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LAD MODELS, TREES AND AN ANALOG OF THE FUNDAMENTAL THEOREM OF ARITHMETIC

NADIA BRAUNER, SYLVAIN GRAVIER, LOUIS-PHILIPPE KRONEK, AND FRÉDÉRIC MEUNIER

ABSTRACT. Motivated by applications of Logical Analysis Data (LAD) in medical contexts, original discrete optimization problems are proposed. When a patient arrives with a presumption of a disease, he is submitted to a sequence of tests. From one patient to another, the tests allowing to detect the disease may vary. A subset of tests whose results detect the disease in a given part of the population is called a *pattern*, which has its own prevalence in the population.

If there is only a limited number of tests that can be done, which ones must be selected in order to maximize the number of spotted patients? Or, if each test has a cost, in which order the tests have to be done, in order to minimize the cost? It is the kind of questions that are investigated in this paper. For various special cases, polynomial algorithms are proposed, especially when the hypergraph whose vertices are the tests and whose edges are the patterns is a tree graph.

One of these questions involves a criterion which is not a number but a sequence of numbers. The objective is then to find the best sequence for the lexicographic order. To solve this question, a new product on finite sequences is defined, namely the *maximum shuffle product*, which maps two sequences to their shuffle that is maximal for the lexicographic order. Surprisingly, this product leads to a theorem similar to the fundamental theorem of arithmetic: every sequence can be written uniquely as the product of *prime sequences*, with the suitable definition of prime sequences.

1. Introduction

1.1. **Context.** Discrete mathematics and medicine see their interactions growing these last years. This is particularly true for the area of classification models, which help practitioners to make diagnosis on the basis of patterns (rules). Formalizing questions arising when one implements a set of patterns (a model), we get nice combinatorial optimization problems.

Logical Analysis of Data (LAD) aims at helping practitioners for decision making when a new patient arrives with a presumption of a disease. It is a rule based method where a classification model is composed of a set of patterns [5]. Assume that we are given possible tests (e.g. age, headache...). From one patient to another, the tests allowing to confirm the disease may vary. A pattern is a boolean conjunction on the results of tests. If a patient can be classified by these results, a positive (resp. negative) class meaning having (resp. not having) the disease, we say that the pattern covers the patient. We shall consider an approximation of the global problem with the following hypothesis: every patient is covered by exactly one pattern.

The problems we are interested in concern for instance the ordering of the tests allowing to classify a new patient as soon as possible, or the selection of a subset of tests if all tests can not be done. To estimate the quality of an ordering, or of a subset, of the tests, a database is used. It is composed of already classified patients, for whom there were a presumption of the disease. The prevalences of the patterns in the database are supposed to be good approximations of the pattern prevalences among the future patients, who will have to be classified. Therefore, when a solution (an ordering, a subset, ...) is proposed, its quality is measured by its results on the database.

A pattern is identified with the subset of tests whose results it takes into account. We will often work with the pattern-test hypergraph H = (V, E), having the tests as vertex set and the patterns

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as edge set. The prevalence of a pattern $e \in E$ is the number p(e) of patients of the database covered by this pattern.

Table 1 presents a LAD model where the patterns are based on 4 tests. The last column gives the number of patients covered by the pattern in the database. In terms of the pattern-test hypergraph, $V = \{1, 2, 3, 4\}$ and $E = \{1, 23, 13, 14\}$, with p(1) = 18, p(23) = 10, p(13) = 70 and p(14) = 38.

| Pattern | Test 1 | Test 2 | Test 3 | Test 4 | class | cover |
|------------------|--------|----------|----------|----------|----------|-------|
| $\overline{P_1}$ | > 30 | | | | positive | 18 |
| $\overline{P_2}$ | | negative | positive | | positive | 10 |
| P_3 | < 15 | | negative | | negative | 70 |
| P_4 | < 15 | | | positive | negative | 38 |

Table 1. A LAD model

Note that when the tests are done one after the other, each time a complete pattern is contained in the tests already done, all the patients in the database covered by this pattern can be classified. If each test v has a specific cost c(v), tests with high costs have to be at the end of the process, since we can then hope that most of the patients have already been spotted. It suggests the following problem.

Problem Testing Cost

INSTANCE: A hypergraph H=(V,E) with n vertices, two maps $p:E\to\mathbb{R}_+$ and $c:V\to\mathbb{R}_+$

SOLUTION: An ordering v_1, \ldots, v_n of the vertices such that

$$\sum_{i=1}^{n} \sum_{e \in E \setminus E[v_1, \dots, v_i]} p(e)c(v_i)$$

is minimum.

This problem was initiated when working on the diagnosis of growth hormone deficiency [9]. During the diagnosis process many practitioners are involved successively (general practitioners, biologists, endocrinologists, radiologists) and the tests (height, bone age, karyotype...) have different performances and costs.

Remark. In this model, any patient whose status has not been settled will be subject to the tests. However, in many cases, the first results of the tests will be sufficient for some patients to predict the result of the next test, and hence that this next test is not necessary for these patients. The modelling of such features deserves future works.

Finding the k tests covering the most patients can be formalized as follows.

Problem Extraction of a sub-model

INSTANCE: A hypergraph H = (V, E) with n vertices, a map $p: E \to \mathbb{R}_+$ and an integer k.

SOLUTION: A subset $S \subseteq V$ of cardinality at most k maximizing p(E[S]).

This problem appeared when working on the diagnosis of pulmonary embolism. The medical question was to identify the first tests to be implemented at the emergency room in order to be able to make rapidly the greatest number of diagnosis for this pathology.

We may want to give a complete ordering v_1, \ldots, v_n of the tests in such a way that if for a reason we do not want to do all tests and we stop them say at v_k , we do not regret our choice of an ordering. We write \leq the lexicographic order on sequences.

Problem Non-dominated ordering

INSTANCE: A hypergraph H = (V, E) with n vertices and a map $p: E \to \mathbb{R}_+$

SOLUTION: An ordering v_1, \ldots, v_n of the vertices such that the sequence h_1, \ldots, h_n , with $h_i = p(E[\{v_1, \ldots, v_i\}])$, is maximal for the lexicographic order.

It is also interesting to compare two sequences with respect to the Pareto-order instead of lexi-cographic order.

A Pareto-optimal ordering v_1, \ldots, v_n is such that for all other ordering v'_1, \ldots, v'_n , there is a $i \in \{1, \ldots, n\}$ with $h'_i < h_i$. A sequence maximal for the lexicographic order is Pareto-optimal, but the converse is not true.

Problem PARETO-OPTIMAL ORDERING

INSTANCE: A hypergraph H = (V, E) with n vertices and a map $p: E \to \mathbb{R}_+$

SOLUTION: An ordering v_1, \ldots, v_n of the vertices such that the sequence h_1, \ldots, h_n , with $h_i = p(E[\{v_1, \ldots, v_i\}])$, is Pareto-optimal.

1.2. Main results. Testing costs, Extraction of a sub-model, and Non-dominated ordering are checked to be NP-hard, even if the hypergraph H is a graph in the case of the two laters, but the complexity of Pareto-optimal ordering is open.

However, we prove

Proposition 1.1. Pareto-optimal ordering is polynomial (in $O(n^2 \log n)$) when the hypergraph is a tree graph.

Theorem 1.2. Non-dominated ordering is polynomial (in $O(n^3 \log n \log d)$) when the hypergraph is a tree graph and d its maximal degree.

The polynomiality stated in Proposition 1.1 is a corollary of Theorem 1.2, since as we have already noted, an ordering of the tests that is maximal for the lexicographic order is necessarily Pareto-optimal. Proposition 1.1 finds its interest in a better complexity $(O(n^2 \log n))$ versus $O(n^3 \log n \log d)$.

The proof of Theorem 1.2 motivates the definition of shuffles of sequences.

A sequence $\mathbf{c} = c_1 \dots c_n$ is a shuffle of s sequences $\mathbf{a}_i = a_{i1} \dots a_{in_i}, i = 1, \dots, s$ if

- $n = \sum_{i=1}^{s} n_i$.
- there are s strictly increasing functions $\alpha_i : [n_i] \to [n]$ (the prints) whose images are pairwise disjoint and such that $c_{\alpha_i(j)} = a_{ij}$ for all i, j.

Denote by ms(a, b) the shuffle of a and b that is maximal for the lexicographic order \leq . We discuss the way for computing ms(a, b) given two sequences a and b. Theorem 1.2 is a consequence of the polynomiality of this operation.

On our track, we discovered a combinatorial analog of the fundamental theorem of arithmetic, which states the uniqueness of decomposition of an integer as product of prime numbers. To our knowledge, this combinatorial analog is completely new. There exist other decompositions theorems which generalize the classical one (see [11] for instance), but they do not contain it.

We define a notion of prime sequences. A prime sequence $\mathbf{p} = p_1 \dots p_n$ is such that $p_i p_{i+1} \dots p_n \succeq \mathbf{p}$ for each $i \geq 1$.

Theorem 1.3. Each sequence **a** can be written as a unique maximum shuffle (up to permutation)

$$oldsymbol{a} = \mathrm{ms}\left(oldsymbol{q}_1^{\delta_1}, oldsymbol{q}_2^{\delta_2}, \ldots, oldsymbol{q}_s^{\delta_s}
ight)$$

where the δ_j are positive integers and the q_j are prime.

Here,
$$q^{\delta}$$
 denotes $\operatorname{ms}(\underbrace{q,\ldots,q}_{\delta \text{ times}})$.

1.3. Notations.

- 1.3.1. Hypergraphs. A hypergraph is denoted H = (V, E). Again, the number n is usually its number of vertices. Two vertices u and v are neighbors is there is an edge $e \in E$ such that $\{u, v\} \subseteq e$. By N(A), for A a subset of vertices, we denote the set of vertices in $V \setminus A$ having at least one neighbor in A. By E[X], for a subset $X \subseteq V$ of vertices, we mean the subset of edges having all their vertices in X.
- 1.3.2. Combinatorics. Let A be an additive semi-group and w be any map from a finite set X into A. For F a subset of X, we denote $\sum_{x \in F} w(x)$ by w(F).
- 1.3.3. Scheduling. We use the classical $\alpha |\beta| \gamma$ notation in scheduling where α denotes the properties of the machines (or the processors), β indicates the constraints on the tasks and γ describes the objective function. In this paper, α will be equal to 1 since we consider one machine problems. The field β contains information on the precedence graph for the tasks (where 'prec' means that the precedence graph can be any acyclic graph). The objective in the field γ is $\sum w_i C_i$ if we want to minimize the total weighed completion time where w_i is the weight of task i and C_i is its completion time. It is $\sum w_i U_i$ when we want to minimize the weighted number of late jobs (where $U_i = 1$ if and only if task i completes after its dead line).
- 1.4. **Plan.** Notations that have not been defined in the introduction are fixed in Section 1.3. We have then four sections (Sections 2, 3, 4 and 5), one for each of these problems, which contain among other things the proofs of Proposition 1.1 and of Theorem 1.2. Section 6 is devoted to the proof of the analog of the fundamental theorem of arithmetic for sequences and shuffles (Theorem 1.3).

2. Testing Cost

We show that the problem Testing Cost is equivalent to a scheduling problem on one machine with a specific precedence graph and the total weighted completion time as the objective function: $1|prec|\sum w_iC_i$ in the usual scheduling notations.

We now define the tasks and the precedence graph.

- To each vertex v in the hypergraph is associated a (vertex-)task of duration $t_v = c(v)$ and of weight $w_v = 0$;
- To each edge e in the hypergraph is associated a (edge-)task of duration $t_e = 0$ and of weight $w_e = p(e)$;

The precedence graph naturally follows: a edge-task can be executed once all the vertex-tasks coming from vertices in the edge are completed. Figure 1 presents the scheduling problem associated to the model from Table 1.

| | Test 1 | Test 2 | Test 3 | Test 4 |
|-------|--------|--------|--------|--------|
| Costs | 1 | 5 | 15 | 8 |

Table 2. Costs for the test of Table 1

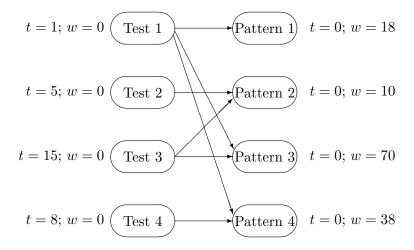


FIGURE 1. The precedence graph associated to the model of Table 1 and the costs of Table 2

Proposition 2.1. Problem Testing Cost is equivalent to the scheduling problem $1|prec| \sum w_i C_i$ with the tasks and the precedence graph defined as above.

Proof. Given a solution of the problem TESTING COST, we get a feasible solution for problem $1|prec|\sum w_iC_i$ of same cost while starting a edge-task as soon as all vertex-tasks it requires are finished. Conversely, an optimal solution of problem $1|prec|\sum w_iC_i$ is such that a edge-task starts as soon as all vertex-task it requires are finished and the sequence of vertices the solution induces is a solution of same cost for TESTING COST.

The precedence graph has the following property: if there is a precedence constraint between tasks i and j of the form $i \to j$, then $w_i = 0$ and $t_j = 0$. This type of graphs is known as red-blue bipartite graphs [7].

The scheduling problem $1|prec|\sum w_iC_i$ is NP-hard for a general precedence graph [8]. We consider the special case of red-blue bipartite graphs for the precedence constraints with the restriction $p_i \in \{0,1\}$ and $w_i \in \{0,1\}$. This theoretical situations corresponds to the very special case where each pattern covers exactly one patient and each test has a unitary cost. It has been proved in [12] that this special case has the same approximation ratio that the general problem. Testing cost is therefore NP-hard.

3. Extraction of a sub-model

- 3.1. The densest subgraph. The special case of the problem EXTRACTION OF A SUB-MODEL when H is a graph has been widely studied and is better known under the name DENSEST k-SUBGRAPH PROBLEM. It is an NP-hard problem since when p(e) = 1 for all e it becomes the problem of selecting the subset X of vertices with |X| = k such that E[X] has maximal cardinality, which contains the maximum clique problem. It is known to be polynomial on trees, with a complexity $O(k^2n)$ [10].
- 3.2. **As a scheduling problem.** The problem EXTRACTION OF A SUB-MODEL can also be linked to a scheduling problem similar to the one of Section 2. There is still one machine and the precedence graph is the same. The tasks are defined as follows:
 - To each vertex v is associated a (vertex-)task of duration $t_v = 1$ and of weight $w_v = 0$ with no deadline $d_v = +\infty$;

• To each edge e is associated a (edge-)task of duration $t_e = 0$ and of weight $w_e = p(e)$ and of deadline $d_e = k$.

The objective function is now the minimization of the number of weighted tasks that finish late, $\sum w_i U_i$ where U_i indicates whether task *i* finishes after its deadline. The edge-tasks that are not late compose the submodel S.

This problem has been widely studied under different forms, see e.g. [2]. For instance, it is equivalent to the tool magazine problem where the tests are the tools, the patterns are the parts to be produced and the prevalence is the demand for the parts. The part-tools matrix indicating for each part the tools needed for its production is then the incidence matrix of the hypergraph H. The value k defines the capacity of the tool magazine and the objective is to select k tools for the magazine that allow the satisfaction of the largest demand. This problem of the tool magazine is known to be NP-hard even if the number of tools for each part is less than 2 [4]. Crama [2] has given linear and non-linear formulations of the problem, proposed solutions methods and listed other applications of this model like repair kit selection or allocation of memory space in a database. It is mentioned in this paper that, those practical problems are solved with ad-hoc heuristics. In [3], the authors present a worst case analysis of the greedy heuristics usually developed for this problem. They also conjecture that there is no polynomial time algorithm with a constant performance ratio.

4. Non-dominated ordering

4.1. First properties. Defining p(e) to be 1 for all $e \in E$, a maximal ordering for the lexicographic order necessarily starts with the vertices of a maximum clique. We have therefore the following complexity result.

Proposition 4.1. Non-dominated ordering is NP-hard even if the hypergraph H is a graph.

As a preliminary remark on the problem, we can notice that

Lemma 4.2. Let v_1, \ldots, v_n be a solution of Non-dominated ordering. For all i < n, if $N(v_1, \ldots, v_i) \neq \emptyset$, then $v_{i+1} \in N(v_1, \ldots, v_i)$.

Proof. Assume that we have i such that $N(v_1, \ldots, v_i) \neq \emptyset$. Choosing v_{i+1} not in $N(v_1, \ldots, v_i)$ leads to $h_{i+1} = h_i$ whereas choosing it in $N(v_1, \ldots, v_i)$ leads $h_{i+1} > h_i$, without changing the values of h_i for i < i.

4.2. A greedy algorithm. We propose now a greedy approach.

At each step, add the minimal number of vertices such that the induced subhypergraph contains a new edge. If several new edges can be obtained in this way, select the one with maximal value p. At the end of each iteration, the hypergraph is updated by merging the edges with the same sets of missing vertices. The new p is the sum of the values of p of the merged edges.

```
Greedy algorithm (a hypergraph H)

// Create an order with a greedy method

Ordering \leftarrow \emptyset and ActiveEdges \leftarrow \emptyset

While Ordering does not contain all the vertices do

ActiveEdges \leftarrow \text{edges from hypergraph } H that have the smallest number of vertices not in Ordering

Add to Ordering the vertices completing an edge in ActiveEdges with the maximal p.

Merge the edges with identical sets of vertices not in Ordering and update the values of p.

end While

Return Ordering
```

Unfortunately, this greedy algorithm does not always lead to an ordering maximal for the lexicographic order ≤. Consider, for example, the model presented in Table 3. The order Test 1, Test 2, Test 3, Test 4, Test 5 can be obtained by the greedy algorithm. It is dominated by the order Test 4, Test 3, Test 5, Test 2, and Test 1. The following proposition gives a sufficient condition for the order produced by the greedy algorithm to be non-dominated.

| Patterns | Test 1 | Test 2 | Test 3 | Test 4 | Test 5 | Cover |
|------------------|--------|--------|--------|--------|--------|-------|
| $\overline{P_1}$ | 1 | 1 | 0 | 0 | 0 | 1 |
| P_2 | 0 | 0 | 1 | 1 | 0 | 1 |
| P_3 | 0 | 0 | 0 | 1 | 1 | 1 |

Table 3. A model for which the greedy algorithm does not give a non-dominated order

Proposition 4.3. If, at each iteration, there is only one active pattern of maximum prevalence, then the result of the greedy algorithm is an optimal solution of NON-DOMINATED ORDERING.

Proof. The proof is achieved by induction on the iterations of the algorithm. At each step, we can not do better than selecting the tests of this active pattern. \Box

The sufficient condition presented in Proposition 4.3 is realistic in practice. The probability is low that there exists at some time in the execution of the algorithm, two sets of distinct patterns that have the same prevalence and that need the realization of a minimal number of tests. Indeed, generically, the hyperplane $\sum_{e \in E} p(e)x_e = 0$ does not contain the vertices of the hypercube $[-1,1]^n$. An interesting question could be to characterize the models that verify the condition of Proposition 4.3.

4.3. **Tree.** The purpose of this subsection is to prove Theorem 1.2. As a by-product, we get some properties of the *shuffle* and the ms operation (defined in the introduction). In Section 6, it will lead to the analog of the fundamental theorem in arithmetic we have mentioned.

4.3.1. Preliminary discussion.

Lemma 4.4. Let $a_1 \ldots a_n$ and $b_1 \ldots b_n$ be two sequences of real numbers (of same length). $a_1 \ldots a_n \leq b_1 \ldots b_n$ if and only if $a_1(a_2 - a_1)(a_3 - a_2) \ldots (a_n - a_{n-1}) \leq b_1(b_2 - b_1)(b_3 - b_2) \ldots (b_n - b_{n-1})$.

Proof. Let
$$i^*$$
 be the first index i such that $b_i \neq a_i$. We have $b_{i^*} > a_{i^*}$ if and only if $b_{i^*} - b_{i^*-1} > a_{i^*-1}$.

Lemma 4.4 combined with Lemma 4.2 shows that finding a maximal ordering for the lexicographic order when the hypergraph H is a tree T reduces to the following problem. Given r a special vertex in T, and a weight function $w: V(T) \to \mathbb{R}_+ \setminus \{0\}$, compute an ordering $r = v_1, \ldots, v_n$ of the vertices such that $v_{i+1} \in N(v_1, \ldots, v_i)$ for all i and such that $w(v_2), \ldots, w(v_n)$ is maximal for the lexicographic order. Doing the same computation for each vertex r of T, and keeping the maximal one gives the solution of the original problem.

Note that the question of finding such an ordering starting from vertex r is easy if all weights in the tree are distinct. Indeed, at each time, selecting the edge with maximal weight among the edges leaving the set of vertices already reached clearly provides the maximal ordering (Proposition 4.3). The difficulty arises precisely when there are many weights that are equal. Choosing then an edge with maximal weight could eventually lead to lower weight later that those that would have been obtained by choosing another edge of the same maximal weight.

Deleting r from T gives raise to say s subtrees T_1, T_2, \ldots, T_s . As we will prove below, once we have a maximal ordering of the vertices for each T_i , it is quite easy to "mix" these orderings to get

the optimal one for T itself. This motivates the following study of sequences and the definition of the shuffle.

4.3.2. Sequences, lexicographic order and shuffles. For $A \subseteq \{1, 2, ..., n\}$, we denote by $\boldsymbol{c}|_A$ the subsequence $(c_i)_{i \in A}$. Given two sequences $\boldsymbol{a} = a_1 ... a_m$ and $\boldsymbol{b} = b_1 ... b_n$, we denote by $\boldsymbol{a} \bullet \boldsymbol{b}$ the sequence $a_1 ... a_m b_1 ... b_n$. The empty sequence is denoted $\boldsymbol{\epsilon}$.

We have the following proposition, which seems to be interesting for its own sake.

Proposition 4.5. Given r sequences \mathbf{a}_i , i = 1, ..., r, of real numbers, it is possible to compute $\operatorname{ms}(\mathbf{a}_1, ..., \mathbf{a}_r)$ in $O(n^2 \log r)$, where n is the sum of the lengths of the \mathbf{a}_i .

Before proving it, we will study some properties of ms. We endow the set of sequences with the lexicographic order \leq .

We will use the properties stated in the following lemma without explicit mention. Indeed, there are all more or less obvious.

Lemma 4.6. We have the following properties.

(1)

$$ms(\boldsymbol{a}, \boldsymbol{b}) = ms(\boldsymbol{b}, \boldsymbol{a}).$$

- (2) For a fixed sequence a, the map $x \mapsto ms(a, x)$ is an increasing map.
- (3) $\operatorname{ms}(\boldsymbol{a}, \boldsymbol{b}) \bullet \operatorname{ms}(\boldsymbol{c}, \boldsymbol{d}) \preceq \operatorname{ms}(\boldsymbol{a} \bullet \boldsymbol{c}, \boldsymbol{b} \bullet \boldsymbol{d}).$
- (4) if $\mathbf{a} \leq \mathbf{b}$ then for any sequence \mathbf{c} we have

$$c \bullet a \prec c \bullet b$$
.

(5)

$$ms(\boldsymbol{a}, ms(\boldsymbol{b}, \boldsymbol{c})) = ms(\boldsymbol{a}, \boldsymbol{b}, \boldsymbol{c}).$$

Proof. Except maybe Points (3) and (5), the other points are straightforward.

Point (3): it is a consequence of the following fact: a shuffle of a and b followed by a shuffle of c and d is a shuffle of $a \cdot c$ and $b \cdot d$.

Point (5): We obviously have

$$ms(\boldsymbol{a}, ms(\boldsymbol{b}, \boldsymbol{c})) \leq ms(\boldsymbol{a}, \boldsymbol{b}, \boldsymbol{c}).$$

Hence, we want to prove the reverse inequality. Let d = ms(a, b, c) and s = ms(b, c). Define β and γ to be respectively the prints of b and c in d. Denote B the image of β and C the image of γ . Then $d|_{B \cup C} \leq s$. Since, $d = ms(a, d|_{B \cup C})$, we conclude with the help of Point (2).

Lemma 4.6 shows that the set of sequences endowed with ms as a product is a monoid.

Lemma 4.7. Let a and b be two sequences such that $a \leq b$. Let c be another sequence. Then

$$ms(a, c \bullet b) \leq ms(c \bullet a, b).$$

Proof. Write $\mathbf{a} = a_1 \dots a_m$, $\mathbf{b} = b_1 \dots b_n$ and $\mathbf{c} = c_1 \dots c_r$.

The proof works by induction on m + n. If m + n = 1, the statement is straightforward.

Let $\mathbf{a}' = a_1 \dots a_k$ and $\mathbf{a}'' = a_{k+1} \dots a_m$ be two sequences such that

$$ms(\boldsymbol{a}, \boldsymbol{c} \bullet \boldsymbol{b}) = ms(\boldsymbol{a}', \boldsymbol{c}) \bullet ms(\boldsymbol{a}'', \boldsymbol{b}).$$

If $a'' = \epsilon$, we define k to be m.

Now, define $b' := b_1 \dots b_{\min(k,n)}$ and b'' such that $b = b' \bullet b''$. If k or n equals 0, we define $b' := \epsilon$. If $b' \succ a'$, then we have

$$\operatorname{ms}(\boldsymbol{a}',\boldsymbol{c}) \bullet \operatorname{ms}(\boldsymbol{a}'',\boldsymbol{b}) \preceq \operatorname{ms}(\boldsymbol{b}',\boldsymbol{c}) \preceq \operatorname{ms}(\boldsymbol{c},\boldsymbol{b}') \bullet \operatorname{ms}(\boldsymbol{a},\boldsymbol{b}'') \preceq \operatorname{ms}(\boldsymbol{c} \bullet \boldsymbol{a},\boldsymbol{b}).$$

Hence, we can assume that b' = a'. Thus, $b'' \succeq a''$. By induction

$$ms(b' \bullet a'', b'') \succeq ms(a'', b' \bullet b'').$$

Using the equality b' = a' and Point (3) of Lemma 4.6, we get finally

$$\begin{array}{rcl} \operatorname{ms}(\boldsymbol{a},\boldsymbol{c} \bullet \boldsymbol{b}) & = & \operatorname{ms}(\boldsymbol{a}',\boldsymbol{c}) \bullet \operatorname{ms}(\boldsymbol{a}'',\boldsymbol{b}' \bullet \boldsymbol{b}'') \\ & \leq & \operatorname{ms}(\boldsymbol{b}',\boldsymbol{c}) \bullet \operatorname{ms}(\boldsymbol{b}' \bullet \boldsymbol{a}'',\boldsymbol{b}'') \\ & = & \operatorname{ms}(\boldsymbol{b}',\boldsymbol{c}) \bullet \operatorname{ms}(\boldsymbol{a}' \bullet \boldsymbol{a}'',\boldsymbol{b}'') \\ & = & \operatorname{ms}(\boldsymbol{c},\boldsymbol{b}') \bullet \operatorname{ms}(\boldsymbol{a},\boldsymbol{b}'') \\ & \leq & \operatorname{ms}(\boldsymbol{c} \bullet \boldsymbol{a},\boldsymbol{b}). \end{array}$$

Proof of Proposition 4.5. Lemma 4.7 implies that if $\mathbf{d} = d_1 \dots d_m$ and $\mathbf{e} = e_1 \dots e_{m'}$ are two sequences such that $\mathbf{e} \succeq \mathbf{d}$, then

$$\operatorname{ms}(\boldsymbol{e}, \boldsymbol{d}) = e_1 \operatorname{ms}(\boldsymbol{d}, e_2 \dots e_{m'}).$$

Indeed, if $e_1 > d_1$, it is obvious, and if not, Lemma 4.7 with $\mathbf{c} := e_1$, $\mathbf{a} := d_2 \dots d_m$ and $\mathbf{b} := e_2 \dots e_m$ implies it.

Combining Point (5) of Lemma 4.6 and this fact, we get that if $a_s \succeq a_i$ for all $i \neq s$, then

$$\operatorname{ms}(\boldsymbol{a}_1,\ldots,\boldsymbol{a}_s)=a_{s1} \bullet \operatorname{ms}(\boldsymbol{a}_1,\ldots,\boldsymbol{a}_{s-1},a_{s2}\ldots a_{sn_s}).$$

This leads directly to an algorithm whose time complexity is $O(n^2 \log s)$:

- sort all sequences: $O(sn \log s)$,
- let $s = \epsilon$,
- repeat n times the following operations: append to s the first term of the maximal sequence a in the list, delete this first term from a and put the new a at the right place in the list (this last operation takes $O(n \log s)$).
- return s.

4.3.3. Proof of Theorem 1.2.

Proof. According to the previous discussion, it remains to prove that the optimal ordering starting at r can be obtained by doing ms on the optimal orderings of the subtrees T_i of T. But it is straightforward: the optimal ordering for T is a shuffle of the orderings of the T_i ; given S orderings on the subtrees T_i , the best shuffle we can get is ms by definition; for each of these orderings, the best is to maximize it with respect to the lexicographic order (ms is increasing).

The whole complexity is computed as follows: for a fixed root r, using the inequalities

$$n^2 \log n \ge \log n \sum_i n_i^2 \ge \sum_i n_i^2 \log n_i$$
 when $\sum_i n_i = n - 1$ and $n_i \ge 0$ for all i ,

we get by induction $O(n^2 \log n \log d)$ where d is the maximum degree of the tree. Doing the computation for each possible root provides the solution.

5. Pareto-optimal ordering

Finding a Pareto-optimal ordering seems to be a difficult task. It is even unclear whether it is possible to check in polynomial time that a given ordering is Pareto-optimal.

In some cases, the following criterion will be used for q=1.

Lemma 5.1. Let $f: \mathbb{R}^n \to \mathbb{R}^q$ be a strictly increasing map (ie $\mathbf{x} \neq \mathbf{x}'$ are such that $x_i \leq x_i'$ for all i, then $f(\mathbf{x}) \neq f(\mathbf{x}')$ and $f(\mathbf{x})_k \leq f(\mathbf{x}')_k$ for all k = 1, ..., q). Let \mathcal{S} be a subset of \mathbb{R}^n . If $f(\mathbf{x}^*)$ is Pareto-optimal on $f(\mathcal{S})$, then \mathbf{x}^* is Pareto-optimal on \mathcal{S} .

Table 4. Evaluation vector of an ordering for the graph of Figure 2

Proof. Let $y \in \mathbb{R}^n$ distinct from x^* . We have $f(y)_k < f(x^*)_k$ for a certain k. Since f is strictly increasing, we cannot have $x_i^* \leq y_i$ for all i. Therefore, x^* is Pareto-optimal.

In the following proposition, the search for a Pareto-optimal ordering is related to the testing cost problem.

Proposition 5.2. Any optimal solution of Testing Cost with c(v) = 1 for all $v \in V$ is a Pareto-optimal solution of Pareto-Optimal ordering.

Proof. Given an ordering v_1, \ldots, v_n , we have the following equality

$$\sum_{i=1}^{n} \sum_{e \in E \setminus E[v_1, \dots, v_i]} p(e)c(v_i) = |E|p(E) - \sum_{i=1}^{n} h_i.$$

Using the map $f(x_1, ..., x_n) = \sum_{i=1}^n x_i$ in Lemma 5.1, we get the required conclusion.

5.1. Pareto-optimality in trees. Proposition 1.1 claims that when H is a tree, finding a Paretooptimal ordering can be done in polynomial time. We prove now this proposition. The proof uses Lemma 5.1 and builds a special function f to transform the search of Pareto-optimality into a classical optimization problem.

Proof of Proposition 1.1. Denote by T = (V, E) the tree and assume that there is a weight function $w: E \to \mathbb{R}_+$.

Fix a root r. Define $w': V \to \mathbb{R}_+$ to be such that w'(v) := w(uv) where u is the neighbor of v on the path linking r and v and w'(r) = 0. Consider the problem – which we call P – of finding an ordering v_1, \ldots, v_n of the vertices of T, starting at a fixed vertex $v_1 = r$, such that

- (1) $v_{i+1} \in N(v_1, ..., v_i)$ for all i = 1, ..., n(2) the quantity $\sum_{i=1}^{n} (i-1)w'(v_i)$ is minimal.

Note that minimizing $\sum_{i=1}^{n} (i-1)w'(v_i)$ is equivalent to maximizing $\sum_{i=1}^{n} (n-i+1)w'(v_i)$, which is equal to $\sum_{i=1}^{n} h_i$ with $h_i = \sum_{\ell=1}^{i} w'(v_\ell)$. The sequence (h_i) is the one that has to be Pareto-optimal. Since choosing $v_{i+1} \notin N(v_1, \ldots, v_i)$ for a certain i provides $h_{i+1} = h_i$, we can restrict our search for a Pareto-optimal sequence to the sequences satisfying $v_{i+1} \in N(v_1, \ldots, v_i)$ for all $i=1,\ldots,n$.

If the ordering v_1, v_2, \ldots, v_n is a solution of problem P, Lemma 5.1 with $f(x_2, \ldots, x_n) = \sum_{i=2}^n (i - 1)^{n-1} f(x_i)$ 1) x_i ensures that the ordering is Pareto-optimal.

Horn [6] proved that P can be solved in polynomial time (see [1] for a fast algorithm, in $O(n \log n)$.

Doing it for each possible root $r \in V$ eventually leads to the Pareto-optimal ordering.

Note that the ordering obtained with this method is not necessarily a maximal one for the lexicographic order. Indeed, consider the tree in Figure 2 and the orderings from Tables 4 and 5. The ordering in Table 4 is obtained by applying the greedy algorithm presented earlier. Moreover, since the sufficient condition of Proposition 4.3 is satisfied, this ordering is maximal for the lexicographic order. The value of the objective function $\sum_{i=1}^{n} h_i$ for this ordering is 104. For the ordering presented in Table 5 the value is 116. The ordering obtained by the method described in the proof is therefore not necessarily maximal for the lexicographic order.

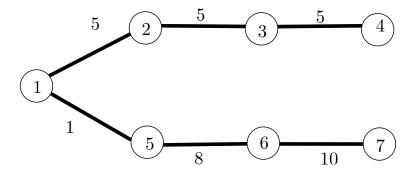


FIGURE 2. A rooted tree

Table 5. Evaluation vector of an ordering for the graph of Figure 2

6. A COMBINATORIAL ANALOG OF THE FUNDAMENTAL THEOREM IN ARITHMETICS

We prove now the combinatorial analog of the fundamental theorem of arithmetic (Theorem 1.3). As in Subsection 4.3.2, we forget the initial motivation and deal only with sequences. Recall that a sequence $\mathbf{p} = p_1 \dots p_n$ is prime if $p_i p_{i+1} \dots p_n \succeq \mathbf{p}$ for each $i \geq 1$, where \leq is the lexicographic order on the sequences.

We have an easy lemma

Lemma 6.1. Let p and p' be two prime sequences such that $p \leq p'$. Then $ms(p, p') = p' \cdot p$.

Proof. It is a consequence of the algorithm described in the proof of Proposition 4.5. \Box

Proof of Theorem 1.3. Let us write $\mathbf{a} = a_1 \dots a_n$. Define $i_1 := 1$ and $\mathbf{a}_1 := \mathbf{a}$. Then define i_2 to be the first index $i > i_1$ such that $a_i a_{i+1} \dots a_n \prec \mathbf{a}$, let $\mathbf{a}_2 := a_{i_2} a_{i_2+1} \dots a_n$. Define i_3 to be the first index $i > i_2$ such that $a_i a_{i+1} \dots a_n \prec \mathbf{a}_2$ and let $\mathbf{a}_3 := a_{i_3} a_{i_3+1} \dots a_n$. And so on. Let r be the index of the last \mathbf{a}_j defined (and by convention $i_{r+1} := n+1$).

Defining p_j to be $a_{i_j}a_{i_j+1}\dots a_{i_{j+1}-1}$, we get

$$a = p_1 \bullet p_2 \bullet \ldots \bullet p_r$$
.

Now, we have the following claim.

CLAIM. The p_i are all prime and we have

$$p_1 \succeq p_2 \succeq \ldots \succeq p_r$$
.

Let us first prove that p_j is prime for any j. Actually, we will prove it for p_1 , the proof being exactly the same for the other indices.

Let us suppose that p_1 is not prime. Then there is an $i \in \{2, ..., i_2 - 1\}$ such that

$$a_i a_{i+1} \dots a_{i_2-1} \prec a_1 a_2 \dots a_{i_2-1}.$$
 (1)

By definition of i_2 , we have also

$$a_i a_{i+1} \dots a_{i_2-1} a_{i_2} a_{i_2+1} \dots a_n \succeq a_1 a_2 \dots a_{i_2-1} a_{i_2} a_{i_2+1} \dots a_n.$$
 (2)

Equation (1) implies that $a_i \leq a_1$. Equation (2) implies that $a_i \geq a_1$. Together, they imply that $a_i = a_1$. We can therefore delete a_1 and a_i from their first places. Again, Equation (1) implies that $a_{i+1} \leq a_2$ and Equation (2) implies that $a_{i+1} \geq a_2$, which in turn leads to $a_{i+1} = a_2$. Going on in the same way, we get eventually:

$$a_i = a_1, a_{i+1} = a_2, \dots, a_{i_2-1} = a_{i_2-i}.$$

Using these equalities in Equation (2), we get that

$$a_{i_2}a_{i_2+1}\ldots a_n \succeq a_{i_2-i+1}a_{i_2-i+2}\ldots a_n$$
.

But, by definition of i_2 , we have $\mathbf{a} \succ a_{i_2} a_{i_2+1} \dots a_n$ and $\mathbf{a} \preceq a_{i_2-i+1} a_{i_2-i+2} \dots a_n$. A contradiction. Let us now prove the chain of inequalities. Again we will only prove that $\mathbf{p}_1 \succeq \mathbf{p}_2$, since the proof of the other inequalities is strictly the same.

Denote $p_1 = p_{11}p_{12} \dots p_{1n_1}$ and $p_2 = p_{21}p_{22} \dots p_{2n_2}$.

Assume for a contradiction that $p_1 \prec p_2$. By construction, we have

$$p_1 \bullet p_2 \bullet \ldots \bullet p_r \succ p_2 \bullet \ldots \bullet p_r$$
.

Thus, if $n_2 \le n_1$, we would have $p_{11} = p_{21}$, $p_{12} = p_{22}$, ..., $p_{1n_2} = p_{2n_2}$, which contradicts $\boldsymbol{p}_1 \prec \boldsymbol{p}_2$. Thus $n_2 > n_1$. But then we have $p_{11} = p_{21}$, $p_{12} = p_{22}$, ..., $p_{1n_1} = p_{2n_1}$ and

$$p_2 \bullet \ldots \bullet p_r \succ p_{2n_1+1}p_{2n_1+2}\ldots p_{2n_2} \bullet p_3 \bullet \ldots \bullet p_r,$$

which is in contradiction with the definition of i_3 .

The claim is proved.

Thus, we have according to Lemma 6.1, $a = ms(p_1, ..., p_r)$. The existence claimed by the theorem is proved. It remains to prove the uniqueness of the decomposition.

Assume for a contradiction that there are also r' prime sequences

$$p_1' \succeq p_2' \succeq \ldots \succeq p_{r'}'$$

such that $a = p'_1 \bullet p'_2 \bullet \ldots \bullet p'_{r'}$ and such that w.l.o.g. the length of p'_1 is strictly larger that the length of p_1 . We can then write

$$\mathbf{p}_1' = \mathbf{p}_1 \bullet a_{i_2} a_{i_2+1} \dots a_i,$$

for some index $i \geq i_2$. Since p'_1 is prime, we have

$$a_{i_2}a_{i_2+1}\ldots a_i\succeq \boldsymbol{p}_1\bullet a_{i_2}a_{i_2+1}\ldots a_i,$$

and since the length of the sequence on the left side is smaller than the length of the right's one, we have

$$a_{i_2}a_{i_2+1}\ldots a_ia_{i+1}\ldots a_n\succeq \boldsymbol{p}_1\bullet a_{i_2}a_{i_2+1}\ldots a_ia_{i+1}\ldots a_n,$$

which contradicts the definition of i_2 .

7. CONCLUSION AND OPEN QUESTIONS

Starting from a practical motivation, we have defined several challenging problems, which mix several features: scheduling, special order. Polynomial algorithms have been proposed in some special cases. One of these algorithms has motivated the notion of a new product – the *maximum shuffle product* – defined on the set of sequences of real numbers. It appeared that this product has a nice property, namely an analog of the fundamental theorem in arithmetic.

Open questions have already been outlined in the paper. But other questions remain open. For instance, there is the question of larger special cases for which the problems remain polynomial. Polynomial algorithms have been proposed when the hypergraph is a tree graph, but there may exist larger classes of hypergraphs for which these problems are polynomial.

It would also be interesting to find a suitable framework for the maximum shuffle product ms, in which we would have more than a monoid. For instance, if ms is the product, what is the addition?

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