


Fast confidence bounds for the false discovery proportion over a path of hypotheses

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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 $V_{\mathfrak{R}}^*$ construction is based on a reference family \mathfrak{R} with a forest structure à la [Durand et al. \(2020\)](#). By an entire curve, we mean the values $V_{\mathfrak{R}}^*(S_1), \dots, V_{\mathfrak{R}}^*(S_m)$ computed on a path of increasing selection sets $S_1 \subsetneq \dots \subsetneq S_m, |S_t| = t$. The new algorithm leverages the fact that going from S_t to S_{t+1} is done by adding only one hypothesis. Compared to a more naive approach, the new algorithm has a complexity in $O(|\mathcal{K}|m)$ instead of $O(|\mathcal{K}|m^2)$, where $|\mathcal{K}|$ is the cardinality of the family.

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

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1 Introduction

Multiple testing theory is often used for exploratory analysis, like in 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 confidence upper bounds for the number of false discoveries, or, equivalently, for the False Discovery Proportion (FDP). This approach is notably advocated in the context of exploratory research by ([Goeman and Solari, 2011](#), Section 1).

Mathematically speaking, assume that we observe some data X that is formally a random variable defined on some probability space equipped of the probability measure \mathbb{P} , and that the distribution of X , denoted by P , belongs to a model \mathfrak{F} . We want to test m null hypotheses $H_{0,1}, \dots, H_{0,m} \subset \mathfrak{F}$. A confidence upper bound (also frequently named post hoc bound, post selection bound or confidence envelope) is then a function $\hat{V} : \mathcal{P}(\mathbb{N}_m^*) \rightarrow \mathbb{N}_m$, where $\mathbb{N}_m = \{0, \dots, m\}$, $\mathbb{N}_m^* = \{1, \dots, m\}$, such that

$$\forall \alpha \in (0, 1), \mathbb{P} \left(\forall S \subseteq \mathbb{N}_m^*, |S \cap \mathcal{H}_0| \leq \hat{V}(S) \right) \geq 1 - \alpha. \quad (1)$$

Here, α is a target error rate and $\mathcal{H}_0 = \{i : P \in H_{0,i}\}$ is the set of indices of the null hypotheses that are true. Note that the construction of \hat{V} depends on α and on the 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 for the number of null hypotheses in S for any selection set $S \subseteq \mathbb{N}_m^*$, that is, the number $|S \cap \mathcal{H}_0|$ of false discoveries in S . This 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:

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

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 construction is the following: one can compute the largest S such that $\frac{\hat{V}(S)}{|S| \vee 1}$

is less than or equal to a nominal level q , and (2) ensures that, with high probability, the FDP of S is upper-bounded by q . The control of the FDP with high probability is sometimes called False Discovery Exceedance (FDX) control.

Post hoc bounds have notably been applied to genetic data. For example, in [Goeman and Solari \(2011\)](#) and [Enjalbert-Courrech and Neuvial \(2022\)](#), the authors apply post hoc bounds to an Urothelial Bladder Carcinoma RNA sequencing dataset to detect genes differentially expressed between stage II and stage III of the disease. Furthermore, the R ([R Core Team, 2024](#)) package `IIDEA` ([Enjalbert Courrech and Neuvial \(2025\)](#), see also [Enjalbert Courrech \(2024\)](#), Chapter 3) implements a user-friendly shiny application ([Chang et al., 2025](#)) that computes post hoc bounds for differential expression analyses, where the user can upload their own microarray or bulk RNAseq data file (the application also comes with the aforementioned dataset).

Another field where post hoc bounds have been successfully applied is functional Magnetic Resonance Imaging (fMRI) studies, where each voxel of an image is tested to detect activation of the corresponding brain region during a given task. Using the aforementioned FDP-controlling construction, in [Blain et al. \(2022\)](#) and [Blain et al. \(2023\)](#), the authors construct rejection sets with a high number of true positives.

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 $\{i \in \mathbb{N}_m : p_i \leq s\}$ where p_i is the p -value associated to the null hypothesis $H_{0,i}$ and $s \in [0, 1]$ is a threshold. In [Goeman and Solari \(2011\)](#) the authors re-wrote the generic construction of [Genovese and Wasserman \(2006\)](#) in terms of closed testing (a framework first introduced for the FWER control by [Marcus et al. \(1976\)](#)), proposed several practical constructions and sparked a new interest in multiple testing procedures based on confidence bounds. 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 to construct post hoc bounds, and show the links between this meta-technique and the closed testing one, along with new bounds. Reference families are families of couples $(R_k, \zeta_k)_{k \in \mathcal{K}}$ where R_k is a subset of hypotheses (called a region), and ζ_k an over-estimator of the number of null hypotheses inside R_k , that is, of $|R_k \cap \mathcal{H}_0|$. From a statistical guarantee on the family, called the Joint Error Rate (JER) control, one is able to build a post hoc bound, denoted $V_{\mathfrak{R}}^*$ in the remainder, by interpolation (see Section 2.2 for all the details).

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 $V_{\mathfrak{R}}^*(S)$ for a given, single selection set S . The constraint, named “forest structure”, is that two regions of hypotheses R_k and $R_{k'}$ are either disjoint, or nested: $R_k \cap R_{k'} \in \{R_k, R_{k'}, \emptyset\}$. This structure arises when the object of study naturally presents different levels of hierarchy. For example, in genomic studies, where each hypothesis tests the association of a Single Nucleotide Polymorphism (SNP) with a given character, we can exploit the natural grouping of SNPs into genes or intergenic regions, and then the grouping of genes into genomic pathways, or into chromosomes. In proteomic studies, where the smallest unit is usually the peptide, we can exploit the natural grouping of peptides into proteins, and the grouping of proteins into proteomic pathways. In brain imagery, known brain anatomy can be used to build the regions.

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: $S_t = \{\sigma(1), \dots, \sigma(t)\}$, where σ is a (random) permutation ordering the p -values: $p_{\sigma(1)} \leq \dots \leq p_{\sigma(m)}$. Whereas the algorithm provided in the aforementioned work ([Durand et al., 2020](#), Algorithm 1), which is reproduced here (see ?) 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 $S_1 \subsetneq \dots \subsetneq S_m$ and $|S_t| = t$, there is a way to

efficiently compute $(V_{\mathfrak{N}}^*(S_t))_{t \in \mathbb{N}_m}$ by leveraging the nested structure.

This is the main contribution of the present paper: a new and fast algorithm (?) computing the curve $(V_{\mathfrak{N}}^*(S_t))_{t \in \mathbb{N}_m}$ for a nested path of selection sets, that is presented in Section 3.2. An additional pruning 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. Notably, a detailed example illustrating how the new algorithms work is provided in Section 3.3. In Section 2, 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), with an example code. A few numerical experiments are presented in Section 5 to demonstrate the computation time gain. We reproduce here, in Figure 1, the striking results of one of those experiments, where the combination of the two new algorithms improves the computation time by a factor 33000. Finally, after some concluding remarks in Section 6, the proofs of all results, including the proof that ? indeed computes correctly the curve, are presented in Section 7.

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 \mathfrak{F} on a measurable space $(\mathcal{X}, \mathfrak{X})$, and data that is represented by a random variable $X : (\Omega, \mathcal{A}) \rightarrow (\mathcal{X}, \mathfrak{X})$ with $X \sim P \in \mathfrak{F}$, that is, the law of X is comprised in the model \mathfrak{F} .

Then we consider $m \geq 1$ null hypotheses $H_{0,1}, \dots, H_{0,m}$ which formally are submodels, that is subsets of \mathfrak{F} . The associated alternative hypotheses $H_{1,1}, \dots, H_{1,m}$ are submodels such that $H_{0,i} \cap H_{1,i} = \emptyset$ for all $i \in \mathbb{N}_m^*$. We denote by $\mathcal{H}_0 = \mathcal{H}_0(P)$ (the dependence in P will be dropped when there is no ambiguity) the set of all null hypotheses that are true, that is $\mathcal{H}_0(P) = \{i \in \mathbb{N}_m^* : P \in H_{0,i}\}$. In other words, $H_{0,i}$ is true if and only if $i \in \mathcal{H}_0$. 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 $i \in \mathcal{H}_0$, then the law of p_i is super-uniform, which is sometimes denoted $\mathcal{L}(p_i) \succeq \mathcal{U}([0, 1])$. 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 $U \sim \mathcal{U}([0, 1])$:

$$\forall x \in \mathbb{R}, \mathbb{P}(p_i \leq x) \leq \mathbb{P}(U \leq x) = 0 \vee (x \wedge 1). \quad (3)$$

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 discoveries, or false positives, 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 re-appear slightly simplified in Section 5.

Example 2.1 (Gaussian one-sided). In this case we assume that $X = (X_1, \dots, X_m)$ 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 $p_i(X) = 1 - \Phi(X_i)$, where Φ denotes the cdf of $\mathcal{N}(0, 1)$.

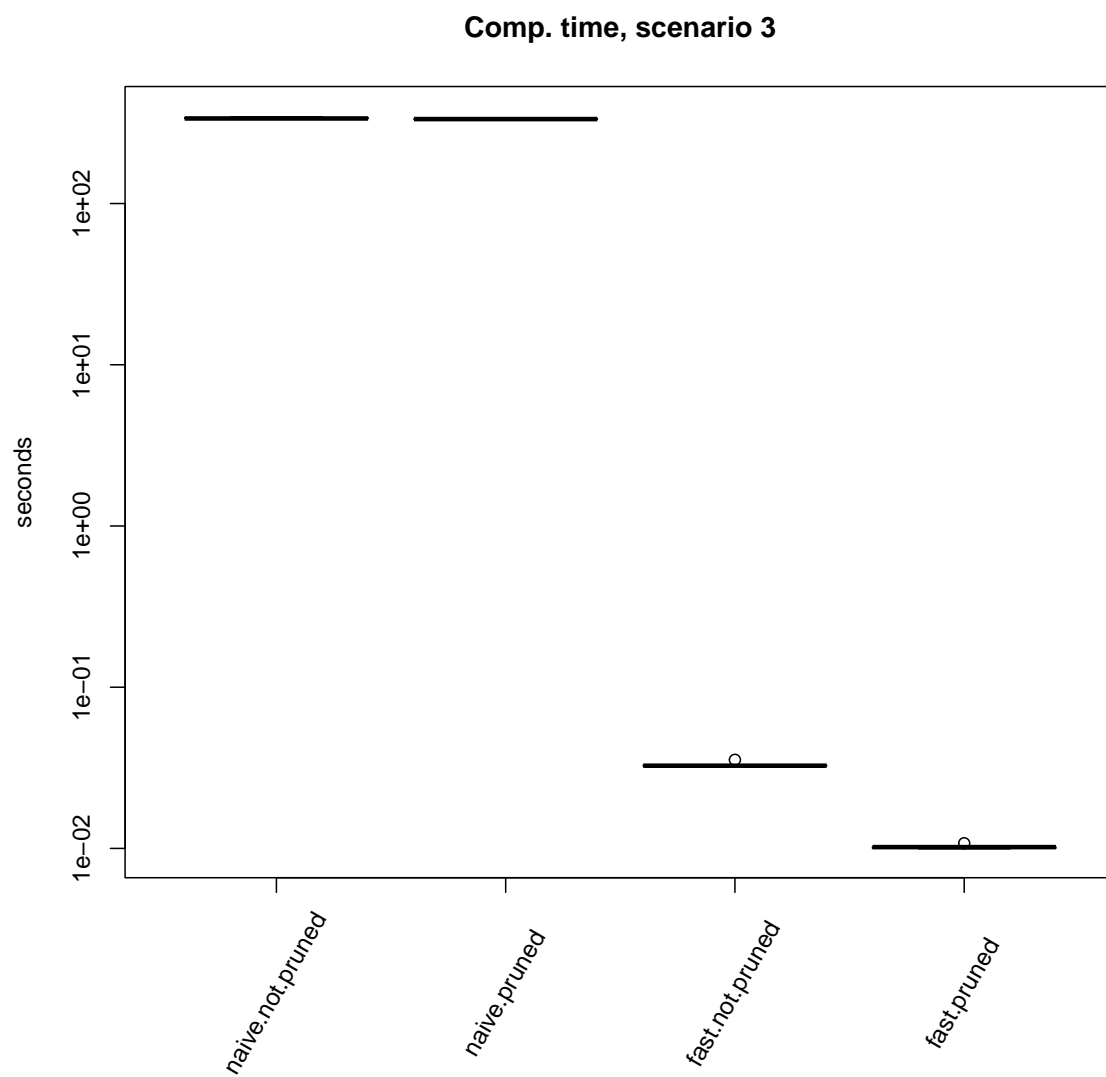


Figure 1: Computation time in scenario 3 for the new fast algorithm versus the previous, naive approach, in seconds (using a logarithmic scale)

2.2 Post hoc bounds with reference families

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

$$\forall P \in \mathcal{P}, \forall X \sim P, \forall \alpha \in (0, 1), \mathbb{P} \left(\forall S \subseteq \mathbb{N}_m^*, V(S) \leq \widehat{V}(X, \alpha)(S) \right) \geq 1 - \alpha. \quad (4)$$

In the remainder, the dependence in (X, α) will be dropped when there is no ambiguity and $\widehat{V}(X, \alpha)(\cdot)$ will simply be written \widehat{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 finite family $\mathfrak{R} = \mathfrak{R}(X, \alpha) = (R_k, \zeta_k)_{k \in \mathcal{K}}$ with $R_k \subseteq \mathbb{N}_m^*$, $\zeta_k \in \{0, \dots, |R_k|\}$ where everything (that is, \mathcal{K} and all the R_k and ζ_k) depends on (X, α) but the dependency is not explicitly written. The R_k are assumed all distinct almost surely (see Remark 2.2).

The intuition of the concept of reference families is the following. The R_k 's are subsets, or regions, of hypotheses, and the associated ζ_k 's are over-estimators of $V(R_k)$. That is, building a reference family amounts to building a collection of regions, that can be much smaller than all possible subsets of hypotheses (note that $|\mathcal{P}(\mathbb{N}_m^*)| = 2^m$ is likely to be very large), for which we have a confidence upper bound. This bound, holding only on the R_k 's in the first place, will then be extended to a simultaneous confidence bound over all subsets S (in the sense of Equation (4)) by an interpolation scheme explained below.

The statistical guarantee over the ζ_k 's, as over-estimators of $V(R_k)$, is written in terms of the following error criterion for a reference family, named Joint Error Rate (JER):

$$\text{JER}(\mathfrak{R}) = \mathbb{P}(\exists k \in \mathcal{K}, |R_k \cap \mathcal{H}_0| > \zeta_k) = \mathbb{P}(\exists k \in \mathcal{K}, V(R_k) > \zeta_k). \quad (5)$$

We say that the reference family \mathfrak{R} controls the JER if the following is true:

$$\forall P \in \mathcal{P}, \forall X \sim P, \forall \alpha \in (0, 1), 1 - \text{JER}(\mathfrak{R}(X, \alpha)) = \mathbb{P}(\forall k \in \mathcal{K}, V(R_k) \leq \zeta_k) \geq 1 - \alpha. \quad (6)$$

Note that Equation (6) is, as foretold, 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 the JER-controlling reference family with the following two steps. First let

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

The JER control says that, with high probability, $\mathcal{H}_0 \in \mathcal{A}(\mathfrak{R})$. We then leverage this information by interpolation, with the following construction:

$$V_{\mathfrak{R}}^*(S) = \max_{A \in \mathcal{A}(\mathfrak{R})} |S \cap A|. \quad (8)$$

By Proposition 2.1 of [Blanchard et al. \(2020\)](#), the JER control of the family in Equation (6) implies that $V_{\mathfrak{R}}^*$ is indeed a confidence bound as required by Equation (4). The same Proposition also establishes that $V_{\mathfrak{R}}^*$ optimally uses the information provided by the JER control of the reference family.

Note that, because of the $\max_{A \in \mathcal{A}(\mathfrak{R})}$, the computation of $V_{\mathfrak{R}}^*(S)$ 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 the remainder of this paper.

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. \(2020\)](#), [Blain et al. \(2022\)](#) or [Meah et al. \(2024\)](#), for example.

Remark 2.2. Some reference family constructions can yield $R_k = R_{k'}$ for $k \neq k'$, for example in the setting of [Meah et al. \(2024\)](#) with discrete p -values. But in that scenario we can always prune the duplicate and keep only one index so that $k \mapsto R_k$ is injective. We implicitly consider that this operation is always done in practice and in the remainder of this article. Of course, if $R_k = R_{k'}$ with $k \neq k'$, we keep the index with the lower value of ζ , that is we keep $\tilde{k} \in \arg \min_{\ell \in \{k, k'\}} \zeta_\ell$ (not doing so would change the bound defined by (8) and decrease its power in terms of type-II error). Similarly, some constructions can yield empty regions, which can always be pruned without changing the bound. This will also be assumed to be the case in the following. Finally, note that the constraint that $\zeta_k \leq |R_k|$ always hold no matter how ζ_k was computed, up to replacing ζ_k by $\zeta_k \wedge |R_k|$: it is clear on Equation (7) that this doesn't change the bound.

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:

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

Representing the R_k 's with a directed graph, where there is an oriented edge $R_k \leftarrow R_{k'}$ if and only if $R_k \subseteq R_{k'}$ and there is no $R_{k''}$ such that $R_k \subsetneq R_{k''} \subsetneq R_{k'}$ gives a forest, hence the name. See [Example 2.2](#) and its representation in [Figure 2](#).

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

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

This definition matches the intuition of depth because we assumed the R_k are distinct, see [Remark 2.2](#).

In all the remainder, H refers to the maximum depth in the structure: $H = \max_{k \in \mathcal{K}} \phi(k)$.

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

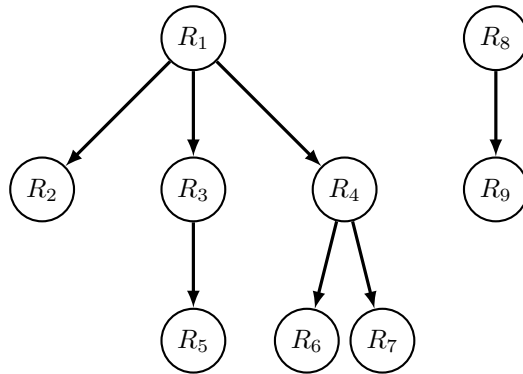


Figure 2: The regions of [Example 2.2](#).

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

$\{\emptyset, \{i, \dots, j\}, \{i', \dots, j'\}\}$. With this identification, each $R_k = R_{(i,j)}$ can be written as $P_{i:j} = \bigcup_{i \leq n \leq j} P_n$ where $(P_n)_{1 \leq n \leq N}$ 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 leaves. 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 3. 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).

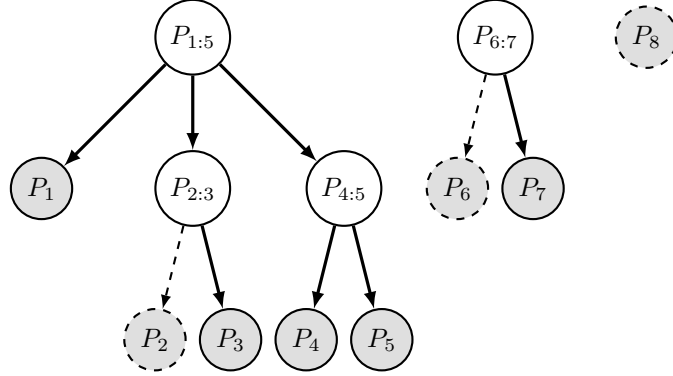


Figure 3: The regions of Example 2.2 but with the labels of Example 2.3.

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 $V_{\mathfrak{A}}^*$. 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:

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

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

$$V_{\mathfrak{A}}^*(S) = \min_{Q \in \mathfrak{P}} \sum_{k' \in Q} \zeta_{k'} \wedge |S \cap R_{k'}|, \quad (12)$$

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 elements $Q \subseteq \mathcal{K}$ 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 $V_{\mathfrak{A}}^*(S)$ for a single $S \subseteq \mathbb{N}_m^*$, which we reproduce here in ?. The family is assumed complete, otherwise the first step would be to complete it. In the original paper, \mathcal{K}^h used to designate the elements of \mathcal{K} at depth h plus the atoms at depth $\leq h$. Actually, including those atoms is not needed for this algorithm to perform exactly the same, and produces redundant computations. If we don't include them, the only difference 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 \mathcal{K}^h as only the elements of \mathcal{K} at depth h (the previous intricate definition may still be necessary for the proof of Theorem 1 of Durand et al. (2020)): $\mathcal{K}^h = \{(i, j) \in \mathcal{K} : \phi(i, j) = h\}, h \geq 1$.

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 \mathcal{K}^1 are the roots of the different trees making up the forest.

Algorithm 1 Computation of a given $V_{\mathfrak{R}}^*(S)$ with a complete family

```

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

```

A step by step description of ? is provided at the end of Section 3 and in the Figure 9 of [Durand et al. \(2020\)](#).

The computation time of the algorithm is in $O(|\mathcal{K}||S|)$, 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 $O(|\mathcal{K}|m^2)$, which is not desirable and makes computations difficult in practice, hence the need for a new, faster algorithm.

Tip 1

In the practical implementation of this algorithm (and of the following ?), Vec and $newVec$ are always of size N (the number of leaves) instead of the cardinality of \mathcal{K}^h . And the sum $\sum_{k' \in Succ_k} Vec_{k'}$ is really easy to compute: if $R_k = R_{(i_0, i_p-1)} = \bigcup_{j=1}^p R_{(i_{j-1}, i_j-1)} = \bigcup_{i_0 \leq n \leq i_p-1} P_n \in \mathcal{K}^h$ 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 $1 \leq j \leq p$, 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.

Furthermore, computing $|S \cap R_k|$ for each k is not necessary, it is sufficient to compute $|S \cap P_n|$ for each leaf P_n , which can be done in $O(N|S|)$.

By the two previous points, we can actually refine the complexity result of ?: it is in $O(N|S| + |\mathcal{K}|) \leq O(|\mathcal{K}||S|)$ because $N \leq |\mathcal{K}|$ for a complete forest, and if the forest is not complete, the Algorithm 2 of [Durand et al. \(2020\)](#) constructs a partition $(P_n)_{1 \leq n \leq N}$ such that $N \leq |\mathcal{K}|$, and so the cardinality of the completed forest is $\leq 2|\mathcal{K}|$.

Speaking of complexity, we have the following result regarding m , $|\mathcal{K}|$ and N :

Proposition 2.1. *For any reference family $(R_k, \zeta_k)_{k \in \mathcal{K}}$ with a forest structure, we have $N \leq m$,*

$H \leq N$, $|\mathcal{K}| \leq 2N - 1$ and these three bounds can be achieved simultaneously. In particular, $|\mathcal{K}| \leq 2m - 1$.

The proof of Proposition 2.1 is given in Section 7.3.

3 New algorithms

3.1 Pruning the forest

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

Definition 3.1 (Pruning). We define by \mathcal{K}^{pr} (\mathcal{K} pruned) the set of elements of \mathcal{K} from which we removed all (i, i') such that 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 $1 \leq j \leq p$ and finally $\zeta_{(i,i')} = \zeta_{(i_0,i_p-1)} \geq \sum_{j=1}^p \zeta_{(i_{j-1}, i_j-1)}$.

An important note is that for a removed $(i, i') \in \mathcal{K} \setminus \mathcal{K}^{\text{pr}}$, we can always choose the indices i_1, \dots, i_{p-1} such that actually $(i_j, i_{j+1} - 1) \in \mathcal{K}^{\text{pr}}$ and not only \mathcal{K} , because if $(i_j, i_{j+1} - 1) \in \mathcal{K} \setminus \mathcal{K}^{\text{pr}}$ 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 have at hand a forest structure, that is, the reference family defined by $\mathfrak{R}^{\text{pr}} = (R_k, \zeta_k)_{k \in \mathcal{K}^{\text{pr}}}$ has a forest structure. Because pruning a forest structure does not touch the atoms, note finally that if \mathcal{K} is complete then so is \mathcal{K}^{pr} .

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

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

The proof of Proposition 3.1 is given in Section 7.1.1.

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

Furthermore, pruning can be done really simply by following ? for $S = \mathbb{N}_m^*$, and pruning when appropriate. This gives the following ?, assuming, for simplicity, that the family is complete. Note that the only differences between ? and ? are the pruning step and ζ_k replacing $\zeta_k \wedge |S \cap R_k|$, because $\zeta_k \leq |R_k|$ and $S = \mathbb{N}_m^*$ here, 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 ? indeed produces the pruned region as in Definition 3.1.

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

The proof of Proposition 3.2 is given in Section 7.1.2.

Tip 2

We saw that ? has $O(N|S| + |\mathcal{K}|)$ complexity, the $N|S|$ term coming from the evaluation of the $|S \cap P_i|$ terms, $1 \leq i \leq N$. Here, $|S \cap P_i| = |\mathbb{N}_m^* \cap P_i| = |P_i|$ can be accessed in $O(1)$, and $N \leq |\mathcal{K}|$ for a complete family, so ? simply has $O(|\mathcal{K}|)$ complexity.

Algorithm 2 Pruning of a complete \mathfrak{R}

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

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

Let (i_1, \dots, i_m) a permutation of \mathbb{N}_m^* , eventually random, and, for all $t \in \mathbb{N}_m^*$, let $S_t = \{i_1, \dots, i_t\}$ 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 $t \in \{0, \dots, m\}$, this is what we call the curve of confidence bounds indexed by (i_1, \dots, i_m) . Applying ? to compute $V_{\mathfrak{R}}^*(S_t)$ for a given t has complexity $O(|\mathcal{K}|t)$, so using it to sequentially compute the full curve has complexity $O(|\mathcal{K}| \sum_{t=0}^m t) = O(|\mathcal{K}|m^2)$. In this section, we present a new algorithm that computes the curve with a $O(|\mathcal{K}|m)$ complexity. The algorithm will need that \mathfrak{R} is complete, so if that is not the case we first need to complete \mathfrak{R} following the Algorithm 2 of [Durand et al. \(2020\)](#), which has a $O(|\mathcal{K}|m)$ complexity. In the remainder of this section we assume that \mathfrak{R} is complete.

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

We will now present the new algorithm and the proof that it computes the curve $(V_{\mathfrak{R}}^*(S_t))_{t \in \mathbb{N}_m^*}$. We present two versions of the algorithm (strictly equivalent): one very formal (?), to introduce

additional notation used in the proof of Theorem 3.1, and, later, a simpler version that is the one actually implemented (?). Recall that a detailed illustration of the steps of the algorithms will be provided in Section 3.3.

In addition to the computation of all $V_{\mathfrak{R}}^*(S_t)$, ? also computes partitions \mathcal{P}^t that realize the minimum in (12) for $V_{\mathfrak{R}}^*(S_t)$. The initialization of \mathcal{P}^t has to be done carefully, for that we let $\mathcal{K}_0^- = \{k \in \mathcal{K} : \zeta_k = 0\}$ and

$$E = \{k \in \mathcal{K}_0^- : \forall k' \in \mathcal{K}_0^-, R_k \subseteq R_{k'} \Rightarrow k' = k\}, \quad (13)$$

$$F = \{(i, i), 1 \leq i \leq N : \forall k \in \mathcal{K}_0^-, R_{(i,i)} \not\subseteq R_k\}, \quad (14)$$

and finally

$$\mathcal{P}^0 = E \cup F. \quad (15)$$

E is the set of indices of the maximal elements k such that $\zeta_k = 0$, F is the set of indices of all leaves that are not a subset of a region indexed by E . \mathcal{P}^0 is the disjoint union of the two.

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{P}^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 $(\mathcal{K}_t)_{0 \leq t \leq m}$ is non-increasing for the inclusion relation, and that $\mathcal{K}_0 = \mathcal{K}$.

Theorem 3.1 (Fast curve computation). *Assume that \mathfrak{R} is complete in the sense of Section 2.3.*

Let any $t \in \mathbb{N}_m$. Then, $\mathcal{P}^t \in \mathfrak{P}$, and for all $k \in \mathcal{K}_t$, we have

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

and

$$V_{\mathfrak{R}}^*(S_t \cap R_k) = \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}|. \quad (17)$$

Furthermore,

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

The proof of this Theorem is given in Section 7.2. The first equality of Equation (18) states that the minimum in (12) is indeed 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). *Assume that \mathfrak{R} is complete in the sense of Section 2.3.*

For $t \in \{0, \dots, m-1\}$, $V_{\mathfrak{R}}^(S_{t+1}) = V_{\mathfrak{R}}^*(S_t)$ if $i_{t+1} \in \bigcup_{k \in \mathcal{K}_t^-} R_k$, and $V_{\mathfrak{R}}^*(S_{t+1}) = V_{\mathfrak{R}}^*(S_t) + 1$ if not.*

Algorithm 3 Formal computation of $(V_{\mathfrak{R}}^*(S_t))_{0 \leq t \leq m}$ with a complete family

```

1: procedure CURVE( $\mathfrak{R} = (R_k, \zeta_k)_{k \in \mathcal{K}}$  with  $\mathfrak{R}$  complete, path  $(S_t)_{1 \leq t \leq m}$  with  $S_t = \{i_1, \dots, i_t\}$ )
2:    $\mathcal{P}^0 \leftarrow E \cup F$  ▷ see (13) and (14)
3:    $\mathcal{K}_0^- \leftarrow \{k \in \mathcal{K} : \zeta_k = 0\}$ 
4:    $\eta_k^0 \leftarrow 0$  for all  $k \in \mathcal{K}$ 
5:   for  $t = 1, \dots, m$  do
6:     if  $i_t \in \bigcup_{k \in \mathcal{K}_{t-1}^-} R_k$  then
7:        $\mathcal{P}^t \leftarrow \mathcal{P}^{t-1}$ 
8:        $\mathcal{K}_t^- \leftarrow \mathcal{K}_{t-1}^-$ 
9:        $\eta_k^t \leftarrow \eta_k^{t-1}$  for all  $k \in \mathcal{K}$ 
10:    else
11:      for  $h = 1, \dots, h_{\max}(t)$  do
12:         $\eta_{k^{(t,h)}}^t \leftarrow \eta_{k^{(t,h)}}^{t-1} + 1$ 
13:        if  $\eta_{k^{(t,h)}}^t < \zeta_k$  then
14:          Pass
15:        else
16:           $h_t^f \leftarrow h$  ▷ final depth
17:           $\mathcal{P}^t \leftarrow \left( \mathcal{P}^{t-1} \setminus \{k \in \mathcal{P}^{t-1} : R_k \subseteq R_{k^{(t,h_t^f)}}\} \right) \cup \{k^{(t,h_t^f)}\}$ 
18:           $\mathcal{K}_t^- \leftarrow \mathcal{K}_{t-1}^- \cup \{k^{(t,h_t^f)}\}$ 
19:          Break the loop
20:        end if
21:      end for
22:      if the loop has been broken then
23:         $\eta_k^t \leftarrow \eta_k^{t-1}$  for all  $k \in \mathcal{K}$  not visited during the loop, that is all  $k \notin \{k^{(t,h)}, 1 \leq$ 
24:         $h \leq h_t^f\}$ 
25:      else
26:         $\mathcal{P}^t \leftarrow \mathcal{P}^{t-1}$ 
27:         $\mathcal{K}_t^- \leftarrow \mathcal{K}_{t-1}^-$ 
28:         $\eta_k^t \leftarrow \eta_k^{t-1}$  for all  $k \in \mathcal{K}$  not visited during the loop, that is all  $k \notin \{k^{(t,h)}, 1 \leq$ 
29:         $h \leq h_{\max}(t)\}$ 
30:      end if
31:    end for
32:    return  $\mathcal{P}^t, \eta_k^t$  for all  $t = 1, \dots, m$  and  $k \in \mathcal{K}$ 
33: end procedure

```

Proof. From (18), $V_{\mathfrak{R}}^*(S_{t+1}) = \sum_{k \in \mathcal{K}^1} \eta_k^{t+1}$ and $V_{\mathfrak{R}}^*(S_t) = \sum_{k \in \mathcal{K}^1} \eta_k^t$. 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 $(V_{\mathfrak{R}}^*(S_t))_{1 \leq t \leq m}$, tracking \mathcal{P}^t is actually useless, what is important is to track and update \mathcal{K}_t^- correctly. Hence the simpler, alternative ?. Note that ? is less formal than ? : as in ? and ?, it recycles notation (mimicking the actual code implementation) so the t subscript or superscript is dropped from all \mathcal{K}_t^- and η_k^t . In ?, the notation V_t is actually equal to $V_{\mathfrak{R}}^*(S_t)$ by Corollary 3.1.

Stocking, for each i_t , the indices k such that $i_t \in R_k$, is done by scanning the forest structure so it has complexity in $O(|\mathcal{K}|)$. Once this information is available, finding $k^{(t,h)}$ and updating \mathcal{K}^- can be done

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

```
1: procedure CURVE( $\mathfrak{R} = (R_k, \zeta_k)_{k \in \mathcal{K}}$  with  $\mathfrak{R}$  complete, path  $(S_t)_{1 \leq t \leq m}$  with  $S_t = \{i_1, \dots, i_t\}$ )
2:    $V_0 \leftarrow 0$ 
3:    $\mathcal{K}^- \leftarrow \{k \in \mathcal{K} : \zeta_k = 0\}$ 
4:    $\eta_k \leftarrow 0$  for all  $k \in \mathcal{K}$ 
5:   for  $t = 1, \dots, m$  do
6:     if  $i_t \in \bigcup_{k \in \mathcal{K}^-} R_k$  then
7:        $V_t \leftarrow V_{t-1}$ 
8:     else
9:       for  $h = 1, \dots, h_{\max}(t)$  do
10:        find  $k^{(t,h)} \in \mathcal{K}^h$  such that  $i_t \in R_{k^{(t,h)}}$ 
11:         $\eta_{k^{(t,h)}} \leftarrow \eta_{k^{(t,h)}} + 1$ 
12:        if  $\eta_{k^{(t,h)}} < \zeta_k$  then
13:          pass
14:        else
15:           $\mathcal{K}^- \leftarrow \mathcal{K}^- \cup \{k^{(t,h)}\}$ 
16:          break the loop
17:        end if
18:      end for
19:       $V_t \leftarrow V_{t-1} + 1$ 
20:    end if
21:  end for
22:  return  $(V_t)_{1 \leq t \leq m}$ 
23: end procedure
```

in $O(1)$. Then each step t of the for loop consists in two successive scans of the $k^{(t,h)}$, $1 \leq h \leq H$, the first to check if $i_t \in \bigcup_{k \in \mathcal{K}^-} R_k$, and the second to update the $\eta_{k^{(t,h)}}$ if $i_t \notin \bigcup_{k \in \mathcal{K}^-} R_k$. So each step has complexity in $O(H)$ and finally the complexity of ? is in $O(Hm + |\mathcal{K}|) \leq O(|\mathcal{K}|m)$, and even only $O(Hm)$ after the first call if the necessary information has been stocked.

3.3 Illustration on a detailed example

In this section, we follow ? during its first steps in a detailed fashion.

We keep the structure of Example 2.2 and Example 2.3. Recall that $m = 25$, $P_{1:5} = R_1 = \{1, \dots, 20\}$, $P_1 = R_2 = \{1, 2\}$, $P_{2:3} = R_3 = \{3, \dots, 10\}$, $P_{4:5} = R_4 = \{11, \dots, 20\}$, $P_2 = \{3, 4\}$, $P_3 = R_5 = \{5, \dots, 10\}$, $P_4 = R_6 = \{11, \dots, 16\}$, $P_5 = R_7 = \{17, \dots, 20\}$, $P_{6:7} = R_8 = \{21, 22\}$, $P_6 = \{21\}$, $P_7 = R_9 = \{22\}$ and $P_8 = \{23, 24, 25\}$.

Now assume that we have the following values for the ζ_k 's: $\zeta_{(1,5)} = 5$, $\zeta_{(1,1)} = 2$, $\zeta_{(2,3)} = 0$, $\zeta_{(3,3)} = 0$, $\zeta_{(4,5)} = 4$, $\zeta_{(4,4)} = 2$, $\zeta_{(5,5)} = 3$, $\zeta_{(6,7)} = 2$, $\zeta_{(7,7)} = 0$. 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 4.

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 $i_1 = 11, i_2 = 17, i_3 = 12, i_4 = 13, i_5 = 18, i_6 = 24, i_7 = 19, i_8 = 22$ and $i_9 = 5$.

First, we apply ? to the family. This results in pruning $P_{6:7}$ (and only this region), because $2 = \zeta_{(6,7)} \geq \zeta_{(6,6)} + \zeta_{(7,7)} = 1 + 0$. This gives Figure 5.

Now we initialize ?, that is we let $t = 0$. Because $\zeta_{(2,3)} = \zeta_{(3,3)} = \zeta_{(7,7)} = 0$, $(2, 3)$, $(3, 3)$ and $(7, 7)$ are added to \mathcal{K}_t^- : $\mathcal{K}_0^- = \{(2, 3), (3, 3), (7, 7)\}$. We define \mathcal{P}^0 according to (13), (14) and (15). Here,

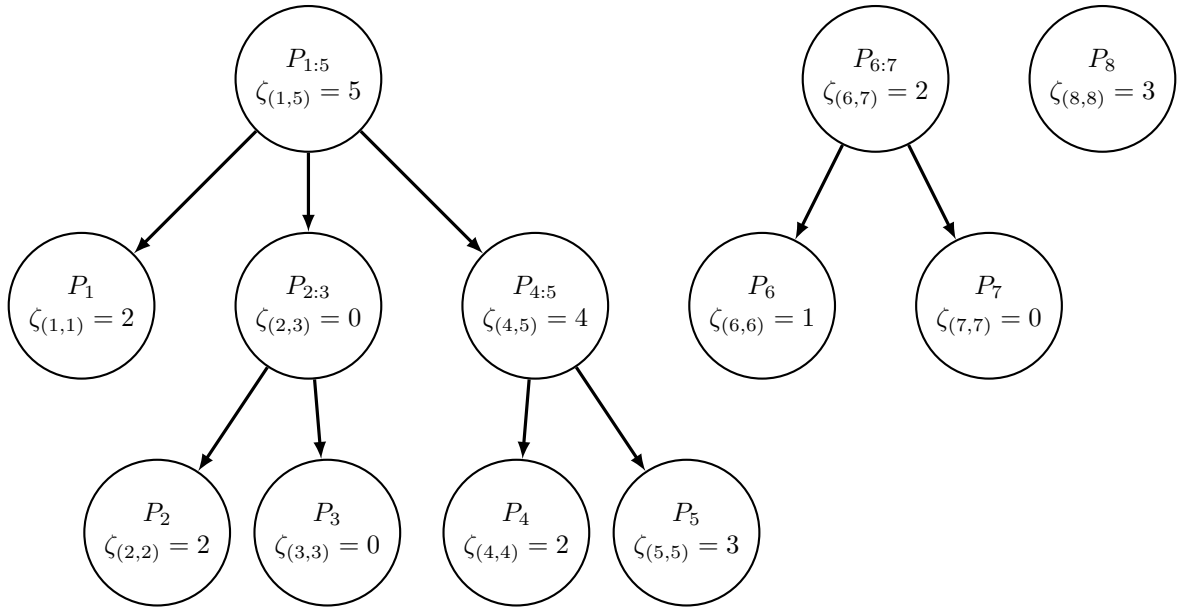


Figure 4: The regions of Example 2.2 with the ζ_k values.

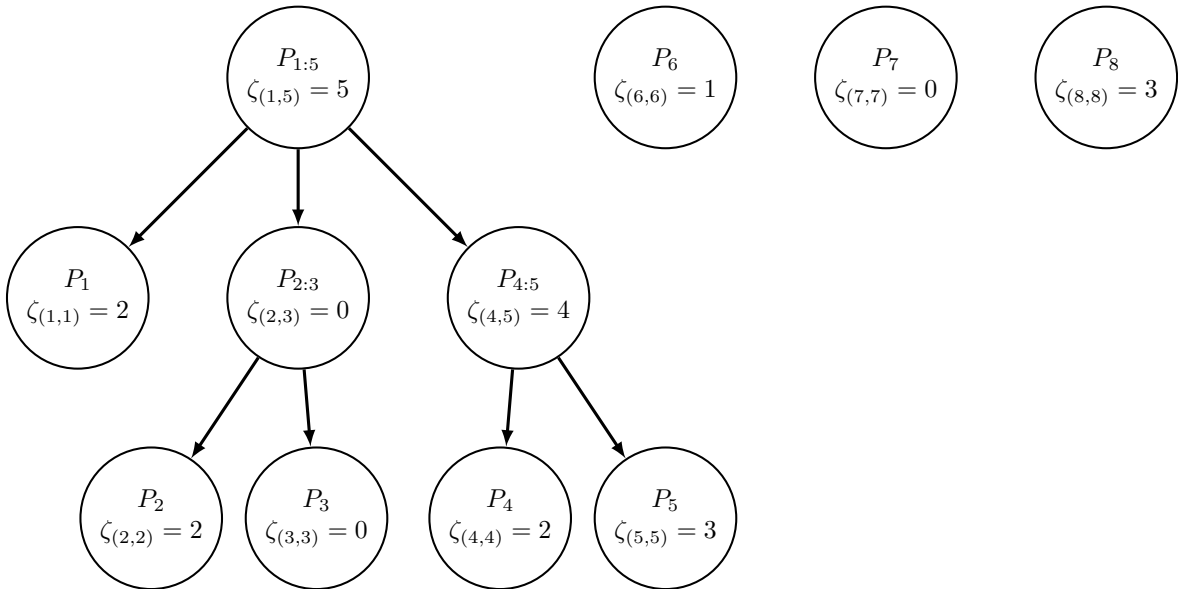


Figure 5: The regions of Example 2.2 after pruning.

$E = \{(2, 3), (7, 7)\}$ and so $F = \{(1, 1), (4, 4), (5, 5), (6, 6), (8, 8)\}$ and $\mathcal{P}^0 = E \cup F$. Furthermore, all η_k^t are set to 0. The initial state of ? is shown in Figure 6, with the elements of \mathcal{K}_t^- being in red to show that they will not contribute to the computations, and the elements of \mathcal{P}^t as squares.

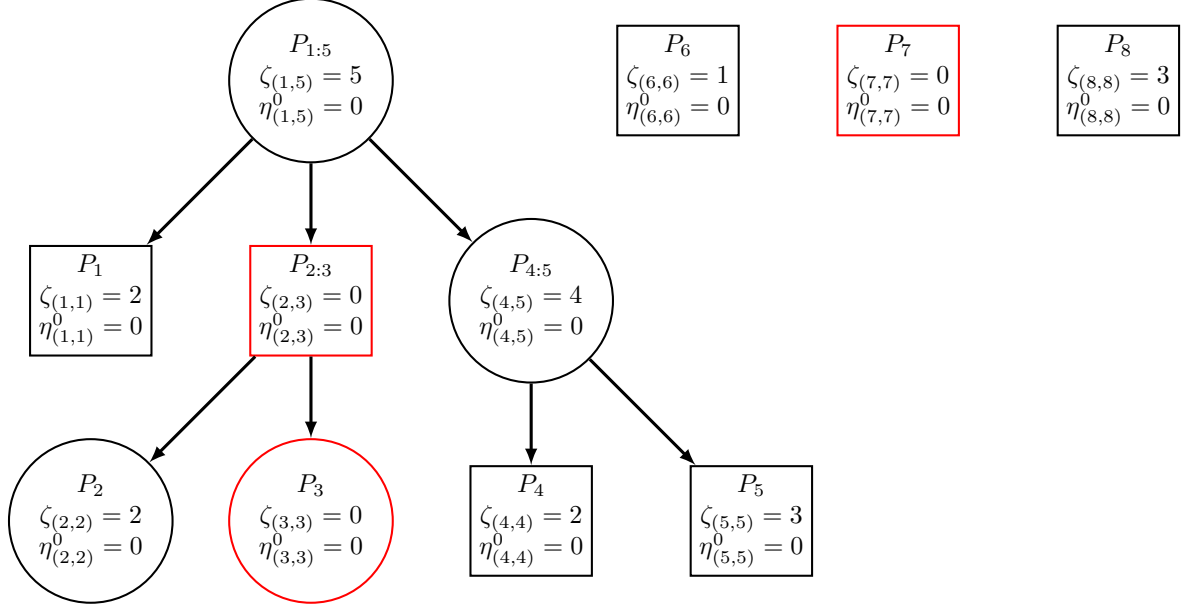


Figure 6: The regions of Example 2.2 at $t = 0$ in ?.

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 η_k^t are increased by one, and by (18) we have $V_{\mathfrak{N}}^*(S_1) = \eta_{(1,5)}^1 + \eta_{(6,6)}^1 + \eta_{(7,7)}^1 + \eta_{(8,8)}^1 = 1 + 0 + 0 + 0 = 1$. The state of the step is summarized in Figure 7.

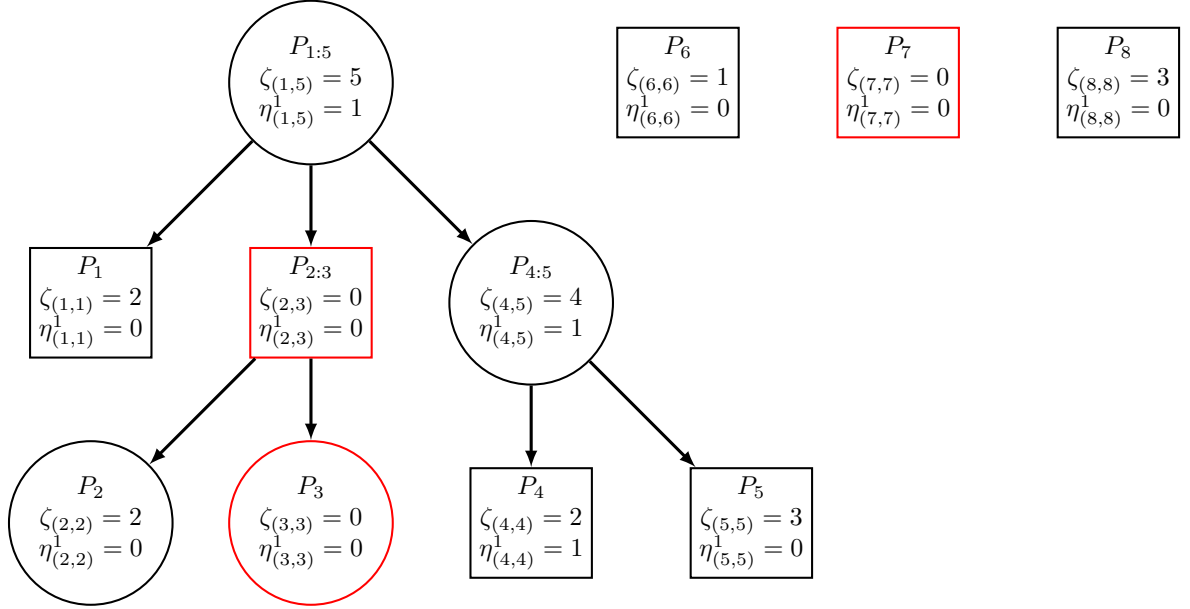


Figure 7: The regions of Example 2.2 at $t = 1$ in ?.

We move on to $t = 2$, with $i_2 = 17$. $i_2 \in P_5 \subseteq P_{4:5} \subseteq P_{1:5}$. The appropriate η_k^t are increased by one, and by (18) we have $V_{\mathfrak{N}}^*(S_2) = 2$. The state of the step is summarized in Figure 8.

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

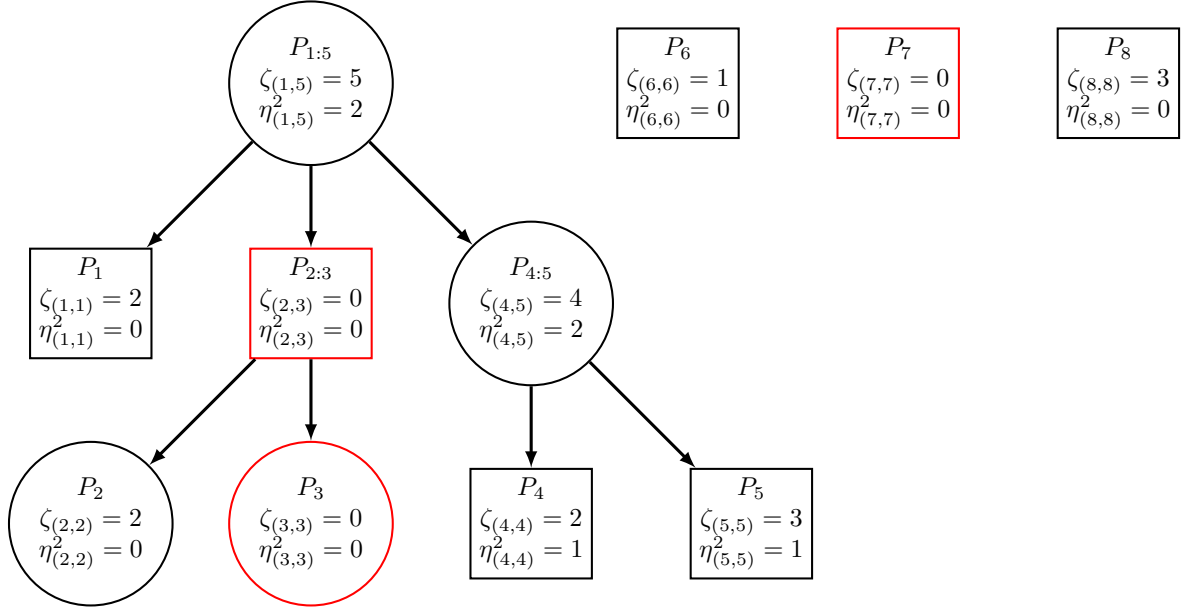


Figure 8: The regions of Example 2.2 at $t = 2$ in ?.

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^- : $\mathcal{K}_3^- = \{(2, 3), (3, 3), (4, 4), (7, 7)\}$. Following line 17 of ?, \mathcal{P}^t does not change (we remove then add $(4, 4)$ from it) and $\mathcal{P}^3 = \mathcal{P}^0$. By (18), we have $V_{\mathfrak{R}}^*(S_3) = 3$. The state of the step is summarized in Figure 9, with P_4 now also in red.

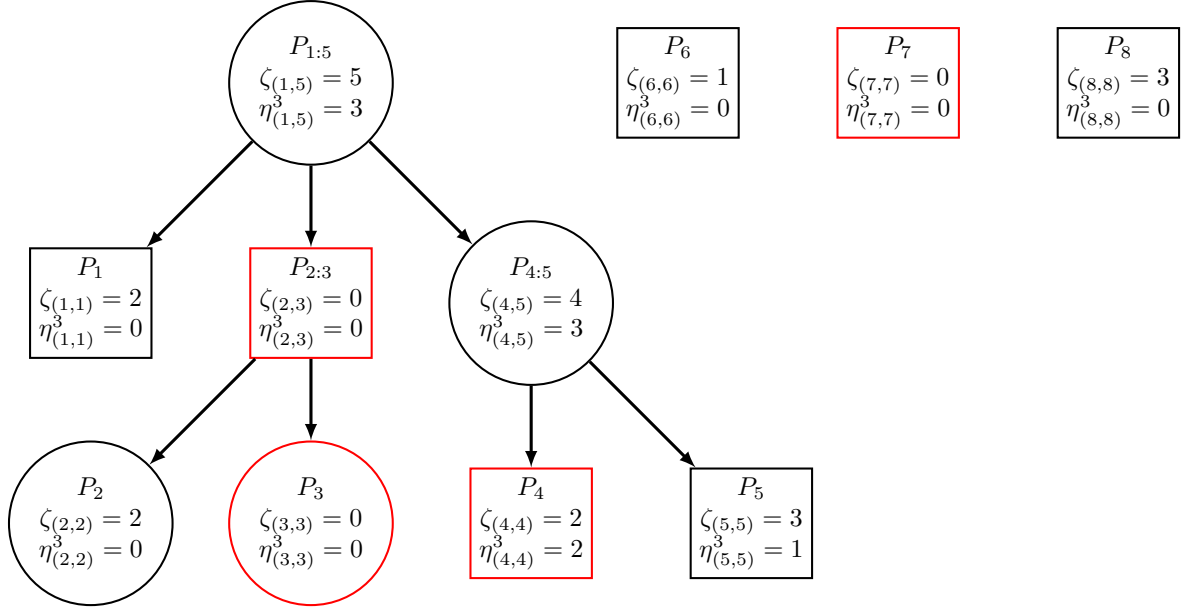


Figure 9: The regions of Example 2.2 at $t = 3$ in ?.

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

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 $\eta_{(1,5)}^t$: $\eta_{(1,5)}^5 = 4 < \zeta_{(1,5)}$, then $\eta_{(4,5)}^t$: $\eta_{(4,5)}^5 = 4$, and we stop there because $\eta_{(4,5)}^5 = 4 = \zeta_{(4,5)}$. $P_{4:5}$ will stop contributing, we add it to \mathcal{K}_t^- : $\mathcal{K}_5^- = \{(2, 3), (4, 5), (3, 3), (4, 4), (7, 7)\}$. We also add $(4, 5)$ and

remove $(4, 4)$ and $(5, 5)$ from \mathcal{P}^t : $\mathcal{P}^5 = \{(1, 1), (2, 3), (4, 5), (6, 6), (7, 7), (8, 8)\}$. Note that $\eta_{(5,5)}^t$ is not updated because we stopped the loop before, see line 23 of ?. By (18), we have $V_{\mathfrak{A}}^*(S_5) = 4$. The state of the step is summarized in Figure 10, with $P_{4:5}$ now also in red.

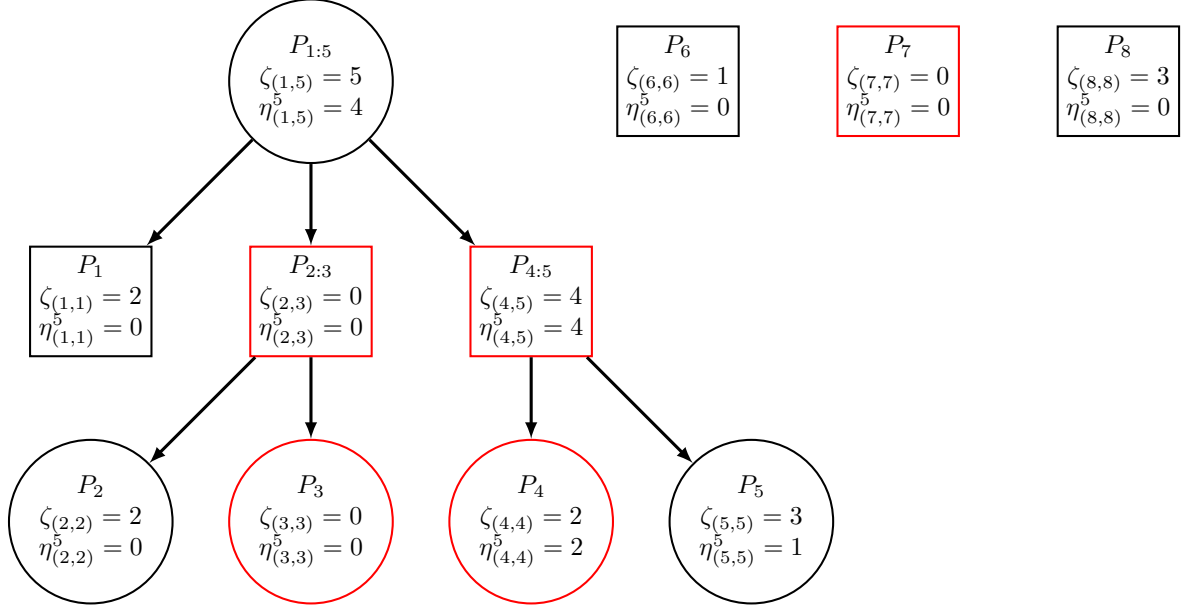


Figure 10: The regions of Example 2.2 at $t = 5$ in ?.

We move on to $t = 6$, with $i_6 = 24$. $i_6 \in P_8$. The appropriate η_k^t is increased by one: $\eta_{(8,8)}^6 = 1 < \zeta_{(8,8)}$, and by (18) we have $V_{\mathfrak{A}}^*(S_6) = \eta_{(1,5)}^6 + \eta_{(6,6)}^6 + \eta_{(7,7)}^6 + \eta_{(8,8)}^6 = 4 + 0 + 0 + 1 = 5$. The state of the step is summarized in Figure 11.

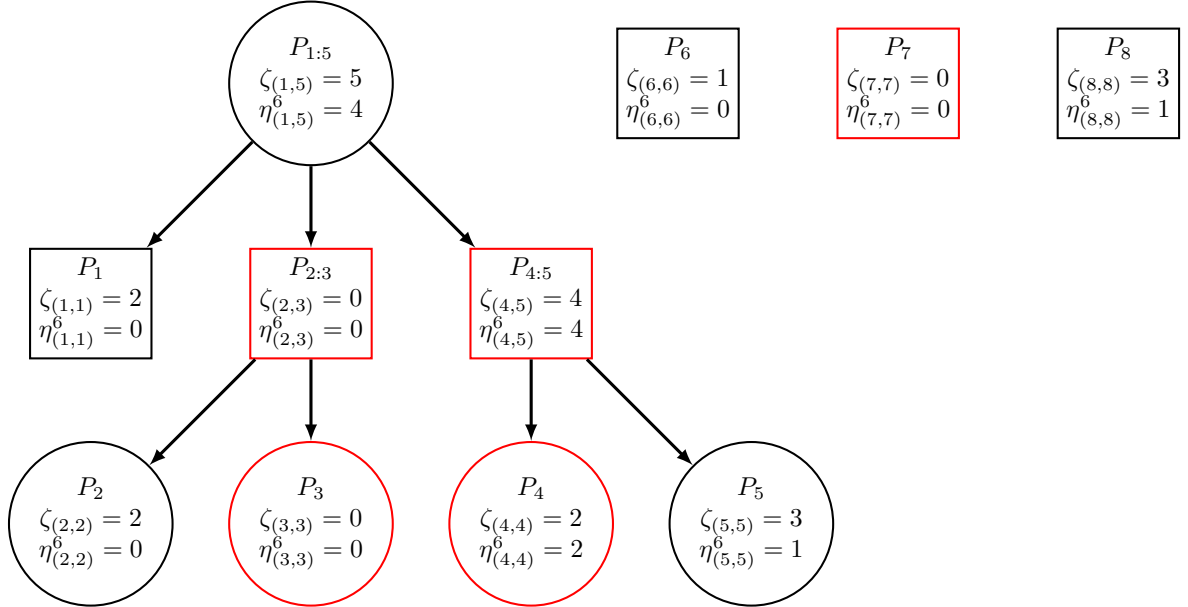


Figure 11: The regions of Example 2.2 at $t = 6$ in ?.

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 ?), and by (18), we have $V_{\mathfrak{A}}^*(S_7) = V_{\mathfrak{A}}^*(S_8) = V_{\mathfrak{A}}^*(S_9) = 5$.

4 Implementation

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

Speaking of the data structure, we briefly describe it, with an example. We represent $(R_k)_{k \in \mathcal{K}}$ by two lists, `C` and `leaf_list`. `leaf_list` is a list of vectors, where `leaf_list[[i]]` is the vector listing the hypotheses in the atom P_i . `C` is a list of lists. For $1 \leq h \leq H$, `C[[h]]` lists the regions at depth h , using the index bounds of the atoms they are composed of. That is, the elements of the list `C[[h]]` are vectors of size two, and if there is k, i and j such that `C[[h]][[k]] = c(i, j)`, it means that $(i, j) \in \mathcal{K}$, or in other words that $R_{(i,j)} = P_{i:j}$ is one of the regions, and that $\phi((i, j)) = h$.

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

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

Caution 1

We emphasize that the 1D structure of the hypotheses has to be respected by the user as the current implementation implicitly uses it: that is, P_1 has to contain the hypotheses labeled $1, 2, \dots, p$, P_2 has to contain the hypotheses labeled $p + 1, \dots$, and so on. Also, the hypotheses have to be in increasing order: `leaf_list[[1]]` has to be equal to `c(1, 2, 3, ..., p)` and not, say, `c(2, 1, 3, ..., p)`.

Tip 3

We see that the current implementation requires to provide a partition $(P_n)_{1 \leq n \leq N}$ of leaves compatible with the reference family. So, in a way, we always provide a complete family. However, not all `c(i, i)` have to be in `C` for the various algorithms implemented to function properly. It is not needed for the implementations of `?` and `?` given that, as stated in Tip 1, they start by computing $|S \cap P_n|$ for each leaf P_n anyway. Having a complete `C` is not needed either for `?`. Indeed, if R_k is a region such that $\zeta_k = |R_k|$, the condition of line 12 of `?` is always true, except for the last element of R_k that is added to S_t . At that point, because the elements of R_k have been exhausted, R_k won't ever be visited again, so adding it to \mathcal{K}^- is irrelevant. In the

end, tracking R_k and η_k for such k was not necessary in the first place. Furthermore, if the implementation does not find $k^{(t,h)}$ (see line 10), it simply pass to the next iteration of the for loop.

The functions `dyadic.from.leaf_list`, `dyadic.from.window.size`, and `dyadic.from.height` return the appropriate data structure to represent a \mathcal{K} that can be described as a dyadic tree, based on some entry parameters that can be inferred from the names of the functions. As said in Tip 3, the completion of \mathcal{C} given `leaf_list` is not necessary, but can be done by the `forest.completion` function. Finally, the ζ_k 's are computed as in Durand et al. (2020) by the `zetas.tree` function with `method=zeta.DKWM`. Using `method=zeta.trivial` just yields $\zeta_k = |R_k|$.

The following R snippet constructs the family of Example 2.2, draws uniform p -values p_1, \dots, p_{25} , computes the ζ_k 's with `method=zeta.trivial`, completes the family with `forest.completion` to get the family of Example 2.3, allowing to verify the claim of Example 4.1, prunes it with `pruning` (which prunes everything except the leaves because of `zeta.trivial`), and finally computes the curve of confidence bounds on the path $S_t = \{\sigma(1), \dots, \sigma(t)\}$, using both the pruned and non-pruned complete family, where σ is a permutation ordering the p -values, using the fast implementation `curve.V.star.forest.fast`.

```
library(sanssouci)

leaf_list <- list(c(1, 2),
                 c(3, 4),
                 c(5, 6, 7, 8, 9, 10),
                 c(11, 12, 13, 14, 15, 16),
                 c(17, 18, 19, 20),
                 21,
                 22,
                 c(23, 24, 25))
C <- list(list(c(1, 5), c(6,7)),
          list(c(1, 1), c(2, 3), c(4, 5), c(7, 7)),
          list(c(3, 3), c(4, 4), c(5, 5)))
pvalues <- runif(25)
o <- order(pvalues)
ZL <- zetas.tree(C, leaf_list, zeta.trivial, pvalues, alpha = 0.05)
complete.res <- forest.completion(C, ZL, leaf_list)
curve.V.star.forest.fast(o, complete.res$C, complete.res$ZL, leaf_list)

[1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25

pruning.res <- pruning(complete.res$C, complete.res$ZL, leaf_list, delete.gaps = TRUE)
curve.V.star.forest.fast(o, pruning.res$C, pruning.res$ZL, leaf_list)

[1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25
```

5 Numerical experiments

In this Section, we present some numerical experiments aiming to demonstrate the impact of the pruning of ? (using the `delete.gaps` option mentioned in Section 4) and of the fast ?, 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 ?.

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 0.14.1, 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. The p -values are generated in a gaussian one-sided fashion (see Example 2.1) where $H_{0,i} = \{\mathcal{N}(\boldsymbol{\mu}, \text{Id}) : \mu_i = 0\}$, $H_{1,i} = \{\mathcal{N}(\boldsymbol{\mu}, \text{Id}) : \mu_i = 4\}$. \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 $(V_{\mathfrak{A}}^*(\{1, \dots, t\}))_{t \in \mathbb{N}_m^*}$ is computed. For the experiments including pruning, the pruning is done once before the `n_repl` replications, to mimic 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	1024	1024	10240	10240
zeta computation	trivial	DKWM	trivial	DKWM
<code>n_repl</code>	100	100	10	10

For the trivial ζ_k computation of scenarios 1 and 3, the pruning obviously deletes all non-atom regions so $|\mathcal{K}^{\text{pr}}| = 512$. Whereas, for the particular instance $\omega \in \Omega$ in the experiments, $|\mathcal{K}^{\text{pr}}| = 541$ for scenario 2, and $|\mathcal{K}^{\text{pr}}| = 573$ 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)).

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 12. The time unit is the second.

Table 2: Scenario 1

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	3.6708287	3.8028650	3.8149199	3.8221756	3.8362092	3.9022797	100
naive.pruned	3.3147519	3.4198975	3.4353463	3.4470054	3.4657886	3.5459636	100
fast.not.pruned	0.0035286	0.0035779	0.0046194	0.0036011	0.0036321	0.1014023	100
fast.pruned	0.0011960	0.0012314	0.0012535	0.0012430	0.0012703	0.0013603	100

Table 3: Scenario 2

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	3.7152477	3.8110591	3.9803535	3.8483790	3.9549886	10.1338336	100
naive.pruned	3.3277028	3.4592016	3.5465768	3.5060270	3.6059210	5.4159371	100
fast.not.pruned	0.0032789	0.0033216	0.0067553	0.0033482	0.0033857	0.1978229	100
fast.pruned	0.0013597	0.0013884	0.0014134	0.0014056	0.0014298	0.0017731	100

Table 4: Scenario 3

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	336.0473732	336.7254511	338.6804399	337.0286221	340.9009506	344.4716282	10
naive.pruned	332.4762463	332.8188433	334.4660587	334.1282526	335.5376761	337.6580202	10
fast.not.pruned	0.0323725	0.0324755	0.0328789	0.0325803	0.0328097	0.0354455	10
fast.pruned	0.0099485	0.0100272	0.0101948	0.0101886	0.0102164	0.0107677	10

Table 5: Scenario 4

expr	min	lq	mean	median	uq	max	neval
naive.not.pruned	340.4702704	341.4280632	344.8181652	344.3519587	348.6074564	350.4574742	10
naive.pruned	337.2238865	338.1905987	340.4203488	340.4743933	342.7488030	344.1039957	10
fast.not.pruned	0.0294732	0.0296390	0.0299436	0.0298885	0.0300172	0.0307673	10
fast.pruned	0.0124157	0.0126186	0.0137188	0.0127803	0.0130546	0.0194847	10

In each scenario, using the fast algorithm is much faster than the naive approach, with a speed factor of at least 1000. Using the naive approach, pruning always gives a slight improvement over not pruning. Using the fast algorithm, the benefits of pruning are significant, with a speed factor of at least 2, and sometimes 3.

Comparing scenarios 1 and 2 first, we see that, as expected, there is no significant change in computation time for `naive.not.pruned`. Methods `naive.pruned` and `fast.pruned` are faster in scenario 1, given that we prune more. But, on the other hand, `fast.not.pruned` is slightly faster in scenario 2. 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. Although, with only `n_rep1=10`, the statistics seem less accurate, this can be confirmed with additional experiments (`n_rep1` can also be set to 100 without problem if 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 ~ 10 when using ? and by ~ 100 when using ? naively, which illustrates the stated complexities of $O(|\mathcal{K}|m)$ and $O(|\mathcal{K}|m^2)$, respectively.

6 Conclusion

In conclusion, we effectively introduced a new algorithm to compute a curve of confidence upper bounds for the false discoveries, or, equivalently, for the FDP, that is much faster than the previous

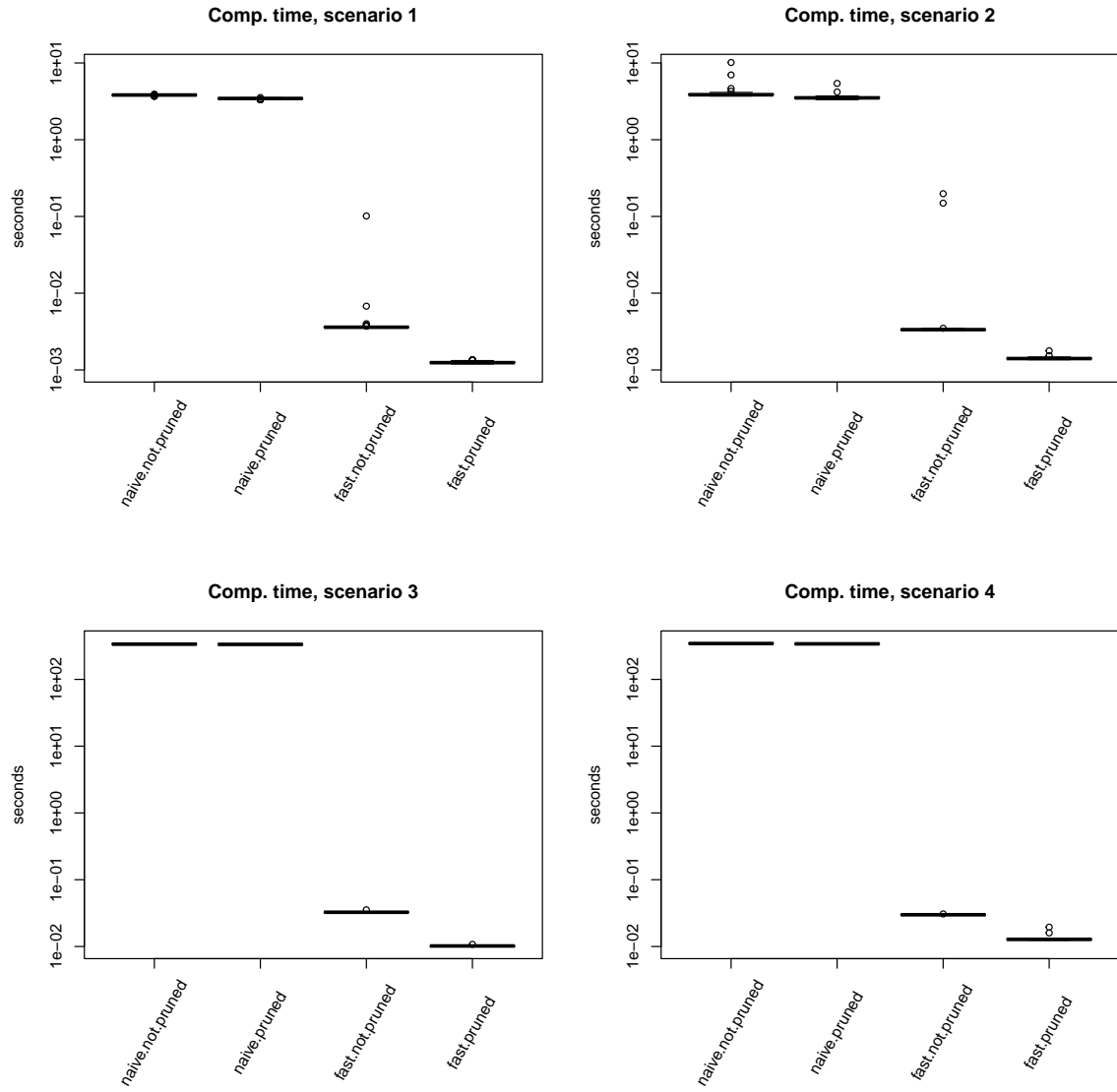


Figure 12: Computation times in each scenario, in seconds (using a logarithmic scale)

alternative, with one power of m less in the complexity. This algorithm can be applied as soon as the confidence upper bound is built according to the JER framework, when the reference family exhibit a forest structure.

To develop new confidence upper bound methods 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 theoretically proven 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 Proofs

7.1 Proofs of Section 3.1

7.1.1 Proof of Proposition 3.1

Recall Equation (11) and, because \mathfrak{R}^{pr} also has a forest structure,

$$V_{\mathfrak{R}^{\text{pr}}}^*(S) = \min_{Q \subseteq \mathcal{K}^{\text{pr}}} \left(\sum_{k' \in Q} \zeta_{k'} \wedge |S \cap R_{k'}| + \left| S \setminus \bigcup_{k' \in Q} R_{k'} \right| \right), \quad (19)$$

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

Let any $Q \subseteq \mathcal{K}$. We split Q in A elements of $\mathcal{K} \setminus \mathcal{K}^{\text{pr}}$, 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}^{pr} , simply denoted k_b , $1 \leq b \leq B$. By the definition of \mathcal{K}^{pr} and the previous remarks, for any $1 \leq a \leq A$, there exist integers $i_{1,a}, \dots, i_{p_a-1,a}$ such that $i_{0,a} < i_{1,a} < \dots < i_{p_a-1,a} < i_{p_a,a}$, $(i_{j-1,a}, i_{j,a} - 1) \in \mathcal{K}^{\text{pr}}$ 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

$$Q^{\text{pr}} = \{k_b : 1 \leq b \leq B\} \cup \{(i_{j-1,a}, i_{j,a} - 1) : 1 \leq a \leq A, 1 \leq j \leq p_a\}. \quad (20)$$

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

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

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

$$\begin{aligned}\zeta_{(i_0,a,i_{p_a,a}-1)} &\geq \sum_{j=1}^{p_a} \zeta_{(i_{j-1,a},i_{j,a}-1)} \\ &\geq \sum_{j=1}^{p_a} \zeta_{(i_{j-1,a},i_{j,a}-1)} \wedge |S \cap R_{(i_{j-1,a},i_{j,a}-1)}|,\end{aligned}$$

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

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

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

$$\begin{aligned}\sum_{k \in Q} \zeta_k \wedge |S \cap R_k| + \left| S \setminus \bigcup_{k \in Q} R_k \right| &\geq \sum_{k \in Q^{\text{pr}}} \zeta_k \wedge |S \cap R_k| + \left| S \setminus \bigcup_{k \in Q^{\text{pr}}} R_k \right| \\ &\geq V_{\mathfrak{R}^{\text{pr}}}^*(S).\end{aligned}\tag{21}$$

Note that Equation (21) 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 $V_{\mathfrak{R}}^*(S) \geq V_{\mathfrak{R}^{\text{pr}}}^*(S)$.

7.1.2 Proof of Proposition 3.2

First, $\mathcal{K} \setminus \mathcal{L} \subseteq \mathcal{K} \setminus \mathcal{K}^{\text{pr}}$ is trivial: a k such that $\zeta_k \geq \sum_{k' \in \text{Succ}_k} \text{Vec}_{k'}$ obviously satisfies the condition of Definition 3.1 to be pruned.

Now let $(i, i') \in \mathcal{K} \setminus \mathcal{K}^{\text{pr}}$ 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 $1 \leq j \leq p$ and finally $\zeta_{(i,i')} = \zeta_{(i_0,i_p-1)} \geq \sum_{j=1}^p \zeta_{(i_{j-1},i_j-1)}$. Then by the proof of Theorem 1 of Durand et al. (2020) but applied to $S = R_{(i,i')}$ we have that $\sum_{j=1}^p \zeta_{(i_{j-1},i_j-1)} \geq \sum_{k' \in \text{Succ}_{(i,i')}} \text{Vec}_{k'}$ (see the unnumbered line just above Equation (A4) in that paper) and so $\zeta_{(i,i')} \geq \sum_{k' \in \text{Succ}_{(i,i')}} \text{Vec}_{k'}$ hence (i, i') is pruned by ? and $\mathcal{K} \setminus \mathcal{K}^{\text{pr}} \subseteq \mathcal{K} \setminus \mathcal{L}$.

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

7.2 Proof of Theorem 3.1

In this section, every reference to a line is a reference to a line of ?.

7.2.1 Derivation of (18)

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

$$Q = \bigcup_{k \in \mathcal{K}^1} \{k' \in Q : R_{k'} \subseteq R_k\} \tag{22}$$

and the union is disjoint. From (12), let $Q^* \in \mathfrak{P}$ such that $V_{\mathfrak{A}}^*(S_t) = \sum_{k' \in Q^*} \zeta_{k'} \wedge |S_t \cap R_{k'}|$. Then by (22),

$$\begin{aligned}
V_{\mathfrak{A}}^*(S_t) &= \sum_{k' \in Q^*} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\
&= \sum_{k \in \mathcal{K}^1} \sum_{\substack{k' \in Q^* \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\
&= \sum_{k \in \mathcal{K}^1} \sum_{\substack{k' \in Q^* \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap (R_k \cap R_{k'})| \\
&= \sum_{k \in \mathcal{K}^1} \sum_{k' \in Q^*} \zeta_{k'} \wedge |(S_t \cap R_k) \cap R_{k'}| \tag{23} \\
&\geq \sum_{k \in \mathcal{K}^1} V_{\mathfrak{A}}^*(S_t \cap R_k), \tag{24}
\end{aligned}$$

where the equality in (23) comes from the fact that if $R_{k'} \not\subseteq R_k$, then $R_{k'} \cap R_k = \emptyset$, that is, $R_k \subsetneq R_{k'}$ is impossible because $k \in \mathcal{K}^1$. Furthermore, (24) holds again by (12).

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

$$\begin{aligned}
\sum_{k \in \mathcal{K}^1} V_{\mathfrak{A}}^*(S_t \cap R_k) &= \sum_{k \in \mathcal{K}^1} \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\
&= \sum_{k \in \mathcal{P}^t} \zeta_k \wedge |S_t \cap R_k| \text{ by (22)} \\
&\geq V_{\mathfrak{A}}^*(S_t) \text{ by (12)}.
\end{aligned}$$

So we proved that $V_{\mathfrak{A}}^*(S_t) = \sum_{k \in \mathcal{P}^t} \zeta_k \wedge |S_t \cap R_k| = \sum_{k \in \mathcal{K}^1} V_{\mathfrak{A}}^*(S_t \cap R_k)$ and finally $V_{\mathfrak{A}}^*(S_t) = \sum_{k \in \mathcal{K}^1} V_{\mathfrak{A}}^*(S_t \cap R_k) = \sum_{k \in \mathcal{K}^1} \eta_k^t$ by (16), again because $\mathcal{K}^1 \subseteq \mathcal{K}_t$. Every equality in (18) is proven.

7.2.2 Proof that $\mathcal{P}^t \in \mathfrak{P}$

By completeness, $F \subseteq \mathcal{K}$ and so $\mathcal{P}^0 \subseteq \mathcal{K}$. First we show that $\mathbb{N}_m^* = \bigcup_{k \in \mathcal{P}^0} R_k$. Let $j \in \mathbb{N}_m^*$ and $i \in \mathbb{N}_N^*$ such that $j \in R_{(i,i)}$. Let $G = \{k \in \mathcal{K}_0^- : j \in R_k\}$. If $G = \emptyset$, for any $k \in \mathcal{K}_0^-$, $R_{(i,i)} \subseteq R_k$ would imply that $j \in R_k$ and $k \in G$, hence a contradiction, and so $(i, i) \in F$ and $j \in \bigcup_{k \in \mathcal{P}^0} R_k$. If $G \neq \emptyset$, for any $k, k' \in G$, $j \in R_k \cap R_{k'}$ so, by forest structure, $R_k \subseteq R_{k'}$ or $R_{k'} \subseteq R_k$, hence \subseteq is a total order on the finite, non-empty set $\{R_k : k \in G\}$, so the latter has a maximum and there exists a unique $k^* \in G$ such that $R_{k^*} = \max_{k \in G} R_k$. Let us show that, as an element of \mathcal{K}_0^- , k^* is maximal, which will imply that $k^* \in E$ and that $j \in \bigcup_{k \in \mathcal{P}^0} R_k$. Let any $k' \in \mathcal{K}_0^-$ such that $R_{k^*} \subseteq R_{k'}$, then $j \in R_{k'}$, so $k' \in G$ and so $R_{k'} \subseteq \max_{k \in G} R_k = R_{k^*}$. Hence $R_{k^*} = R_{k'}$ and $k^* = k'$ (see Remark 2.2), k^* is indeed maximal, $k^* \in E$ and $j \in \bigcup_{k \in \mathcal{P}^0} R_k$. This proves that $\mathbb{N}_m^* = \bigcup_{k \in \mathcal{P}^0} R_k$.

Now let us prove that the elements \mathcal{P}^0 index disjoint sets. Let $k, k' \in \mathcal{P}^0$ such that there exists $j \in R_k \cap R_{k'}$. By forest structure, $R_k \subseteq R_{k'}$ or $R_{k'} \subseteq R_k$. Now we separate four cases. The case $k \in F, k' \in E$ is impossible, because k would be the index of an atom, so we would imperatively have $R_k \subseteq R_{k'}$, which would contradict the definition of F . Similarly, $k \in F', k' \in E$ is impossible. If $k, k' \in E$, then $k = k'$ by the very definition of E . Finally, if $k \in F, k' \in F'$, R_k and $R_{k'}$ are both atoms, and because the atoms realize a partition of \mathbb{N}_m^* , then $k = k'$. In all cases, $k = k'$ which concludes.

The $R_k, k \in \mathcal{P}^0$, are disjoint, non-empty (see Remark 2.2), and cover \mathbb{N}_m^* , so they form a partition of \mathbb{N}_m^* , in other words, $\mathcal{P}^0 \in \mathfrak{P}$.

We then show that $\mathcal{P}^t \in \mathfrak{P}$ by recursion. We just showed that $\mathcal{P}^0 \in \mathfrak{P}$. Let $t \in \{0, \dots, m-1\}$ and assume that $\mathcal{P}^t \in \mathfrak{P}$. In many cases, $\mathcal{P}^{t+1} = \mathcal{P}^t$ and so $\mathcal{P}^{t+1} \in \mathfrak{P}$ by the recursion hypothesis. Otherwise, \mathcal{P}^{t+1} is given by the adjustment in line 17, in which case we have

$$\mathcal{P}^{t+1} = \left(\mathcal{P}^t \setminus \{k \in \mathcal{P}^t, R_k \subseteq R_{k^{(t+1, h_{t+1}^f)}}\} \right) \cup \{k^{(t+1, h_{t+1}^f)}\}. \quad (25)$$

Note that this imply that $i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$ (see lines 6 and 10).

Let $j \in \mathbb{N}_m^*$ and $k \in \mathcal{P}^t$ such that $j \in R_k$. If $R_k \subseteq R_{k^{(t+1, h_{t+1}^f)}}$, then $j \in R_{k^{(t+1, h_{t+1}^f)}} \subseteq \bigcup_{\kappa \in \mathcal{P}^{t+1}} R_\kappa$, and if not, $k \in \mathcal{P}^{t+1}$ and again $j \in \bigcup_{\kappa \in \mathcal{P}^{t+1}} R_\kappa$.

Now let $k, k' \in \mathcal{P}^{t+1}$, $k \neq k'$. If both are different from $k^{(t+1, h_{t+1}^f)}$, then they are both in $\mathcal{P}^t \in \mathfrak{P}$ so $R_k \cap R_{k'} = \emptyset$. Assume that $k' = k^{(t+1, h_{t+1}^f)}$. By the forest structure, $R_k \cap R_{k'} = \emptyset$, or $R_{k^{(t+1, h_{t+1}^f)}}$, or R_k . By (25), the latter is impossible. It remains to show that $R_k \cap R_{k'} = R_{k^{(t+1, h_{t+1}^f)}}$ is also not possible, in other words that we can't have $R_{k^{(t+1, h_{t+1}^f)}} \subsetneq R_k$. If that was the case, because of the strict inclusion, we would have $k \notin F$ because R_k could not be an atom, so we would have either $k \in E \subseteq \mathcal{K}_0^- \subseteq \mathcal{K}_t^-$ or $k \in \mathcal{P}^t \setminus \mathcal{P}^0$. In the second case, k would have been added to $\mathcal{P}^{t'}$ at a previous step $t' \leq t$ of the algorithm, but in that case it would also have been added to $\mathcal{K}_{t'}^- \subseteq \mathcal{K}_t^-$ (see lines 17 and 18). So in the end, in both cases, we would have

$$i_{t+1} \in R_{k^{(t+1, h_{t+1}^f)}} \subsetneq R_k \subseteq \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$$

which is a contradiction with the fact that $i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$, and so $R_k \cap R_{k'} = \emptyset$, and finally $\mathcal{P}^{t+1} \in \mathfrak{P}$.

7.2.3 Proof of (16) and (17)

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

So we let $t \in \{0, \dots, m-1\}$, and assume that $\mathcal{P}^{t'} \in \mathfrak{P}$ and that (16) and (17) hold for all $t' \leq t$. In all the following, \tilde{k} is the element of \mathcal{P}^t such that $i_{t+1} \in R_{\tilde{k}}$. We will distinguish two cases: if $i_{t+1} \in \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$ or not. First we show an inequality that will be used in both cases. We have, for all $k \in \mathcal{K}_t$,

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

Indeed, by (12),

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

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 $R_k \subsetneq R_{k'}$ 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 $R_k \subsetneq R_{k'}$ would entail $R_{\tilde{k}} \subsetneq R_{k'}$ 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 (26).

7.2.3.1 First case: $i_{t+1} \in \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$

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

$$\begin{aligned} \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| &= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\ &= V_{\mathfrak{R}}^*(S_t \cap R_k) \text{ by (17)} \\ &= \eta_k^t \text{ by (16)} \\ &= \eta_k^{t+1} \end{aligned}$$

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 $\zeta_{\bar{k}} \leq |S_t \cap R_{\bar{k}}|$, 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 (see line 17), or if not. Note that \bar{k} being added to \mathcal{P}^t during a previous step means that there exists $t', 1 \leq t' \leq t$, such that $h_{t'}^f$ is defined and $\bar{k} = k^{(t', h_{t'}^f)}$. The contrary means that for all $t', 1 \leq t' \leq t$ such that $h_{t'}^f$ is defined, $\bar{k} \neq k^{(t', h_{t'}^f)}$, and also that $\bar{k} \in \mathcal{P}^{t'}$ for all $t' \in \{0, \dots, t\}$.

7.2.3.1.1 First subcase: \bar{k} has never been added during the process of line 17

As remarked just above, $\bar{k} \in \mathcal{P}^0 = E \cup F$. Our goal is to show that $\bar{k} \notin F$. This will imply that $\bar{k} \in E \subseteq \mathcal{K}_0^-$ and so that $\zeta_{\bar{k}} = 0 \leq |S_t \cap R_{\bar{k}}|$ as desired.

Let us assume that $\bar{k} \in F$ and find a contradiction. \bar{k} is then the index of an atom that contains i_{t+1} , and $i_{t+1} \in \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$, so there exists $k' \in \mathcal{K}_t^-$ such that $i_{t+1} \in R_{k'}$ and then, by atomicity, $R_{\bar{k}} \subseteq R_{k'}$. By definition of F , k' cannot be in \mathcal{K}_0^- , so there exists $t' \in \{1, \dots, t\}$ such that k' has been added to $\mathcal{K}_{t'}^-$ by the process of line 18, and so, by lines 16 and 17, $k' = k^{(t', h_{t'}^f)}$ and $k' \in \mathcal{P}^{t'}$. But we also have $\bar{k} \in \mathcal{P}^{t'}$, and so by $\mathcal{P}^{t'}$ realizing a partition, the inclusion $R_{\bar{k}} \subseteq R_{k'}$ is actually an equality, so $\bar{k} = k' = k^{(t', h_{t'}^f)}$, which is not possible by assumption of this first subcase.

7.2.3.1.2 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 $\bar{k} = k^{(t', h_{t'}^f)}$ and that at that step $\eta_{\bar{k}}^{t'} \geq \zeta_{\bar{k}}$, because the if condition in line 13 failed. Also $\bar{k} \in \mathcal{P}^{t'}$ so $\bar{k} \in \mathcal{K}_{t'}$ so we can write

$$\begin{aligned} \zeta_{\bar{k}} &\leq \eta_{\bar{k}}^{t'} \\ &= V_{\mathfrak{R}}^*(S_{t'} \cap R_{\bar{k}}) \text{ by (16)} \\ &\leq |S_{t'} \cap R_{\bar{k}}| \\ &\leq |S_t \cap R_{\bar{k}}|. \end{aligned}$$

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 $R_{\bar{k}} \subseteq R_k$.

7.2.3.1.3 End of the first case

We write the following chain of claims:

$$\begin{aligned}
V_{\mathfrak{A}}^*(S_{t+1} \cap R_k) &\leq \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \text{ by (26) and } \mathcal{K}_{t+1} \subseteq \mathcal{K}_t \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_{t+1} \cap R_{\bar{k}}| \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge (|S_t \cap R_{\bar{k}}| + 1) \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_t \cap R_{\bar{k}}| \text{ because } \zeta_{\bar{k}} \leq |S_t \cap R_{\bar{k}}| \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| = \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| \\
&= V_{\mathfrak{A}}^*(S_t \cap R_k) \text{ by (17)} \\
&= \eta_k^t \text{ by (16)} \\
&= \eta_k^{t+1}.
\end{aligned}$$

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

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

which concludes this first case.

7.2.3.2 Second case: $i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$

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

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

we directly have

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

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 ,

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

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

$$\begin{aligned}
V_{\mathfrak{R}}^*(S_{t+1} \cap R_k) &\leq \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| \text{ by (26) and } \mathcal{K}_{t+1} \subseteq \mathcal{K}_t \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_{t+1} \cap R_{\bar{k}}| \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge (|S_t \cap R_{\bar{k}}| + 1) \\
&\leq \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k \\ k' \neq \bar{k}}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + \zeta_{\bar{k}} \wedge |S_t \cap R_{\bar{k}}| + 1 \\
&= \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_t \cap R_{k'}| + 1 \\
&= V_{\mathfrak{R}}^*(S_t \cap R_k) + 1 \text{ by (17)}.
\end{aligned} \tag{27}$$

Note that by the joint construction of \mathcal{K}_t^- and \mathcal{P}^t on lines 17 and 18, the fact that $i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$ implies that $\bar{k} \in F$, so \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 $k^{(t+1, h)}$, $1 \leq h \leq \phi(\bar{k})$. We now prove that for all of them, $V_{\mathfrak{R}}^*(S_{t+1} \cap R_k) \geq V_{\mathfrak{R}}^*(S_t \cap R_k) + 1$, 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 $S \subseteq \mathbb{N}_m$:

$$V_{\mathfrak{R}}^*(S) = \max_{\substack{A \subseteq \mathbb{N}_m \\ \forall k' \in \mathcal{K}, |A \cap R_{k'}| \leq \zeta_{k'}}} |A \cap S| = \max_{\substack{A \subseteq S \\ \forall k' \in \mathcal{K}, |A \cap R_{k'}| \leq \zeta_{k'}}} |A|. \tag{28}$$

By (28), 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 $V_{\mathfrak{R}}^*(S_t \cap R_{k^{(t+1, \phi(\bar{k})-1)}}) - V_{\mathfrak{R}}^*(S_t \cap R_{k^{(t+1, \phi(\bar{k})}}) = |B| - |A_{\phi(\bar{k})}|$ 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 (28).

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 $k' \in \mathcal{K}$

such that $R_{k^{(t+1, \phi(\bar{k})-1)}} \cap R_{k'} = \emptyset$, we have $|A_{\phi(\bar{k})-1} \cap R_{k'}| = 0 \leq \zeta_k'$. Furthermore,

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

by construction of $A_{\phi(\bar{k})}$. Finally, for all k' such that $R_{k^{(t+1, \phi(\bar{k})-1)}} \subseteq R_{k'}$, $|A_{\phi(\bar{k})-1} \cap R_{k'}| = |A_{\phi(\bar{k})-1}| = V_{\mathfrak{N}}^*(S_t \cap R_{k^{(t+1, \phi(\bar{k})-1)}}) = |B|$ 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 (28) 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 $A_{\phi(\bar{k})} \subseteq \dots \subseteq A_1$ such that for all $\ell \in \{1, \dots, \phi(\bar{k})\}$ and $k' \in \mathcal{K}$, $A_\ell \subseteq S_t \cap R_{k^{(t+1, \ell)}}$, $V_{\mathfrak{N}}^*(S_t \cap R_{k^{(t+1, \ell)}}) = |A_\ell|$, 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 $i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$ implies that $\eta_{k^{(t+1, \ell)}}^t < \zeta_{k^{(t+1, \ell)}}$ for all $\ell \in \{1, \dots, \phi(\bar{k})\}$. So by (16), $|A_\ell| < \zeta_{k^{(t+1, \ell)}}$.

Let, for any $\ell \in \{1, \dots, \phi(\bar{k})\}$, $\tilde{A}_\ell = A_\ell \cup \{i_{t+1}\}$. Given that $A_\ell \subseteq S_t \cap R_{k^{(t+1, \ell)}}$ and that $i_{t+1} \in S_{t+1} \setminus S_t$, $\tilde{A}_\ell \subseteq S_{t+1} \cap R_{k^{(t+1, \ell)}}$, $|\tilde{A}_\ell| = |A_\ell| + 1$, and for all $\ell' \in \{1, \dots, \phi(\bar{k})\}$, $|\tilde{A}_\ell \cap R_{k^{(t+1, \ell')}}| = |A_\ell \cap R_{k^{(t+1, \ell')}}| + 1$. Note that if, furthermore, $\ell \geq \ell'$, then $A_\ell \subseteq A_{\ell'}$, so

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

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

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

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

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

$$\begin{aligned} V_{\mathfrak{N}}^*(S_{t+1} \cap R_{k^{(t+1, \ell)}}) &\geq |\tilde{A}_\ell| \text{ by (28)} \\ &= |A_\ell| + 1 \\ &= V_{\mathfrak{N}}^*(S_t \cap R_{k^{(t+1, \ell)}}) + 1. \end{aligned}$$

So, as we wanted, $V_{\mathfrak{N}}^*(S_{t+1} \cap R_k) \geq V_{\mathfrak{N}}^*(S_t \cap R_k) + 1$ 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 (27) becomes an equality and we have proven that

$$V_{\mathfrak{N}}^*(S_{t+1} \cap R_k) = V_{\mathfrak{N}}^*(S_t \cap R_k) + 1 = \eta_k^t + 1 = \eta_k^{t+1},$$

that is, (16) is true at $t + 1$. Looking at the first line of (27), we also proved that

$$V_{\mathfrak{N}}^*(S_{t+1} \cap R_k) = \sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|. \quad (29)$$

The only thing left to prove is that (29) 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

$$\sum_{\substack{k' \in \mathcal{P}^t \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| = \sum_{\substack{k' \in \mathcal{P}^{t+1} \\ R_{k'} \subseteq R_k}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}|. \quad (30)$$

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

Now assume that h_{t+1}^f exists. So (25) holds. We will split each term in (30) in a sum of two terms. First, note that by (25), 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,

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

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

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

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.

Let us show that

$$\eta_{k^{(t+1, h_{t+1}^f)}}^{t+1} = \zeta_{k^{(t+1, h_{t+1}^f)}}. \quad (31)$$

Due to h_{t+1}^f existing, we broke the loop in line 19, so the condition in line 13 was false, so $\eta_{k^{(t+1, h_{t+1}^f)}}^{t+1} \geq \zeta_{k^{(t+1, h_{t+1}^f)}}$. We show by recursion over $t' \in \{0, \dots, t\}$ that $\eta_{k^{(t+1, h_{t+1}^f)}}^{t'} < \zeta_{k^{(t+1, h_{t+1}^f)}}$. Given that

$i_{t+1} \notin \bigcup_{\kappa \in \mathcal{K}_t^-} R_\kappa$ and $i_{t+1} \in R_{k^{(t+1, h_{t+1}^f)}}$, $k^{(t+1, h_{t+1}^f)} \notin \mathcal{K}_t^-$ and in particular $k^{(t+1, h_{t+1}^f)} \notin \mathcal{K}_0^-$ so $\eta_{k^{(t+1, h_{t+1}^f)}}^0 = 0 < \zeta_{k^{(t+1, h_{t+1}^f)}}$. Now let $t' < t$ and assume that $\eta_{k^{(t+1, h_{t+1}^f)}}^{t'} < \zeta_{k^{(t+1, h_{t+1}^f)}}$. We distinguish two cases. If $i_{t'+1} \notin R_{k^{(t+1, h_{t+1}^f)}}$, $k^{(t+1, h_{t+1}^f)}$ is not visited at step $t' + 1$ of the algorithm, so $\eta_{k^{(t+1, h_{t+1}^f)}}^{t'+1} = \eta_{k^{(t+1, h_{t+1}^f)}}^{t'} < \zeta_{k^{(t+1, h_{t+1}^f)}}$ by the recursion hypothesis. If $i_{t'+1} \in R_{k^{(t+1, h_{t+1}^f)}}$, all $k^{(t'+1, h)}$, $h \leq h_{t+1}^f$, are visited, and for all of them the condition in line 13 is true, otherwise we would have, for some $h \leq h_{t+1}^f$, $k^{(t'+1, h)} \in \mathcal{K}_{t'+1}^- \subseteq \mathcal{K}_t^-$. Noting that, necessarily, $k^{(t'+1, h)} = k^{(t+1, h)}$, we would finally have $i_{t+1} \in R_{k^{(t+1, h_{t+1}^f)}} \subseteq R_{k^{(t+1, h)}}$ which is a contradiction. This completes the recursion. Specifically for $t' = t$, $\eta_{k^{(t+1, h_{t+1}^f)}}^t \leq \zeta_{k^{(t+1, h_{t+1}^f)}} - 1$ and so $\eta_{k^{(t+1, h_{t+1}^f)}}^{t+1} \leq \eta_{k^{(t+1, h_{t+1}^f)}}^t + 1 \leq \zeta_{k^{(t+1, h_{t+1}^f)}}$ and so (31) holds.

Also note that $k^{(t+1, h_{t+1}^f)} \in \mathcal{P}^{t+1} \subseteq \mathcal{K}_{t+1}$, which implies two things. Firstly, because (16) holds at $t + 1$, $\eta_{k^{(t+1, h_{t+1}^f)}}^{t+1} = V_{\mathfrak{R}}^*(S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}})$. Secondly, by applying (29), we get that $\sum_{R_{k'} \subseteq R_{k^{(t+1, h_{t+1}^f)}}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| = V_{\mathfrak{R}}^*(S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}})$. Wrapping all these assertions:

$$\begin{aligned} \sum_{\substack{R_{k'} \subseteq R_{k^{(t+1, h_{t+1}^f)}} \\ k' \in \mathcal{P}^t}} \zeta_{k'} \wedge |S_{t+1} \cap R_{k'}| &= V_{\mathfrak{R}}^*(S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}}) \\ &= V_{\mathfrak{R}}^*(S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}}) \wedge |S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}}| \\ &= \eta_{k^{(t+1, h_{t+1}^f)}}^{t+1} \wedge |S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}}| \\ &= \zeta_{k^{(t+1, h_{t+1}^f)}} \wedge |S_{t+1} \cap R_{k^{(t+1, h_{t+1}^f)}}| \text{ by (31),} \end{aligned}$$

which achieves the second case and so the proof of Theorem 3.1.

7.3 Proof of Proposition 2.1

Recall that the R_k are assumed to be all non-empty and distinct by Remark 2.2.

The P_n , $n \in \mathbb{N}_N^*$, form a partition of \mathbb{N}_m^* , so $N \leq m$.

There is a sequence $R^{(H)} \subsetneq \dots \subsetneq R^{(1)}$. There exists $n_H \in \mathbb{N}_N^*$ such that $P_{n_H} \subseteq R^{(H)}$ and for $1 \leq h \leq H - 1$, there exists $n_h \in \mathbb{N}_N^*$ such that $P_{n_h} \subseteq R^{(h)} \setminus R^{(h+1)}$. For h_1, h_2 such that $1 \leq h_1 < h_2 \leq H$, we have $P_{n_{h_2}} \subseteq R^{(h_2)}$ and $P_{n_{h_1}} \subseteq R^{(h_1)} \setminus R^{(h_1+1)} \subseteq R^{(h_1)} \setminus R^{(h_2)}$ so $P_{n_{h_1}} \cap P_{n_{h_2}} = \emptyset$ and in particular $n_{h_1} \neq n_{h_2}$, so $h \mapsto n_h$ is an injection and $H \leq N$.

To show that the three bounds can be achieved simultaneously, we let $P_i = \{i\}$ for all $i \in \mathbb{N}_m^*$, $\mathcal{K} = \{(i, i), i \in \mathbb{N}_m^*\} \cup \{(1, i), i \in \mathbb{N}_m^*\}$, and, as usual, for $(i, j) \in \mathcal{K}$, $R_{(i, j)} = P_{i:j} = \bigcup_{i \leq n \leq j} P_n$.

Finally we show by induction over $N \geq 1$ that for any family $(R_k)_{k \in \mathcal{K}}$ with a forest structure and N leaves, $|\mathcal{K}| \leq 2N - 1$. For $N = 1$ it is trivial, necessarily $P_1 = \mathbb{N}_m^*$ and then the only possible set R_k is also \mathbb{N}_m^* so $|\mathcal{K}| = 1 = 2N - 1$. Let $N \geq 1$. Assume that for any family $(R_k)_{k \in \mathcal{K}}$ with a forest structure and N compatible leaves, $|\mathcal{K}| \leq 2N - 1$. Let $(R_k)_{k \in \mathcal{K}}$ a forest structure with $N + 1$ leaves, let H its maximum depth, let P_1, \dots, P_{N+1} the leaves. If $H = 1$, all the regions are two-by-two disjoint so there is an injection from the regions to the leaves and so $|\mathcal{K}| \leq N + 1 \leq 2(N + 1) - 1$. From now on we assume $H \geq 2$ and we distinguish two cases.

In the first case, assume that there exists \hat{k} of depth H , that is $\phi(\hat{k}) = H$, such that $R_{\hat{k}}$ is comprised of at least two leaves: there exist \tilde{i} and \tilde{j} with $\tilde{j} \geq \tilde{i} + 1$ such that $R_{\hat{k}} = \bigcup_{\tilde{i} \leq n \leq \tilde{j}} P_n$. Let $\mathcal{K}^- =$

$\mathcal{K} \setminus \{(\tilde{i}, \tilde{i}), (\tilde{i}+1, \tilde{i}+1)\}$, $(R_k)_{k \in \mathcal{K}^-}$ has also a forest structure, and we show that $P_1, \dots, P_{i-1}, P_i \cup P_{i+1}, P_{i+2}, \dots, P_{N+1}$ is a sequence of N leaves that are compatible with this family. First note that they well define a partition of \mathbb{N}_m^* . Let $k \in \mathcal{K}^-$, we just have to prove that if $P_i \subseteq R_k$ or $P_{i+1} \subseteq R_k$, then $P_i \cup P_{i+1} \subseteq R_k$. If that's the case, then $R_k \cap R_{\tilde{k}} \neq \emptyset$, and by the forest structure property of \mathcal{K} , $R_k \subsetneq R_{\tilde{k}}$ or $R_{\tilde{k}} \subseteq R_k$, but actually $R_k \subsetneq R_{\tilde{k}}$ is impossible because $\phi(\hat{k}) = H$ which is the maximal depth. So $R_{\tilde{k}} \subseteq R_k$, noticing that $P_i \cup P_{i+1} \subseteq R_{\tilde{k}}$, we get the desired result.

On the contrary, in the second case, assume that for all $k \in \mathcal{K}$ of height H , R_k is a leaf. Let $\hat{k} \in \mathcal{K}$ of depth H , that is $\phi(\hat{k}) = H$, and let $\tilde{k} \in \mathcal{K}$ the element of depth $H-1$ such that $R_{\hat{k}} \subsetneq R_{\tilde{k}}$. \tilde{k} exists because $H \geq 2$. Identify \tilde{k} to $(\tilde{i}, \tilde{j}) \in (\mathbb{N}_N^*)^2$ such that $R_{\tilde{k}} = \bigcup_{\tilde{i} \leq n \leq \tilde{j}} P_n$. If $\tilde{j} = \tilde{i}$, then $R_{\tilde{k}} = P_{\tilde{i}}$, then we have also $R_{\hat{k}} = P_{\tilde{i}}$, and there is a contradiction with the fact that $R_{\hat{k}} \subsetneq R_{\tilde{k}}$. So $\tilde{j} \geq \tilde{i}+1$. Let again $\mathcal{K}^- = \mathcal{K} \setminus \{(\tilde{i}, \tilde{i}), (\tilde{i}+1, \tilde{i}+1)\}$, and let us show again that $P_1, \dots, P_{i-1}, P_i \cup P_{i+1}, P_{i+2}, \dots, P_{N+1}$ is a sequence of N leaves that are compatible with $(R_k)_{k \in \mathcal{K}^-}$. The reasoning is the same as in the first case, but working with \tilde{k} instead of \hat{k} . Let $k \in \mathcal{K}^-$, such that $P_i \subseteq R_k$ or $P_{i+1} \subseteq R_k$. Then $R_k \cap R_{\tilde{k}} \neq \emptyset$, and by the forest structure property of \mathcal{K} , $R_k \subsetneq R_{\tilde{k}}$ or $R_{\tilde{k}} \subseteq R_k$. But actually $R_k \subsetneq R_{\tilde{k}}$ is impossible, because this implies that $\phi(k) = H$, so R_k is a leaf, so necessarily $R_k = P_i$ or $R_k = P_{i+1}$, but $k \in \mathcal{K}^-$ so $k \neq (\tilde{i}, \tilde{i})$ and $k \neq (\tilde{i}+1, \tilde{i}+1)$, hence a contradiction. So $R_{\tilde{k}} \subseteq R_k$, noticing that $P_i \cup P_{i+1} \subseteq R_{\tilde{k}}$, we get the desired result.

In both cases, we have constructed a forest structure $(R_k)_{k \in \mathcal{K}^-}$ with N compatible leaves. By the induction hypothesis, $|\mathcal{K}^-| \leq 2N-1$ and so $|\mathcal{K}| \leq |\mathcal{K}^-| + 2 \leq 2(N+1)-1$ which concludes.

A direct, alternative proof that $|\mathcal{K}| \leq 2m-1$ is given in next section.

7.3.1 Direct proof that $|\mathcal{K}| \leq 2m-1$

We show by induction on $m \geq 1$ that, for a family of subsets $(R_k)_{k \in \mathcal{K}}$ with a forest structure such that the R_k are all non-empty and distinct, $|\mathcal{K}| \leq 2m-1$. For $m=1$ it is trivial, the only subset possible is $\{1\}$. Now let $m \geq 1$ and assume that the result is true for m . Let $(R_k)_{k \in \mathcal{K}}$ a family of non-empty and distinct subsets of \mathbb{N}_{m+1}^* with a forest structure.

Let k_1, \dots, k_D , $D \leq H$, the indices of the regions including $m+1$ (possibly non-existent, in which case $D=0$), ordered such that $R_{k_1} \subsetneq \dots \subsetneq R_{k_D}$. Let $\tilde{\mathcal{K}} = \mathcal{K} \setminus \{k_1, \dots, k_D\}$, and let $\mathcal{K}' = \mathcal{K} \setminus \{k_1, k_2\} = \tilde{\mathcal{K}} \cup \{k_3, \dots, k_D\}$. For $k \in \tilde{\mathcal{K}}$, we let $R'_k = R_k$, and for $k \in \{k_3, \dots, k_D\}$, we let $R'_k = R_k \setminus \{m+1\}$. The rest of the proof consists in proving that $(R'_k)_{k \in \mathcal{K}'}$ is a family of non-empty and distinct subsets of \mathbb{N}_m^* with a forest structure. Once this is proven, by induction hypothesis we will have $|\mathcal{K}'| \leq 2m-1$, and finally $|\mathcal{K}| \leq |\mathcal{K}'| + 2 \leq 2m-1+2 = 2(m+1)-1$.

First, any R'_k , $k \in \mathcal{K}'$, is non-empty, because if $k \in \tilde{\mathcal{K}}$, $R'_k = R_k \neq \emptyset$, and if $k = k_d$ with $d \geq 3$, $R_{k_1} \subsetneq R_{k_2} \subsetneq R_k$ so $|R_k| \geq 3$ and then $|R'_k| = |R_k \setminus \{m+1\}| \geq 2$.

To prove that $(R'_k)_{k \in \mathcal{K}'}$ is a family of distinct subsets of \mathbb{N}_m^* with a forest structure, we need to take $k, k' \in \mathcal{K}'$, $k \neq k'$, and show that $R'_k \neq R'_{k'}$ and $R'_k \cap R'_{k'} \in \{\emptyset, R'_k, R'_{k'}\}$.

If $|\tilde{\mathcal{K}}| \geq 2$, let $k, k' \in \tilde{\mathcal{K}}$, $k \neq k'$. We have $R'_k = R_k$ and $R'_{k'} = R_{k'}$, so $R'_k \neq R'_{k'}$ and $R'_k \cap R'_{k'} \in \{\emptyset, R_k, R_{k'}\} = \{\emptyset, R'_k, R'_{k'}\}$.

If $D \geq 4$, let $i, j \in \{3, \dots, D\}$, $i < j$. We have $R'_{k_i} = R_{k_i} \setminus \{m+1\}$ and $R'_{k_j} = R_{k_j} \setminus \{m+1\}$ with $R_{k_i} \subsetneq R_{k_j}$, so $R'_{k_i} \neq R'_{k_j}$ and $R'_{k_i} \cap R'_{k_j} = R_{k_i} \setminus \{m+1\} = R'_{k_i}$.

If $D \geq 3$ and $|\tilde{\mathcal{K}}| \geq 1$, let $i \in \{3, \dots, D\}$ and $k \in \tilde{\mathcal{K}}$. We have $R'_{k_i} = R_{k_i} \setminus \{m+1\}$ and $R'_k = R_k$.

$$\begin{aligned} R'_{k_i} \cap R'_k &= (R_{k_i} \setminus \{m+1\}) \cap R_k \\ &= R_{k_i} \cap R_k \text{ because } m+1 \notin R_k \\ &\in \{\emptyset, R_{k_i}, R_k\} \text{ by the property of forest structure} \end{aligned}$$

Given that $R'_{k_i} \subsetneq R_{k_i}$, $R'_{k_i} \cap R'_k = R_{k_i}$ is impossible, so $R'_{k_i} \cap R'_k \in \{\emptyset, R_k\} = \{\emptyset, R'_k\}$ so the only thing remaining to prove is that R'_{k_i} and R'_k are distinct. We prove that by showing that if $R'_k = R'_{k_i}$, there is a contradiction. Indeed, then $R_k = R_{k_i} \setminus \{m+1\}$, and we can study $R_{k_2} \cap R_k$. On the one hand, $R_{k_2} \cap R_k \in \{\emptyset, R_{k_2}, R_k\} = \{\emptyset, R_{k_2}, R_{k_i} \setminus \{m+1\}\}$ by forest structure. On the other hand,

$$\begin{aligned} R_{k_2} \cap R_k &= (R_{k_2} \setminus \{m+1\}) \cap R_k \text{ because } m+1 \notin R_k \\ &= (R_{k_2} \setminus \{m+1\}) \cap (R_{k_i} \setminus \{m+1\}) \\ &= R_{k_2} \setminus \{m+1\}. \end{aligned}$$

So $R_{k_2} \cap R_k = R_{k_2}$ is impossible. Furthermore, $R_{k_2} \cap R_k = R_k$ is also impossible because $R_{k_2} \subsetneq R_{k_i}$ and $m+1 \in R_{k_2}$ hence $R_{k_2} \setminus \{m+1\} \subsetneq R_{k_i} \setminus \{m+1\}$. So $R_{k_2} \cap R_k = \emptyset$, that is $R_{k_2} \setminus \{m+1\} = \emptyset$, so $R_{k_2} = \{m+1\}$ and the contradiction is the following: $\{m+1\} \subseteq R_{k_1} \subsetneq R_{k_2} = \{m+1\}$.

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Session information

R version 4.5.1 (2025-06-13)

Platform: x86_64-pc-linux-gnu

Running under: Ubuntu 24.04.3 LTS

Matrix products: default

BLAS: /usr/lib/x86_64-linux-gnu/blas/libblas.so.3.12.0

LAPACK: /usr/lib/x86_64-linux-gnu/lapack/liblapack.so.3.12.0 LAPACK version 3.12.0

locale:

[1] LC_CTYPE=C.UTF-8	LC_NUMERIC=C	LC_TIME=C.UTF-8
[4] LC_COLLATE=C.UTF-8	LC_MONETARY=C.UTF-8	LC_MESSAGES=C.UTF-8
[7] LC_PAPER=C.UTF-8	LC_NAME=C	LC_ADDRESS=C
[10] LC_TELEPHONE=C	LC_MEASUREMENT=C.UTF-8	LC_IDENTIFICATION=C

time zone: Etc/UTC

tzcode source: system (glibc)

attached base packages:

[1] stats graphics grDevices datasets utils methods base

other attached packages:

[1] sanssouci_0.16.2 microbenchmark_1.5.0

loaded via a namespace (and not attached):

[1] digest_0.6.37	fastmap_1.2.0	xfun_0.53	Matrix_1.7-3
[5] lattice_0.22-5	matrixStats_1.5.0	knitr_1.50	htmltools_0.5.8.1
[9] generics_0.1.4	rmarkdown_2.29	cli_3.6.5	grid_4.5.1
[13] renv_1.1.5	matrixTests_0.2.3	compiler_4.5.1	tools_4.5.1
[17] evaluate_1.0.4	yaml_2.3.10	rlang_1.1.6	jsonlite_2.0.0