k})-1}\cap R_{k'}|\leq \zeta_{k'}$. In the end, $|A_{\phi(\bar{k})-1}\cap R_{k'}|\leq \zeta_{k'}$ for all $k'\in \mathcal{K}$, so $A_{\phi(\bar{k})-1}$ realizes the maximum in (25) for $S_t\cap R_{k^{(t+1,\phi(\bar{k})-1)}}$. By applying exactly the same method, we recursively construct a non-increasing sequence $A_{\phi(\bar{k})}\subseteq \cdots\subseteq A_1$ such that for all $\ell\in\{1,\ldots,\phi(\bar{k})\}$ and $k'\in \mathcal{K}$, $A_\ell\subseteq S_t\cap R_{k^{(t+1,\ell)}}$, $V_{\mathfrak{R}}^*(S_t\cap R_{k^{(t+1,\ell)}})=|A_\ell|$, and $|A_\ell\cap R_{k'}|\leq \zeta_{k'}$. Furthermore for $\ell'>\ell$, $A_\ell\setminus A_{\ell'}\subseteq S_t\cap R_{k^{(t+1,\ell)}}\setminus S_t\cap R_{k^{(t+1,\ell)}}$. Also note that the fact that $i_{t+1}\notin \bigcup_{k\in \mathcal{K}_t^-}R_k$ implies that $\eta_{k^{(t+1,\ell)}}^t<\zeta_{k^{(t+1,\ell)}}$ for all $\ell\in\{1,\ldots,\phi(\bar{k})\}$. So by (16), $|A_\ell|<\zeta_{k^{(t+1,\ell)}}$.

 $\widetilde{A}_{\ell} \subseteq S_{t+1} \cap R_{k^{(t+1,\ell)}}, |\widetilde{A}_{\ell}| = |A_{\ell}| + 1$, and for all $\ell' \in \{1, \dots, \phi(\overline{k})\}, |\widetilde{A}_{\ell} \cap R_{k^{(t+1,\ell')}}| = |A_{\ell} \cap R_{k^{(t+1,\ell')}}| + 1$. Note that if, furthermore, $\ell \ge \ell'$, then $A_{\ell} \subseteq A_{\ell'}$, so

$$\begin{split} |\widetilde{A}_{\ell} \cap R_{k^{(t+1,\ell')}}| &= |A_{\ell} \cap R_{k^{(t+1,\ell')}}| + 1 \\ &\leq |A_{\ell'} \cap R_{k^{(t+1,t')}}| + 1 \\ &= |A_{\ell'}| + 1 \\ &< \zeta_{k^{(t+1,\ell')}} + 1. \end{split}$$

On the contrary, if $\ell < \ell'$, we write that

$$\begin{split} |\widetilde{A}_{\ell} \cap R_{k^{(t+1,\ell')}}| &= |A_{\ell} \cap R_{k^{(t+1,\ell')}}| + 1 \\ &= |(A_{\ell} \setminus A_{\ell'}) \cap R_{k^{(t+1,\ell')}}| + |A_{\ell'} \cap R_{k^{(t+1,\ell')}}| + 1 \\ &< 0 + \zeta_{k^{(t+1,\ell')}} + 1, \end{split}$$

because $A_{\ell} \setminus A_{\ell'} \subseteq R_{k^{(t+1,\ell')}} \setminus R_{k^{(t+1,\ell')}}$ hence $(A_{\ell} \setminus A_{\ell'}) \cap R_{k^{(t+1,\ell')}} = \emptyset$. In both cases, $|\widetilde{A}_{\ell} \cap R_{k^{(t+1,\ell')}}| < \zeta_{k^{(t+1,\ell')}} + 1$ so $|\widetilde{A}_{\ell} \cap R_{k^{(t+1,\ell')}}| \le \zeta_{k^{(t+1,\ell')}}$. Additionally, for all $k' \in \mathcal{K}$ such that $i_{t+1} \notin R_{k'}$, $|\widetilde{A}_{\ell} \cap R_{k'}| = |A_{\ell} \cap R_{k'}| \le \zeta_{k'}$.

In the end, $|\widetilde{A}_{\ell} \cap R_{k'}| \leq \zeta_{k'}$ for all $k' \in \mathcal{K}$, so

$$\begin{split} V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_{k^{(t+1,\ell)}}) &\geq |\widetilde{A}_{\ell}| \text{ by (25)} \\ &= |A_{\ell}| + 1 \\ &= V_{\mathfrak{R}}^{\star}(S_{t} \cap R_{k^{(t+1,\ell)}}) + 1. \end{split}$$

So, as we wanted, $V_{\mathfrak{R}}^*(S_{t+1} \cap R_k) \geq V_{\mathfrak{R}}^*(S_t \cap R_k) + 1$ for all $k \in \mathcal{K}_t$ such that $i_{t+1} \in R_k$ and so for all such k that are in \mathcal{K}_{t+1} . So every inequality in (24) becomes an equality and we have proven that

$$V_{\mathfrak{R}}^{*}(S_{t+1} \cap R_{k}) = V_{\mathfrak{R}}^{*}(S_{t} \cap R_{k}) + 1 = \eta_{k}^{t} + 1 = \eta_{k}^{t+1},$$

that is, (16) is true at t + 1. Looking at the first line of (24), we also proved that

$$V_{\mathfrak{R}}^{*}(S_{t+1} \cap R_{k}) = \sum_{\substack{k' \in \mathcal{P}^{t} \\ R_{k'} \subseteq R_{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|.$$
(26)

The only thing left to prove is that (26) is also true with \mathcal{P}^{t+1} instead of \mathcal{P}^t , that is that (17) also holds at t+1, or, put differently, that

$$\sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| = \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|. \tag{27}$$

If h_{t+1}^f does not exist, meaning that we didn't break the loop, $\mathscr{P}^{t+1} = \mathscr{P}^t$ so there is nothing to prove.

Now assume that h_{t+1}^f exists. So (23) holds. We will split each term in (27) in a sum of two terms. First, note that by (23), for any $k' \in \mathcal{K}$ such that $R_{k'} \cap R_{k^{(t+1,h_{t+1}^f)}} = \emptyset$, we have that $k' \in \mathcal{P}^{t+1}$ if and only if $k' \in \mathcal{P}^t$. And so,

$$\begin{split} \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| &= \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \cap R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{k^{(t+1,h_{t+1}^f)}} \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}| \\ &= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \cap R_{k^{(t+1,h_{t+1}^f)}} = \emptyset \\ R_{k'} \cap R_{k^{(t+1,h_{t+1}^f)}}} = \emptyset} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{k^{(t+1,h_{t+1}^f)}} \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}|. \end{split}$$

Recall that we already proved that there is no $k' \in \mathscr{P}^t$ such that $R_{k^{(t+1,h_{l+1}^f)}} \subsetneq R_{k'}$, so for any $k' \in \mathscr{P}^t$, either $R_{k'} \cap R_{k^{(t+1,h_{l+1}^f)}} = \emptyset$ or $R_{k'} \subseteq R_{k^{(t+1,h_{l+1}^f)}}$. Hence the split

$$\begin{split} \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| &= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \cap R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \ + \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \\ &= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \ + \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|, \\ &= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \ + \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|, \end{split}$$

where the last equality comes from the fact that $R_{k^{(t+1,h_{t+1}^f)}} \subseteq R_k$, because $k \in \mathcal{K}_{t+1}$, $i_{t+1} \in R_k$, and $k^{(t+1,h_{t+1}^f)} \in \mathcal{P}^{t+1}$.

Given the two previously made splits, it remains to prove that

$$\sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_{k}^{(t+1,h_{t+1}^f)}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| = \zeta_{k^{(t+1,h_{t+1}^f)}} \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}|.$$

Interestingly, this does not depend on k anymore. By (26), the left-hand side is equal to $V_{\Re}^*(S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}})$. Because we are breaking the loop at step h_{t+1}^f , $\eta_{t+1}^{t+1} = \zeta_{k^{(t+1,h_{t+1}^f)}}$. Finally, because (16) holds at t+1, $\eta_{k^{(t+1,h_{t+1}^f)}}^{t+1} = V_{\Re}^*(S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}})$. Wrapping all these assertions:

$$\begin{split} \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_{k^{(t+1,h_{t+1}^f)}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| &= V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}) \\ &= V_{\mathfrak{R}}^{\star}(S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}) \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}| \\ &= \eta_{k^{(t+1,h_{t+1}^f)}}^{t+1} \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}| \\ &= \zeta_{k^{(t+1,h_{t+1}^f)}} \wedge |S_{t+1} \cap R_{k^{(t+1,h_{t+1}^f)}}|, \end{split}$$

which achieves the second case and so the proof. \Box

4 Implementation

All algorithms discussed in this manuscript are already implemented in the R (R Core Team, 2024) 438 package sanssouci (Neuvial et al., 2024) which is available on GitHub (see the References for the link) 439 and is dedicated to the computation of confidence bounds for the number of false positives. It also 440 hosts the implementation of the methods described in Blanchard et al. (2020) and Enjalbert-Courrech 441 and Neuvial (2022). Algorithm 1 is implemented as the V. star function, Algorithm 2 is implemented 442 as the pruning function, and Algorithm 4 is implemented as the curve. V. star.forest.fast function 443 (whereas the curve. V. star. forest. naive function just repeatedly calls V. star). Note that the 444 pruning function has a delete.gaps option that speeds up the computation even more by removing 445 unnecessary gaps introduced in the data structure by the pruning operation, those gaps being due to 446 the specific structure that is used to store the information of \mathcal{K} . 447

Speaking of the data structure, we briefly describe it, with an example. We represent $(R_k)_{k\in\mathcal{X}}$ by two 448 lists, C and leaf_list. leaf_list is a list of vectors, where leaf_list[[i]] is the vector listing 449 the hypotheses in the atom P_i . C is a list of lists. For $1 \le h \le H$, C[[h]] lists the regions at depth h, 450 using the index bounds of the atoms they are composed of. That is, the elements of the list C[[h]] 451 are vectors of size two, and if there is k, i and j such that C[[h]][[k]] = c(i, j), it means that 452 $(i,j) \in \mathcal{K}$, or in other words that $P_{i:j}$ is one of the regions, and that $\phi((i,j)) = h$. We emphasize that 453 the 1D structure of the hypotheses has to be respected by the user as the current implementation 454 implicitly uses it: that is, P_1 has to contain the hypotheses labeled 1, 2, ..., p, P_2 has to contain the 455 hypotheses labeled p + 1, ..., and so on. Also, the hypotheses have to be in non-decreasing order: 456 leaf_list[[1]] has to be equal to c(1, 2, 3, ..., p) and not, say, c(2, 1, 3, ..., p).

Example 4.1 (Implementation of Example 2.3). For the reference family given in Example 2.2 and completed in Example 2.3, H = 3. For h = 1, we have C[[1]][[1]] = c(1, 5), C[[1]][[2]] = c(6, 7), C[[1]][[3]] = c(8, 8). For h = 2, we have C[[2]][[1]] = c(1, 1), C[[2]][[2]] = c(2, 1)

```
461 3), C[[2]][[3]] = c(4, 5), C[[2]][[4]] = c(6, 6), C[[2]][[5]] = c(7, 7). For h = 3, we
462 have C[[3]][[1]] = c(2, 2), C[[3]][[2]] = c(3, 3), C[[3]][[3]] = c(4, 4), C[[3]][[4]] =
463 c(5, 5).

464 And then for the atoms, we have leaf_list[[1]] = c(1, 2), leaf_list[[2]] = c(3, 4),
465 leaf_list[[3]] = c(5, 6, 7, 8, 9, 10), leaf_list[[4]] = c(11, 12, 13, 14, 15, 16),
466 leaf_list[[5]] = c(17, 18, 19, 20), leaf_list[[6]] = 21, leaf_list[[7]] = 22 and finally
467 leaf_list[[8]] = c(23, 24, 25).

468 The functions dyadic.from.leaf_list, dyadic.from.window.size, and dyadic.from.height re-
469 turn the appropriate data structure to represent a \mathcal{X} that can be described as a dyadic tree, based on
470 some entry parameters that can be inferred from the names of the functions. The completion of a
471 forest structure, mentioned in Section 2.3, is done by the forest.completion function. Finally, the
472 \zeta_k's are computed as in Durand et al. (2020) by the zetas.tree function with method=zeta.DKWM.
```

5 Numerical experiments

473

In this Section, we present some numerical experiments aiming to demonstrate the impact of the pruning of Algorithm 2 (using the delete.gaps option mentioned in Section 4) and of the fast Algorithm 4, in terms of computation time, compared to the only previously available method to compute a curve of confidence bounds. As mentioned in Section 2.3 and Section 4, this naive method simply consisted in a for loop repeatedly applying Algorithm 1.

To compare the computation time, we use the R package microbenchmark version 1.5.0 (Mersmann, 2024) with R version 4.4.0 (2024-04-24) and sanssouci version 0.13.0, on a MacBook Air M1 (2020) running macOS 15.1.1. The package microbenchmark allows to run code snippets a given number n_repl of times, and to compute summary statistics on the computation time. The script executing the computation can be found in the same repository as this manuscript.

Four scenarios are studied, all based on a common setting which we first describe. A number m of hypotheses is tested. We use a reference family (R_k, ζ_k) such that the R_k 's have a forest structure of maximal depth H=10. The graph of the inclusion relations between the R_k 's is a binary tree, hence there are $2^H-1=1023$ R_k 's and in particular $2^{H-1}=512$ atoms. P-values are generated in a gaussian one-sided fashion where $H_{0,i}=\{\mathcal{N}(\mu,\mathrm{Id}): \mu_i=0\}$, $H_{1,i}=\{\mathcal{N}(\mu,\mathrm{Id}): \mu_i=4\}$, and $p_i(X)=1-\Phi(X_i)$. \mathcal{H}_1 is comprised of the leafs 1, 5, 9 and 10, that is $\mathcal{H}_1=P_1\cup P_5\cup P_9\cup P_{10}$. For each scenario, the curve $\left(V_{\mathfrak{R}}^*(\{1,\ldots,t\})\right)_{t\in\mathbb{N}_m^*}$ is computed. For the experiments including pruning, the pruning is done once before the n_rep1 replications, to mimick the practice where pruning only needs to be done once and for all, while the user may be interested in computing multiple bounds and curves after that.

In scenarios 1 and 2, m = 1024 (so the atoms are of size 2), in scenarios 3 and 4, m = 10240 (so the atoms are of size 10). In scenarios 1 and 3, the ζ_k 's are estimated trivially by $\zeta_k = |R_k|$, and in scenarios 2 and 4, they are computed as in Durand et al. (2020) with the DKWM inequality (Dvoretzky et al., 1956; Massart, 1990). Because of the size of m and the poor performances of the naive approach, we set n_repl=100 in scenarios 1 and 2 and n_repl=10 only in scenarios 3 and 4. The differences between the scenarios are summarized in Table 1.

Table 1: Differences between the scenarios

parameter	Scenario 1	Scenario 2	Scenario 3	Scenario 4
m zeta computation	1024	1024	10240	10240
	trivial	DKWM	trivial	DKWM

parameter	Scenario 1	Scenario 2	Scenario 3	Scenario 4
n_repl	100	100	10	10

For the trivial ζ_k computation of scenarios 1 and 3, the pruning obviously deletes all non-atom regions so $|\mathcal{K}^{\mathfrak{pr}}| = 512$. Whereas, for the particular instance $\omega \in \Omega$ in the experiments, $|\mathcal{K}^{\mathfrak{pr}}| = 541$ for scenario 2, and $|\mathcal{K}^{\mathfrak{pr}}| = 573$ for scenario 4. Those results alone illustrate the benefits of pruning with respect to the reduction of the cardinality of the reference family: the regions above atoms with no signal (or no detectable signal in the trivial scenarios) are pruned. The fact that the regions above atoms with detectable signal are not pruned means that they are relevant for the confidences bounds (which had already been demonstrated in the simulation study of Durand et al. (2020)).

501

502

503

The summary statistics of the computation time in each scenario are presented in Table 2, Table 3, Table 4, and Table 5, and they are also presented as boxplots in Figure 11. The time unit is the second in logarithmic scale in the boxplots).

Table 2: Scenario 1

expr	min	lq	mean	median	υ	ıq max	neval
naive.not.pruned	3.6924007	3.7943906	3.8521256	3.8386487	3.878041	2 4.5247099	100
naive.pruned	3.2822354	3.4126177	3.4758338	3.4614076	3.506154	3.8822089	100
fast.not.pruned	0.1332744	0.1367000	0.1383806	0.1385039	0.139270	0.1768691	100
fast.pruned	0.0921422	0.0945472	0.0974025	0.0954231	0.097868	0.1908498	100

Table 3: Scenario 2

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	3.7280744	3.8025695	3.8514710	3.8451367	3.8831009	4.1891831	100
naive.pruned	3.3556131	3.4533210	3.4926114	3.4906796	3.5182172	3.8501820	100
fast.not.pruned	0.1214844	0.1246071	0.1265674	0.1260760	0.1279640	0.1407320	100
fast.pruned	0.0815349	0.0827995	0.0841622	0.0835618	0.0851062	0.0896013	100

Table 4: Scenario 3

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	d 332.1856576	335.5148922	337.9856658	338.2432916	340.3329972	344.6255264	10
naive.pruned	328.3186707	329.3081834	332.1861199	331.4335773	333.3563651	338.7111614	10
fast.not.pruned	1.4881838	1.4966417	1.5066370	1.5078498	1.5151194	1.5217546	10
fast.pruned	0.9354581	0.9418174	0.9498806	0.9512573	0.9550453	0.9675895	10

Table 5: Scenario 4

expr	min	lq	mean	median	uq	max	neval
naive.not.prune	d 331.0124665	335.6357519	349.7740812	337.6459728	342.1652204	401.4881647	10
naive.pruned	331.2567637	332.2215437	363.5822362	333.0651271	335.8347696	493.5124771	10
fast.not.pruned	1.3575818	1.3588461	1.3733567	1.3641336	1.3762178	1.4460291	10

Table 5: Scenario 4

expr	min	lq	mean	median	uq	max	neval
fast.pruned	0.9287399	0.9441687	0.9551275	0.9520622	0.9624959	0.9972532	10

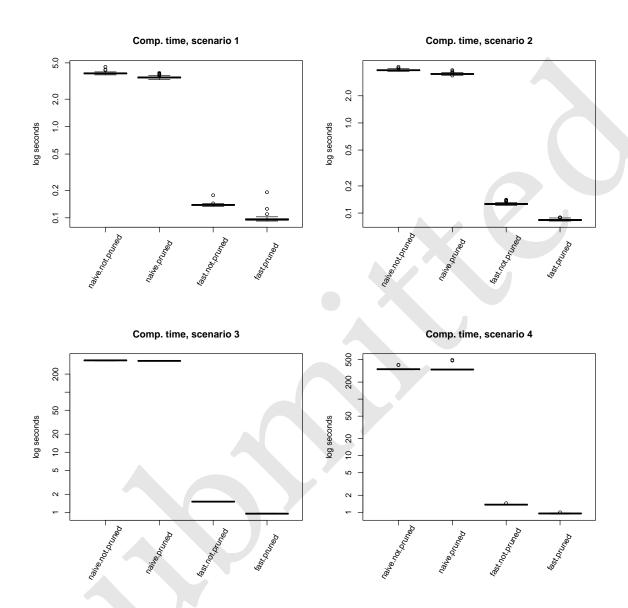


Figure 11: Computation times in scenario, in log seconds

On each scenario, using the fast algorithm is much faster than the naive approach, while pruning always gives a slight improvement over not pruning.

511

512

513

515

Comparing scenarios 1 and 2 first, we see that, as expected, there is no significant change in computation time for naive.not.pruned, while naive.pruned is faster in scenario 1, given that we prune more. But, on the other hand, fast.not.pruned and fast.pruned are both faster in scenario 2, even if we prune less. This is because, for the regions with signal, said signal is detected and so those regions are quickly saturated, in the sense that we quickly have $\eta_k^t = \zeta_k$ and k added to \mathcal{K}_k^- , which saves a lot of time.

The comparison between scenarios 3 and 4 is similar, except that this time we prune even less in scenario 4 and so the effect of the saturation is not enough to compensate. Although, with only n_repl=10, the statistics seem less accurate, this can be confirmed with additional experiments (n_repl can also be set to 100 without problem is we don't include naive methods).

Finally, comparing scenarios 3 & 4 with scenarios 1 & 2, we see that multiplying the number of hypotheses by 10 effectively multiplies the computation time by \sim 10 when using Algorithm 4 and by \sim 100 when using Algorithm 1 naively, which illustrates the stated complexities of $O(m|\mathcal{K}|)$ and $O(m^2|\mathcal{K}|)$, respectively.

526 6 Conclusion

In conclusion, we effectively introduced a new algorithm to compute a curve of confidence upper bounds, much faster the previous alternative, with one power of *m* less in the complexity.

To develop new confidence upper bounds methodology and test them on simulations, it was previously not conceivable to replicate experiments a sufficient number of times while computing whole curves. For instance, in the simulation study of Durand et al. (2020), the number of replications chosen was 10 and the whole curve was not computed, only ten values along the curve were computed, for an m set to 12800, that is 0.078% of the curve had been computed. Now, simulation studies with an adequate number of replications and 100% of the curve become feasible.

A lot of work remains to be done on the sanssouci package. For example, to make the data format of a forest structure $(R_k)_{k\in\mathcal{K}}$ less convoluted and more user-friendly is an interesting project. Another one would be to implement inside the package the methods of the paper Blain et al. (2022), which are currently only available in the Python language (Van Rossum and Drake, 2009), and the methods of the paper Meah et al. (2024).

Other current works include the development of new reference families with theoretical JER control that could better account for realistic models, such as models with dependence between the *p*-values, see for example Perrot-Dockès et al. (2023), or models with discreteness.

7 Acknowledgements

This work has been supported by ANR-20-IDEES-0002 (PIA), ANR-19-CHIA-0021 (BISCOTTE), ANR-23-CE40-0018 (BACKUP) and ANR-21-CE23-0035 (ASCAI). Thanks to Romain Périer for being the first to extensively use the new implemented algorithms. Thanks to Pierre Neuvial for his valuable feedback.

48 References

543

Yoav Benjamini and Yosef Hochberg. Controlling the false discovery rate: a practical and powerful approach to multiple testing. *J. Roy. Statist. Soc. Ser. B*, 57(1):289–300, 1995. ISSN 0035-9246. URL https://www.jstor.org/stable/2346101.

Alexandre Blain, Bertrand Thirion, and Pierre Neuvial. Notip: Non-parametric True Discovery
Proportion control for brain imaging. *Neuroimage*, 260, October 2022. URL https://doi.org/10.1016/j.neuroimage.2022.119492.

Gilles Blanchard, Pierre Neuvial, and Etienne Roquain. Post hoc confidence bounds on false positives using reference families. *Ann. Statist.*, 48(3):1281–1303, 2020. ISSN 0090-5364. doi: 10.1214/19-AOS1847. URL https://doi.org/10.1214/19-AOS1847.

- Małgorzata Bogdan, Ewout van den Berg, Chiara Sabatti, Weijie Su, and Emmanuel J. Candès. SLOPE—
 adaptive variable selection via convex optimization. *Ann. Appl. Stat.*, 9(3):1103–1140, 2015. ISSN
 1932-6157,1941-7330. doi: 10.1214/15-AOAS842. URL https://doi.org/10.1214/15-AOAS842.
- Guillermo Durand, Gilles Blanchard, Pierre Neuvial, and Etienne Roquain. Post hoc false positive control for structured hypotheses. *Scand. J. Stat.*, 47(4):1114–1148, 2020. ISSN 0303-6898. doi: 10.1111/sjos.12453. URL https://doi.org/10.1111/sjos.12453.
- A. Dvoretzky, J. Kiefer, and J. Wolfowitz. Asymptotic minimax character of the sample distribution
 function and of the classical multinomial estimator. *Ann. Math. Statist.*, 27:642–669, 1956. ISSN
 0003-4851. doi: 10.1214/aoms/1177728174. URL https://doi.org/10.1214/aoms/1177728174.
- Nicolas Enjalbert-Courrech and Pierre Neuvial. Powerful and interpretable control of false discoveries in two-group differential expression studies. *Bioinformatics*, 38(23):5214–5221, 10 2022. ISSN 1367-4803. doi: 10.1093/bioinformatics/btac693. URL https://doi.org/10.1093/bioinformatics/btac693.
- Christopher R. Genovese and Larry Wasserman. Exceedance control of the false discovery proportion.
 J. Amer. Statist. Assoc., 101(476):1408–1417, 2006. ISSN 0162-1459. doi: 10.1198/016214506000000339.
 URL https://doi.org/10.1198/016214506000000339.
- Jelle J. Goeman and Aldo Solari. Multiple testing for exploratory research. *Statist. Sci.*, 26(4):584–597, 2011. ISSN 0883-4237. doi: 10.1214/11-STS356. URL https://doi.org/10.1214/11-STS356.
- Ruth Marcus, Eric Peritz, and K. R. Gabriel. On closed testing procedures with special reference to ordered analysis of variance. *Biometrika*, 63(3):655–660, 1976. ISSN 0006-3444. doi: 10.1093/biomet/63.3.655. URL https://doi.org/10.1093/biomet/63.3.655.
- P. Massart. The tight constant in the Dvoretzky-Kiefer-Wolfowitz inequality. *Ann. Probab.*, 18(3):
 1269–1283, 1990. ISSN 0091-1798,2168-894X. URL http://links.jstor.org/sici?sici=0091-1798(199007)
 18:3<1269:TTCITD>2.0.CO;2-Q&origin=MSN.
- Iqraa Meah, Gilles Blanchard, and Etienne Roquain. False discovery proportion envelopes with m-consistency. *Journal of Machine Learning Research*, 25(270):1–52, 2024. URL http://jmlr.org/ papers/v25/23-1025.html.
- Rosa J. Meijer, Thijmen J. P. Krebs, and Jelle J. Goeman. A region-based multiple testing method for
 hypotheses ordered in space or time. *Stat. Appl. Genet. Mol. Biol.*, 14(1):1–19, 2015. ISSN 2194-6302.
 doi: 10.1515/sagmb-2013-0075. URL https://doi.org/10.1515/sagmb-2013-0075.
- Nicolai Meinshausen. False discovery control for multiple tests of association under general dependence. *Scand. J. Statist.*, 33(2):227–237, 2006. ISSN 0303-6898. doi: 10.1111/j.1467-9469.2005.00488.x.
 URL https://doi.org/10.1111/j.1467-9469.2005.00488.x.
- Olaf Mersmann. *microbenchmark: Accurate Timing Functions*, 2024. URL https://CRAN.R-project.org/package=microbenchmark. R package version 1.5.0.
- Pierre Neuvial, Gilles Blanchard, Guillermo Durand, Nicolas Enjalbert-Courrech, and Etienne Roquain.
 sanssouci: Post Hoc Multiple Testing Inference, 2024. URL https://sanssouci-org.github.io/sanssouci.
 R package version 0.13.0.
- Marie Perrot-Dockès, Gilles Blanchard, Pierre Neuvial, and Etienne Roquain. Selective inference for false discovery proportion in a hidden markov model. *TEST*, pages 1–27, 2023.
- R Core Team. *R: A Language and Environment for Statistical Computing*. R Foundation for Statistical Computing, Vienna, Austria, 2024. URL https://www.R-project.org/.

- Guido Van Rossum and Fred L. Drake. Python 3 Reference Manual. CreateSpace, Scotts Valley, CA,
 2009. ISBN 1441412697.
- Anna Vesely, Livio Finos, and Jelle J. Goeman. Permutation-based true discovery guarantee by sum tests. J. R. Stat. Soc. Ser. B. Stat. Methodol., 85(3):664–683, 2023. ISSN 1369-7412,1467-9868. doi: 10.1093/jrsssb/qkad019. URL https://doi.org/10.1093/jrsssb/qkad019.

Session information

```
R version 4.4.1 (2024-06-14)
   Platform: x86_64-pc-linux-gnu
   Running under: Ubuntu 24.04.1 LTS
607
608
   Matrix products: default
            /usr/lib/x86_64-linux-gnu/blas/libblas.so.3.12.0
   BLAS:
   LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.12.0
612
   locale:
613
                                                          LC TIME=C.UTF-8
     [1] LC CTYPE=C.UTF-8
                                 LC NUMERIC=C
614
                                                          LC_MESSAGES=C.UTF-8
     [4] LC_COLLATE=C.UTF-8
                                 LC_MONETARY=C.UTF-8
615
     [7] LC_PAPER=C.UTF-8
                                 LC_NAME=C
                                                          LC_ADDRESS=C
   [10] LC_TELEPHONE=C
                                 LC_MEASUREMENT=C.UTF-8 LC_IDENTIFICATION=C
617
618
   time zone: Etc/UTC
619
   tzcode source: system (glibc)
620
621
   attached base packages:
   [1] stats
                  graphics
                                                   utils
                            grDevices datasets
                                                             methods
                                                                        base
624
   other attached packages:
625
   [1] microbenchmark 1.5.0
626
   loaded via a namespace (and not attached):
     [1] compiler_4.4.1
                            fastmap_1.1.1
                                                cli_3.6.2
                                                                   htmltools_0.5.8.1
629
     [5] tools_4.4.1
                            yaml_2.3.8
                                               rmarkdown_2.26
                                                                   knitr_1.46
630
     [9] jsonlite_1.8.8
                            xfun_0.43
                                               digest_0.6.35
                                                                   rlang_1.1.3
631
   [13] renv_1.0.7
                            evaluate_0.23
632
```