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▶ To cite this version:

Farah Ben-Naoum, Christophe Godin. Algorithmic Height Compression of Unordered Trees. Journal of Theoretical Biology, 2016, 389, pp.237 - 252. 10.1016/j.jtbi.2015.10.030. hal-01412215

HAL Id: hal-01412215 https://hal.science/hal-01412215v1

Submitted on 8 Dec 2016

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Algorithmic Height Compression of Unordered Trees

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Abstract

By nature, tree structures frequently present similarities between their subparts. Making use of this redundancy, different types of tree compression techniques have been designed in the literature to reduce the complexity of tree structures. A popular and efficient way to compress a tree consists of merging its isomorphic subtrees, which produces a directed acyclic graph (DAG) equivalent to the original tree. An important property of this method is that the compressed structure (i.e. the DAG), has the same height as the original tree, thus limiting partially the possibility of compression. In this paper we address the problem of further compressing this DAG in height. The difficulty is that compression must be carried out on substructures that are not exactly isomorphic as they are strictly nested within each-other. We thus introduced a notion of quasi-isomorphism between subtrees, that makes it possible to define similar patterns along any given path in a tree. We then proposed an algorithm to detect these patterns and to merge them, thus leading to compressed structures corresponding to DAGs augmented with return edges. In this way, redundant information is removed from the original tree in both width and height, thus achieving minimal structural compression. The complete compression algorithm is then illustrated on the compression of various plant-like structures.

Keywords: Plants modeling, branching structures, self-nestedness, quasi-isomorphism, height redundancy.

1 1. Introduction

Many plants like trees exhibit complex branching structures. These struc-2 tures contain large numbers of components (branches, leaves, flowers, etc.) 3 whose organization in space may give a mixed feeling of both order and disor-4 der. The impression of disorder often comes from the fact that components 5 apparently do not follow any clear deterministic rule and that spatial dis-6 tribution of organs do not show exact symmetries. However, most of the 7 time, behind this semi-chaos, the same branching structures also show some 8 order in their organization as many sub-structures look similar, hierarchies 9 can be identified among branches and some symmetries in shapes, although 10 none exact-ones, seem to exist. During decades, botanists have strived to 11 identify rules characterizing this order despite the apparent disorder in plant 12 architectures, introducing notions such as repetition, modularity, gradients, 13 symmetry of branching structures, e.g. Troll (1937); Arber (1950); Halle 14 et al. (1978); Harper et al. (1986); Barthelemy (1991). These rules, mainly 15 of qualitative nature, gave key insights on how to address the notion of order 16 in plant architectures. 17

In the early 70's, computational formalisms started to emerge to repre-18 sent formally the architecture of branching structures, e.g. Lindenmayer 19 (1971); Lindenmayer and Rozenberg (1972); Lindenmayer (1975); Honda 20 (1971). These new conceptual tools made it possible to consider the analysis 21 of branching systems organization from a new and quantitative perspective. 22 Many approaches started from there to describe models as finite automata or 23 grammars that could reproduce by simulation the branching organization of 24 plants, e.g. Borchert and Honda (1984); Reffye et al. (1989); Prusinkiewicz 25 and Lindenmayer (1990). Among them, L-systems emerged in the plant mod-26 eling community as the most commonly used paradigm -based on rewriting 27 rules- to model branching system development (Prusinkiewicz (1998)). Most 28 of these approaches started from the design of a formal model embedding 29 biological assumptions, and were defined to produce algorithmically branch-30 ing systems similar to those observed in real plants. By contrast, the reverse 31 (inverse) problem, namely building algorithmically a computational finite 32 representation (model) from given observed branching systems, was much 33 less studied. 34

On fractal images this inverse problem was solved on two dimensional images by Barnsley (1988, 2006) using models of fractal geometry. However, these models rely on iterated function systems (IFS) which correspond to purely geometrical transformations. In particular, they don't take into
 account topology of the studied biological structures.

To account on inference, many approaches based on L-systems were also developed and reviewed later on in (Ben-Naoum (2009)). However, due to the complexity of the general problem, these different approaches make various simplifying assumptions. These assumptions in general are not satisfactory in the context of study of biological organisms such as plants structures.

One particularly important simplifying hypothesis consists of focusing on 45 the result of the developmental process without considering the intermediate 46 stages of development. This leads to a whole class of approaches in which 47 models are constructed to capture plant branching architecture in a minimal 48 way. A first approach was proposed by Viennot et al. (1989) based on a 40 statistical analysis of binary tree organization. Subtrees were characterized 50 by an integer representing their amount of asymmetry (the so called Holton-51 Strahler -HS- number (Deussen and Lintermann (2002))). Then parameters 52 were inferred from data to describe statistically the probability of finding a 53 tree with HS number n knowing the HS number m of the parent tree. A 54 similar statistical but more general approach was later used by Durand et al. 55 (2005, 2007), based on Markovian processes to describe subtrees frequencies 56 at any given node. These inference approaches are aimed to capture the 57 statistical properties of tree organization. Viewed as compression techniques, 58 they provide approximated strategies to compress trees, relying on different 59 types of a priori assumptions (e.g. Markovian property). An exact and 60 deterministic approach was recently introduced in Godin and Ferraro (2010). 61 In this latter work, the authors addressed the problem of recognizing sim-62 ilar, possibly nested, patterns in plant structures, and to compress them as 63 directed acyclic graphs (DAGs) in a reversible way. Trees were represented 64 as unordered labeled tree structures, where no order is assumed on the chil-65 dren of any node of the trees. Trees whose compression are linear DAGs 66 define the class of self-nested trees, i.e. trees that have very high compress-67 ibility. From this, a degree of self-nestedness could be defined for any tree 68

as the normalized edit-distance of this tree to the class of self-nested trees
 embedding the original tree, and a polynomial-time algorithm was designed
 to compute this distance.

The tree compressions techniques defined in Godin and Ferraro (2010) are based on the merging of isomorphic subtrees. Any two subtrees that are merged therefore result in a structure with the same height. As a result, the compressed DAG, where all the isomorphic subtrees have been merged, has



Figure 1: Basic example of width and height tree reduction

the same height as the original tree. If indeed compression occurs in width, no 76 compression occurs in height. However, some trees such as high dichotomic 77 trees or fish bone-like trees have a repetitive structure in height and are 78 therefore poorly compressed by this technique. In this paper, we consider 79 the problem of maximally compressing trees both in width and height with 80 no loss of information. This idea is illustrated in Fig.1. In section 3, we recall 81 the definition of tree reduction and self-nestedness introduced in Godin and 82 Ferraro (2010). We then introduce and study in section 4 a weak extension 83 of tree isomorphism, called quasi-isomorphism, that will make it possible 84 to identify similar (but not identical) tree patterns at different heights of a 85 tree. Based on these definitions and their properties, we present in section 86 5 an algorithm able to compress in height the reduction of any tree. This 87 algorithm is then applied to both theoretical tree-like structures and vegetal 88 branching systems to characterize its compression ability. 80

For seek of clarity all property proofs described in the main text are given in the Online Supplementary Material.

92 2. Notational conventions

An undirected graph, is a pair G = (V, E) where V denotes a finite set 93 of vertices and E a finite set of unordered pairs of vertices called edges. A 94 *complete graph* is a simple undirected graph in which every pair of distinct 95 vertices is connected by a unique edge. A *clique* in an undirected graph 96 G = (V, E) is a subset of the vertex set $C \subseteq V$, such that the subgraph 97 induced by C is complete (in some cases, the term clique may also refer to 98 the subgraph). A maximal clique is a clique that cannot be extended by gc including one more adjacent vertex from the original set of vertices. 100

¹⁰¹ A directed graph, is a pair G = (V, E) where all edges correspond to ¹⁰² ordered pairs of vertices. Let (x, y) be an edge in E, x is called a parent ¹⁰³ of y and y is a *child* of x. A vertex that has no child is called a *leaf*. We ¹⁰⁴ will denote, in the sequel, by *child*(x) the set of all the children of x, and ¹⁰⁵ *parent*(x) the set of all parents vertices of x. In a directed graph G, the ¹⁰⁶ *Indegree* of a vertex v, denoted $deg^{-}(v)$, is the number of its parents, and ¹⁰⁷ its *Outdegree*, denoted $deg^{+}(v)$, is the number of its children. The Indegree ¹⁰⁸ and the Outdegree of G are respectively the maximum Indegree and the ¹⁰⁹ maximum Outdegree of all G vertices.

A path from a vertex x to a vertex y is a (possibly empty) sequence of edges $\{(x_i, x_{i+1})\}_{i=1,M-1}$ such that $x_1 = x$, $x_M = y$, and M correspond to |P| the length of P. For a path P we denote by V_P the set of all vertices that belong to P. In an other hand, $P = \lambda$ if P is an empty path. Where we denote by λ the empty path. Analogously, we say that there are no path between x and y, or there is an empty path.

A vertex x is called an *ancestor* of a vertex y (and y is called a *descendant* of x), noted $x \prec y$, if there exists a path from x to y. Analogically, for the edges e = (x, y) and e' = (x', y') we can say that $e \prec e'$ iff $y \prec x'$.

Given two paths $P_1 = \{(x_i, x_{i+1})\}_{i=1,N-1}$ and $P_2 = \{(y_j, y_{j+1})\}_{j=1,M-1}$:

• for
$$x_N = y_1$$
 we define the paths union
 $P_1 \cup P_2 = \{(x_1, x_2), ..., (x_{N-1}, x_N), (y_1, y_2), ..., (y_{M-1}, y_M)\}$

• we denote $V_{P_1} \cap V_{P_2}$ the vertices paths *intersection* i.e. the subset of vertices common to both P_1 and P_2

- we say that
- $\begin{array}{ll} {}_{125} & P_1 \subset P_2 \text{ if the sequence made by the subset of edges common to} \\ {}_{126} & \text{both } P_1 \text{ and } P_2 \text{ correspond to } P_1 \text{ (path inclusion)} \end{array}$
- $P_{127} P_1 \prec P_2 \Leftrightarrow x_N \prec y_1$

$$-P_1 \multimap P_2 \Leftrightarrow x_N = y_1 \text{ (paths succession)}$$

In a directed graph, we call *cycle* a non empty path whose extremities coincide to the same vertex. If the *path is elementary*, i.e. does not pass twice through the same vertex, it is called *elementary* cycle.

A directed acyclic graph (DAG) is a graph containing no cycle (but which may contain undirected cycles), (Preparata and Yeh (1973)). A linear DAG is a DAG containing at least one path that goes through all its vertices.

A tree T is a connected graph containing neither directed of undirected cycle. A rooted tree is a tree such that there exists a unique vertex, called the root, which has no parent vertex. In a tree, each vertex, different from
the root, has exactly one parent vertex. In the following, a rooted tree is
simply called a tree.

In this paper, we consider *rooted unordered trees*, meaning that the order among the sibling vertices of any given vertex is not significant.

The degree deg of a tree is the maximum number of children of a vertex of T. The notation |T| represents the number of T vertices.

The height h(x) of a vertex x in a DAG is the length of the longest path from x to a leaf. The height h(D) of a DAG D is the height of its root vertex, and its width, denoted l(D), is the maximum number of vertices of the same height in all the DAG.

¹⁴⁸ A subtree is a particular connected subgraph of a tree. Let x be a vertex ¹⁴⁹ of a tree T = (V, E), T[x] is a complete subtree if it is the maximal subtree ¹⁵⁰ rooted in x with:

¹⁵¹ T[x] = (V[x], E[x]), where $V[x] = \{y \in V/x \text{ is the ancestor of } y\}$ and ¹⁵² $E[x] = \{(u, v) \in E/u \in V[x], v \in V[x]\}$. In the sequel, we will only consider ¹⁵³ complete subtrees and use the simpler term "subtree".

Let us consider two trees, $T_1 = (V_1, E_1)$ and $T_2 = (V_2, E_2)$. A bijection ϕ from V_1 to V_2 is a *tree isomorphism* if for each $(x, y) \in E_1$, $(\phi(x), \phi(y)) \in E_2$. If there exists an isomorphism between T_1 and T_2 , the two structures are thus identical up to a relabeling of their components. In this case we say that T_1 is *isomorphic to* T_2 .

¹⁵⁹ A multi-set is a set of typed elements such that the number of elements ¹⁶⁰ of each type is known. It is defined as a set of pairs $M = \{(k, n_k)\}_k$ where ¹⁶¹ k varies over the element types and n_k is the number of occurrences of type ¹⁶² k in the set.

163 **3.** Tree reduction

164 3.1. Definition

We consider here the principle of *tree reduction* as was defined by Godin 165 and Ferraro (2010). The aim of the *reduction* is to transform a rooted un-166 ordered tree T into a directed acyclic graph noted D = (V, E). Each vertex of 167 the D corresponds to an equivalence class of T subtrees. All subtrees belong-168 ing to the same class are *isomorphic*. All DAG vertices (classes) connected 169 by edges are ordered by the *ancestor* partial order relation, noted \prec , between 170 the subtrees which they represent. In other words, let A and B two DAG171 vertices where $A \preceq B$, then the tree of class B is isomorphic to a subtree of 172



Figure 2: a.i. a tree T, and ii. the associated reduction graph $\mathcal{R}(T)$, b.i. the tree T_1 , and ii. its reduction graph $\mathcal{R}(T_1)$

the tree of class A. Each DAG contains a single root vertex corresponding to the class of the entire tree T and one terminal vertex representing the class of all T leaves.

To obtain a graph equivalent to the tree T, additional information is 176 added in the DAG edges, as edges weights. For any edge (c_i, c_j) , its weight 177 $n(c_i, c_i)$ is defined by the number of occurrences of the subtree of class c_i 178 in this of class c_i . The obtained graph is called the reduction graph of T, 179 and noted by $\mathcal{R}(T) = (V, E)$, where E is a multi-set whose elements have the 180 form $((c_i, c_i), n(c_i, c_i))$. The construction of the reduction graph of a tree can 181 be carried out in time $O(|T|^2 deg(T) \log deg(T))$ (Godin and Ferraro (2010)). 182 Fig.2.a shows an example of a tree and its reduction graph. 183

In the following sections we will consider DAGs corresponding to reduction graphs, i.e. DAGs whose edges are augmented by weights. \mathcal{D} denotes the class of all DAGs.

187 3.2. Self-nested trees

Let us define the set of *self-nested* trees, as the trees in which all subtrees with identical height are isomorphic. Godin and Ferraro (2010) showed that self-nested-trees have the following characterizing properties:

- any two subtrees are either isomorphic or included one into another (one is a subtree of the other).
- their reduction $\mathcal{R}(T)$ is a linear *DAG*.

The reduction of the tree in Fig.2.b.i is the linear DAG depicted in Fig.2.b.ii. By contrast, the tree of Fig.2.a.i is not self-nested as its reduction is a non linear DAG (Fig.2.a.ii). ¹⁹⁷ Based on these definitions, the degree of self-nestedness of any tree T can ¹⁹⁸ be quantified by computing the distance between T and the smallest self-¹⁹⁹ nested tree (noted NEST) that embeds it. This distance is null if the tree is ²⁰⁰ in the class of self-nested tree and augments as the tree contains increasingly ²⁰¹ self-nested structures. Godin and Ferraro (2010) showed that this distance ²⁰² and the corresponding NEST of a tree T can be computed in polynomial ²⁰³ time by a DAG linearization algorithm.

An important property of both the reduction graph and the NEST of a 204 tree is that they preserve the height of nodes in the original tree. This means 205 that vertices in either the exact (the reduction graph) or the approximated 206 ones (the NEST) corresponding to vertices in the original tree have exactly 207 the same height. In Fig.2.b for example, the green node in the reduction 208 graph has a height of 3 (Fig.2.b.ii), corresponding exactly to the height of 209 the subtrees that it refers to in the original tree rooted also in green vertices 210 (Fig.2.b.i). The same property is true for all the other colors. As a conse-211 quence, a tree of height N will necessarily have a compression with a number 212 of vertices greater than N. In many cases, when N is of the same order of 213 magnitude as the tree size |T|. This is clearly a limitation of the previous 214 contraction procedures. 215

We are therefore lead to study how to compress trees in height as well as in width. For this, we need to relax the definition of isomorphism between trees to capture the notion of repetition of tree patterns independently of their height.

4. Quasi-periodic paths in a reduction graph

221 4.1. Intuitive idea of height reduction

Intuitively, a tree can be reduced in height if there are subtrees repeated in some way along some paths from the root to the leaves. But how to define exactly such repetitions?

Consider for instance tree of Fig.3.a. We see that the structure including 225 the subtrees originating at vertex B are repeated at different heights of the 226 original tree. The series of strictly nested subtrees $A(3) \subset A(2) \subset A(1) \subset A(2)$ 227 A(0) are similar, but not exactly isomorphic. Actually, if we discard the right-228 hand side subtree of A(i), i = 0, ..., 3 (illustrated in Fig.3.b), on the picture, 229 we remark that the remaining trees are indeed isomorphic. We shall say 230 that these trees are quasi-isomorphic, i.e. isomorphic for almost all of their 231 immediate subtrees except one, and shall show that nested quasi-isomorphic 232



Figure 3: a. a height regular tree T, b. the repeated pattern of T, c. Intuitive elimination of T pattern repetitions, d. the $DAG \mathcal{R}(T)$, e. the subDAG representing the regular repeated pattern, f. The intuitive height reduced DAG of $\mathcal{R}(T)$

sub-trees can be compressed by suppressing their repetition in the original tree and replacing them by loops over the repeated structure, as seen in the compressed tree of Fig.3.c. The objective is then to apply this principle of height compression directly to the associated DAG, as illustrated in graphs of Figs.3.d, 3.e and 3.f.

238 4.2. Vertex and edge signatures

239 **Definition 1 (Vertices signature)** Let $\mathcal{R}(T) = (V, E)$ a reduction graph. 240 We associate with V the matrix $\sigma \in \mathbf{N}^{|V| \times |V|}$ of vertex signature, where each 241 element σ_{xy} is defined as:

$$\sigma_{xy} = \begin{cases} n(x,y) & \forall ((x,y), n(x,y)) \in E \\ 0 & else \ where \end{cases}$$

In the matrix σ , the line associated with vertex x defines the signature σ_x of x. It represents the links of x with all its children vertices, and then describes the subtree T[x].

- **Example 1** Let $\mathcal{R}(T_1) = (V, E)$ the reduction graph of Fig.2.b.ii, where: ₂₄₆ $V = \{A, B, C, D, E\}$ and
- ²⁴⁷ $E = \{((A, B), 1), ((A, C), 2), ((A, D), 1), ((B, C), 1), ((B, E), 1), ((C, D), 1), ((A, C), 2), ((C, E), 1), ((C, D), 1), ((C, D), 1)\}$
- ²⁴⁹ The associated vertex signature matrix is given as follow:

 $A \quad B \quad C \quad D \quad E$ $\sigma_A = \begin{bmatrix} 0 & 1 & 2 & 1 & 0 \\ \sigma_B = \begin{bmatrix} 0 & 0 & 1 & 0 & 1 \\ \sigma_C = \begin{bmatrix} 0 & 0 & 0 & 1 & 1 \\ \sigma_D = \begin{bmatrix} 0 & 0 & 0 & 0 & 1 & 1 \\ \sigma_E = \begin{bmatrix} 0 & 0 & 0 & 0 & 0 & 1 \end{bmatrix}$

Property 1 $\forall x, y \in V, \sigma_x \neq \sigma_y$.

Definition 2 (Atomic path) We define an atomic path in a reduction
graph as a path of length 1.

 $_{254}$ $\langle x, y \rangle$ denotes the atomic path from a vertex x to a vertex y.

Definition 3 Given a vertices signature matrix σ , and given x, y two vertices where $\sigma_{xy} \neq 0$. We define the vector $\sigma_{x,y}$ as the vector σ_x with σ_{xy} is set to 0.

The vector $\sigma_{x,y}$ describes the constitution of the subtree associated with the vertex x up to the subtree structure of its child y. This leads to the following property.

Property 2 $\sigma_{x,y} = \sigma_{x',y'} \Leftrightarrow x$ is isomorphic to x' up to a child node y in xand a child node y' in x'.

Property 2 sets up the basis of a new relation which we call quasi-isomorphism between T[x] and T[x'].

Example 2 Given two trees rooted in X and X', illustrated respectively in Fig.4.a and 4.c, and associated with the vertices x and x' in the corresponding reduction graphs of Fig.4.b and 4.d. Note that the subtrees rooted in Y and Y' are distinct. We obtain:

$$\left. \begin{array}{l} \sigma_{a_1} = \sigma_{a_2} \\ \sigma_{xa_1} = \sigma_{x'a_2} = 1 \\ \sigma_{b_1} = \sigma_{b_2} \\ \sigma_{xb_1} = \sigma_{x'b_2} = 1 \end{array} \right\} \stackrel{Def.3}{\Longleftrightarrow} \sigma_{x,y} = \sigma_{x',y'}$$

Then X and X' are quasi-isomorphic sub-trees regarding their respective descendants Y and Y'.



Figure 4: a. and c. quasi-isomorphic trees T[X] and T[X'], b. and d. the corresponding reduction graphs $\mathcal{R}(T[X])$ and $\mathcal{R}(T[X'])$. Black vertices display their differences

Definition 4 (Edge signature) The signature of an edge (x, y), denoted by $\sigma_{x|y}$, is the line $\sigma_{x,y}$ augmented by the additional value of σ_{xy} on the right side.

Example 3 The edges signatures associated with the graph $\mathcal{R}(T_1)$ of Example 1 are given as follow:

 $\sigma_A = \begin{bmatrix} 0 & 1 & 2 & 1 & 0 \end{bmatrix},$ 276 $\sigma_{A,B} = \begin{bmatrix} 0 & 0 & 2 & 1 & 0 \end{bmatrix}$ and $\sigma_{AB} = 1$ then: 277 $\sigma_{A|B} = \begin{bmatrix} 0 & 0 & 2 & 1 & 0 & 1 \end{bmatrix}$ likewise 278 $\sigma_{A|C} = \begin{bmatrix} 0 & 1 & 0 & 1 & 0 & 2 \end{bmatrix}$ $\sigma_{A|D} = \begin{bmatrix} 0 & 1 & 2 & 0 & 0 & 1 \end{bmatrix}$ $\sigma_{B|C} = \begin{bmatrix} 0 & 0 & 0 & 0 & 1 \end{bmatrix}$ 1 $\sigma_{B|E} = \begin{bmatrix} 0 & 0 & 1 & 0 & 0 \end{bmatrix}$ 1 279 $\sigma_{C|D} = \begin{bmatrix} 0 & 0 & 0 & 0 & 1 \end{bmatrix}$ 1 1 $\sigma_{C|E} = \begin{bmatrix} 0 & 0 & 0 & 1 \end{bmatrix}$ 0 $\sigma_{D|E} = [0 0 0 0 0 1]$

The edges signatures allow us to study the quasi-isomorphism relation in a detailed and formal way on the basis of tree reduction graphs.

282 4.3. Quasi-isomorphism of paths

283 4.3.1. Quasi-isomorphism of atomic paths

²⁸⁴ Definition 5 (Quasi-isomorphic atomic paths) In a reduction graph,

given two distinct atomic paths $\langle x, y \rangle$, $\langle x', y' \rangle$. We say that $\langle x, y \rangle$ and $\langle x', y' \rangle$ are quasi-isomorphic, noted $\langle x, y \rangle \cong \langle x', y' \rangle$, if and only if $\sigma_{x|y} = \sigma_{x'|y'}$.

287 Example 4 A. From Example 2:

$$\begin{array}{c} \sigma_{x,y} = \sigma_{x',y'} \\ \sigma_{xy} = \sigma_{x'y'} = 1 \end{array} \right\} \begin{array}{c} \stackrel{Def.4}{\longleftrightarrow} \sigma_{x|y} = \sigma_{x'|y'} \\ \stackrel{Def.5}{\longleftrightarrow} \langle x, y \rangle \cong \langle x', y' \rangle \end{array}$$

B. Given the set of edges signature computed in Example 3. There exist in this set, two equal edges signatures: $\sigma_{B|C} = \sigma_{C|D}$, which induce that $\langle B, C \rangle \cong \langle C, D \rangle$. This appears clearly in Fig.2.b.i in which the subtree associated with *B* except its child *C*, and the subtree associated with *C* except its child *D* are isomorphic.

Property 3 (Fundamental property) Let e, f be two atomic paths starting from x and let e', f' be two atomic paths starting from x', then

$$((e \cong e') \land (f \cong f')) \Rightarrow ((f = e) \land (f' = e'))$$

In other terms, at most one pair of quasi-isomorphic atomic paths can start from a pair of vertices x and x'.

298 4.3.2. Quasi-isomorphism of paths

²⁹⁹ Let's consider two paths P and Q, where $P = \{e_1, ..., e_n\}$, and $Q = \{f_1, ..., f_m\}$.

Definition 6 (quasi-isomorphic paths) We say that P is quasi-isomorphic to Q (noted $P \cong Q$) if and only if n = m and $e_i \cong f_i$ $(1 \le i \le n)$.

In other words, two paths are quasi-isomorphic if and only if their atomic paths are quasi-isomorphic when compared piecewise in the paths order.

Example 5 A. In the *DAG* $\mathcal{R}(T_1)$ (Fig.2.b) they are two quasi-isomorphic paths: $P = \langle B, C \rangle$ and $P' = \langle C, D \rangle$ from $\sigma_{B|C} = \sigma_{C|D}$.

B. In the *DAG* $\mathcal{R}(T_2)$ (Fig.5.a) they are two quasi-isomorphic paths: $P \cong P'$ with $P = \{\langle B, E \rangle, \langle E, I \rangle\}, P' = \{\langle C, F \rangle, \langle F, H \rangle\}$ from $\sigma_{B|E} = \sigma_{C|F} =$ [000100000001], and $\sigma_{E|I} = \sigma_{F|H} =$ [000000100001].

Notation: A path $P = \{\langle x_1, x_2 \rangle, ..., \langle x_{n-1}, x_n \rangle\}$ can be denoted as: $P = x_1 P x_n$ or $P = x_1 P_1 x_k P_2 x_n$ and so on. If $P = x_1 P_1 x_i P_i x_{i+1} P_n x_n$ and $|P_i| = 1$ we can note $P = x_1 P_1 x_i x_{i+1} P_n x_n$.

313 4.3.3. Quasi-periodic path

A quasi-periodic path is a path decomposable in strictly smaller sub-paths which are all quasi-isomorphic with each other, and which are not themselves strictly decomposable.



Figure 5: a.i. A tree T_2 , ii. its reduction $\mathcal{R}(T_2)$. We surround by dashed lines the two QIPs $P = \{\langle B, E \rangle, \langle E, I \rangle\}$ and $P' = \{\langle C, F \rangle, \langle F, H \rangle\}$, b. Illustration of QPP in a pattern of a DAG where the same color is used for all quasi-isomorphic atomic paths

Definition 7 (quasi-periodic path QPP) Given two vertices x, y with $x \prec y$. We say that xPy is a quasi-periodic path if there exist paths $P_1, ..., P_n$ $(P_i \neq \lambda)$ and $(n \ge 2)$ for which:

 $\bullet P = P_1 \cup \ldots \cup P_n$

• $P_1 \cong \ldots \cong P_n$

• $\forall i \leq n, P_i \text{ is not a quasi-periodic path.}$

P is called QPP(x, y), and the sequence $P_1, ..., P_n$ the normal decomposition of P.

³²⁵ **Property 4** The normal decomposition of P is unique.

Example 6 A. From the graph $\mathcal{R}(T_1)$ of Fig.2.b, the path $P = P_1 \cup P_2$ with P₁ = $\langle B, C \rangle$, $P_2 = \langle C, D \rangle$ is a QPP.

B. Moreover, in $\mathcal{R}(T_2)$ of Fig.5.a.ii, although the paths P and P' are two quasi-isomorphic paths, they do not form a QPP since they are not consecutive.

 $_{331}$ By definition, each QPP is a set of path chunks which are:

- embedded in the same global path,
- all quasi-isomorphic with each other,
- pairwise connected.

Therefore a *QPP* can be considered as a periodic path made by the repetition of quasi-isomorphic paths forming its normal decomposition. This is illustrated in Fig.5.b that shows a *QPP* whose normal decomposition is made up two subsequences. One sequence is indicated by red, blue and yellow.

Let us consider quasi-periodic paths P and Q with identical extremities.

340 **Property 5** Let xPy = QPP(x, y) and xQy = QPP(x, y) then P = Q.

³⁴¹ The property shows that these two paths coincide.

Property 6 Every subsequence of a normal decomposition of a QPP is a
normal decomposition of a smaller QPP.

Let $P_1, ..., P_n$ be a normal decomposition of a QPP P, the path corresponding to a subsequence $P_i, ..., P_{i+k}$ (i < n, 1 < k < n-i) of the normal decomposition of P is itself a QPP.

347 4.3.4. Maximal Quasi-periodic path

- ³⁴⁸ Definition 8 (Maximal quasi-periodic path MQPP) A maximal
- quasi-periodic path (noted MQPP) is a QPP maximum for path inclusion.

Let $P_1, ..., P_n$ be the normal decomposition of a MQPP P, then there is no $QPP Q \ (P \subset Q)$ such that $P_1, ..., P_n$ is a part of the normal decomposition of Q.

Let us denote by $MQPP(\mathcal{R}(T))$ the set of all the MQPPs of a reduction graph $\mathcal{R}(T) = (V, E)$.

Property 7 The set $MQPP(\mathcal{R}(T))$ can be computed in time $O(|E|^2h(\mathcal{R}(T)))$ $l(\mathcal{R}(T))).$

4.3.5. Relative positions of maximal quasi-periodic paths in a reduction graph
Let us now study the relative positions of two MQPPs in a reduction
graph. As for any two paths in a reduction graph, two MQPPs can either *intersect* or *not*, intersecting MQPPs can either be *nested* or *not*.



Figure 6: a. Tree T_3 and its reduction $\mathcal{R}(T_3)$ with two intersecting MQPPs. b. Tree T_4 and its reduction $\mathcal{R}(T_4)$ with two disjoint MQPPs. c. Tree T_5 and its reduction $\mathcal{R}(T_5)$ with a set of nested MQPPs

Example 7 A. In the reduction graph $\mathcal{R}(T_3)$ of Fig.6.a, the paths

³⁶² $P = \{\langle A, B \rangle, \langle B, G \rangle, \langle G, E \rangle\}$ and $P' = \{\langle G, C \rangle, \langle C, D \rangle\}$ are intersecting ³⁶³ MQPPs. They have a common vertex G.

- B. In the reduction graph $\mathcal{R}(T_4)$ of Fig.6.b , the paths $P = \{\langle B, D \rangle, \langle D, E \rangle\}$ and $P' = \{\langle C, G \rangle, \langle G, H \rangle, \langle H, I \rangle\}$ are two disjoint MQPPs.
- ³⁶⁶ C. In the reduction graph $\mathcal{R}(T_5)$ of Fig.6.c:
- $P_{1} = \{ \langle A, B \rangle, \langle B, C \rangle, \langle C, D \rangle \} \cup \{ \langle D, E \rangle, \langle E, F \rangle, \langle F, I \rangle \}$
- $P_2 = \{ \langle A, B \rangle, \langle B, C \rangle \}$
- $P_3 = \{ \langle D, E \rangle, \langle E, F \rangle \}$
- $_{370}$ 3 *MQPPs* with the nesting relations: $P_2 \subset P_1, P_3 \subset P_1$.

Let us describe the general structure of a nested MQPP P^1 . Let us call $P_1^1, ..., P_N^1$ the normal decomposition of P^1 . The motifs P_n^1 are all quasiisomorphic with each other and their union entirely covers P^1 . Therefore, to study the structure of P^1 one only needs to study the structure of one of them, say P_l^1 . Now let us consider the list of MQPPs included in P_l^1 . This list may be empty if P^1 is not nested and the MQPP is said to be simple. Otherwise, P_l^1 contains itself a list of MQPPs denoted P_l^{1j} for j = 1..J. If one of the P_l^{1j} is nested this decomposition recursively continues and P_l^{1j}



Figure 7: Indexing the component of a nested MQPP. Each motif is composed of a series of MQPPs and each MQPP is composed of a series of motifs.

can be decomposed into a list of quasi-isomorphic motifs denoted P_{lm}^{1j} for m = 1..M, and so on (see Fig.7). These series of decomposition forms a tree structure called the MQPP tree of P^1 . Interestingly, this tree contains many identical subtrees as indicated by the following property.

Property 8
$$\forall m \neq m', P_{l \ldots m'}^{1 j \ldots k} \cong P_{l \ldots m}^{1 j \ldots k}$$

Definition 9 (Maximally Nested MQPP) A maximally nested MQPP
 is a MQPP that is not strictly contained within a strictly greater MQPP.

Definition 10 (Normal decomposition of a DAG) The normal decomposition of a DAG D is the set \mathcal{N} of all its maximally nested MQPPs.

Therefore, the normal decomposition of a DAG contains only top level nested MQPPs which are either simple or decomposable in to finer MQPPs.

In general, the relative position of two *MQPPs* must respect constraints as indicated by the following property.

Let $P = \{e_1, ..., e_n\}$ be a path. If $e_n = \langle x, y \rangle$ we say that P terminates through x.

³⁹⁴ **Property 9** Let P and P' be two intersecting MQPPs:

- 1. If P and P' starts from the same vertex then $P \subset P'$ or $P' \subset P$.
- $_{396}$ 2. If P and P' are not nested then either:
- (a) at least one of the MQPPs terminates through the intersection
 point,



Figure 8: a. Patterns of relative positions of two intersecting MQPPs, b. Patterns of impossible situations of two intersecting MQPPs. See text for detail explanation

- (b) the intersection corresponds to a postfix of one of the two paths,
 (c) or both MQPPs belong to the same path.
- **Example 8** Let us illustrate the different situations of two non nested intersecting MQPPs cited in Property 9. Fig.8.a shows some possible situations of two non nested intersecting MQPPs, and Fig.8.b some impossible ones, so in:
- Figs.8.a.i, 8.a.ii, 8.a.iii and 8.a.iv at least one of the *MQPP*s terminates through the intersection point,
- Fig.8.a.v the intersection corresponds to a postfix of one of the two paths,
- Fig.8.a.vi both *MQPP*s belong to the same path,
- Figs.8.b.i, 8.b.ii and 8.b.iii there is no *MQPP* which terminates through the intersection point.



Figure 9: a.i. A height repeated branch, ii. the corresponding $MQPP = P_1 \cup ... \cup P_n$ in the associated DAG, iii. the equivalent height reduced DAG including a return edge and a returns number (n-1), b.i. a DAG, ii. the original DAG augmented with return edges (dashed edges), the result is a DAG with return edges

⁴¹² 5. Height compression of a reduction graph

Each MQPP can be decomposed in a unique manner as a sequence of quasi-isomorphic paths (normal decomposition) $P_1, ..., P_n$ (as in Fig.9.a.ii). We will make use of this repeated pattern to compress these MQPPs. This will be made by allowing introduction of loops in the original DAGs, as in Fig.9.a.iii. Let us introduce the notion of DAG with return edges.

⁴¹⁸ 5.1. DAGs with Return Edges

⁴¹⁹ A DAG with return edges is constructed from a DAG by adding to it ⁴²⁰ edges that induce oriented cycles, Fig.9.b.

⁴²¹ Definition 11 (DAG with return edges) Let D = (V, E) be a DAG. ⁴²² We consider a set E' of edges in $V \times V$ such that:

•
$$\forall (x,y) \in E', y \prec x \text{ in } D.$$

• As for any edge of D, we associate a weight n(x, y) with each edge (x, y) $\in E'$.

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• In addition we associate with each edge $(x, y) \in E'$ one of the children z of x if any, such that n(x, y) = n(x, z), z is called the exit vertex of the cycle induced by the return edge (x, y) and denoted ex(x, y).

The triplet D' = (V, E, E') verifying these conditions is called DAG with return edges, where E' is the set of return edges of D'. \mathcal{D}_{\uparrow} denotes the class of all DAGs with return edges.

Property 10 Let $B = (e_1, ..., e_n)$ an elementary cycle in a DAG with return edges D = (V, E, E'). There is a unique return edge $e_k \in E'$ in B denoted re(B). We called the path $\{e_{k+1}, ..., e_n, e_1, ..., e_{k-1}\}$ the principal path of B denoted pp(B), and the vertex $ex(e_k) = z$ the exit vertex of the cycle B, denoted ex(B).

Furthermore, we say that $x \in pp(B)$, for each vertex x in pp(B). In the DAG with return edges of Fig.10.e, for the elementary cycle $B_2 = ((x, y),$ (y, z), (z, w), (w, x): $re(B) = (w, x), pp(B) = \{(x, y), (y, z), (z, w)\}$ and ex(B) = e.

The MQPPs correspond to maximally repeated patterns on the paths of a DAG. It is thus possible to compress the DAG in height by compressing these repeated motifs. This will lead to construct DAGs with return edges representing the compressed trees. As we will show, the loop in these graphs can only be either nested or disjoint. This leads to introduce a special kind of DAGs with return edges, namely DAGs with nested returns.

447 5.2. DAG with nested returns

Definition 12 (DAG with nested returns) We define a DAG with nested returns as a DAG with return edges in which any two elementary cycles B_1 and B_2 (with respective vertex sets V_1 and V_2) are either nested ($V_1 \cap V_2 = V_1$ or $V_1 \cap V_2 = V_2$) or disjoint ($V_1 \cap V_2 = \emptyset$).

⁴⁵² We denote the class of all *DAGs* with nested returns by $\mathcal{D}_{\uparrow\uparrow}$. Note that ⁴⁵³ $\mathcal{D}_{\uparrow\uparrow} \subset \mathcal{D}_{\uparrow}$.

Example 9 Fig.11.a shows examples of patterns of vertices of $DAGs \in \mathcal{D}_{\uparrow\uparrow}$ and in Fig.11.b $DAGs \notin \mathcal{D}_{\uparrow\uparrow}$.



Figure 10: a. Cycle B_1 in a pattern of a DAG with returns D, b. The multiplicities of the set of entering edges in B_1 (Edges in bold), c. Cycle B_2 in the same graph D, d. The multiplicities of the set of entering edges in B_2 , e. the DAG D with the nested cycles B_1 and B_2 , f. the merged sets of multiplicities of entering edges in cycles B_1 and B_2



Figure 11: a. Patterns of valid DAGs in $\mathcal{D}_{\uparrow\uparrow}$, b. Patterns of DAGs not in $\mathcal{D}_{\uparrow\uparrow}$, (Dashed edges represent return edges)

Let $\mathcal{B}(D)$ denote the set of all cycles in a given $D \in \mathcal{D}_{\uparrow\uparrow}$. If $B_1, B_2 \in \mathcal{B}(D)$, the inclusion relation is denoted by $B_1 \subset B_2$.

Definition 13 (Entering edges in a cycle) Given D = (V, E, E') in $\mathcal{D}_{\uparrow\uparrow}$. Let B a cycle in $\mathcal{B}(D)$. The set of entering edges in B, denoted $\mathcal{E}(B)$, is the set of all edges $(x, y) \in E$ such that $y \in pp(B)$ and $x \notin pp(B)$.

⁴⁶¹ Definition 14 (Multiplicity of a cycle for an edge) Given $D \in \mathcal{D}_{\uparrow\uparrow}$ and ⁴⁶² let $B \in \mathcal{B}(D)$. With each edge e in $\mathcal{E}(B)$ we associate an integer μ_B called ⁴⁶³ the multiplicity of B for e.

⁴⁶⁴ From Fig.10.b, μ_{B_1} for (a, x) is 4, while it is 6 for edge (b, y).

The intuitive idea is to define on each edge entering a loop the number of 465 times this loop must be scanned, and these is the multiplicity of the loop for 466 this edge, Figs.10.b and 10.d. However, Fig.10.e shows a DAG with nested 467 returns composed by the fusion of DAGs with return edges of Figs.10.a and 468 10.c, where $B_1 \subset B_2$. Thus, the set of entering edges in B_1 is a subset of 469 entering edges in B_2 , Figs.10.b and 10.d. So the multiplicity of each entering 470 edge in simultaneously B_1 and B_2 correspond to the list of the multiplicities 471 of B_1 and B_2 ordered according to the nesting order of the loops starting 472 from the inner one, Fig. 10.f. More details are given in Fig. 12.a. 473

Property 11 Given D = (V, E, E') in $\mathcal{D}_{\uparrow\uparrow}$, $\forall x \in V$ there is at most an ordered list of elementary nested cycles containing x, noted $\mathcal{B}(x)$ where $\mathcal{B}(x) = \{B_1, ..., B_{b(x)}\}, b(x) = |\mathcal{B}(x)|$ and $B_1 \subset ... \subset B_{b(x)}$.

In the *DAG* with nested returns of Fig.10.e there is two elementary cycles $B_1 = \{(x, y), (y, x)\}$ and $B_2 = \{(x, y), (y, z), (z, w), (w, x)\}$ with $B_1 \subset B_2$. Then $\mathcal{B}(x) = \mathcal{B}(y) = \{B_1, B_2\}, \ \mathcal{B}(z) = \mathcal{B}(w) = \{B_2\}$ and $\mathcal{B}(e) = \emptyset$.

Our aim is to show that DAGs with nested returns are in general compressed versions of DAGs, where the cycles can be unfolded to construct DAGs. Expanding a cycle in a DAG with nested returns corresponds to unfolding the return edge by successive repetitions of the principal path of the cycle. Let us illustrate this process on the graph of Fig.12.b from which, at first, the loop B_1 is expanded. Let $\mu_{max} = 2$ be the maximal multiplicity of all entering edges in B_1 . Expanding B_1 consists of:

• unfolding its return edge by the repetition of the principal path of B_1 μ_{max} times. Let $P_1, ..., P_{\mu_{max}+1}$ be the obtained repeated patterns with $P_1 = pp(B_1).$



Figure 12: a. A general pattern of a formal detailed DAG with nested returns, b. A pattern of a labeled DAG with nested returns D, c. expansion of the inner cycle of D, d. completely expanded DAG. Vertices from which start thick edges of the same color have a common set of descendant vertices

• redirect each entering edge in B_1 , of multiplicity μ_{max} , into the equivalent entering edge in the new expanded path, i.e. edge of the same pattern in $P_{\mu_{max}-\mu+1}$.

Fig.12.c shows the resulting DAG with nested returns. The expansion of the remaining loop B_2 is performed in the same manner, Fig.12.d.

⁴⁹⁵ **Property 12** Given D = (V, E, E') in $\mathcal{D}_{\uparrow\uparrow}$, it is possible to unfold D into a ⁴⁹⁶ DAG in time of $O(\mathcal{U} | V | | E' | deg^+(\mathcal{D}_{\uparrow\uparrow}))$.

⁴⁹⁷ \mathcal{U} is the maximal multiplicity of all cycles of $\mathcal{D}_{\uparrow\uparrow}$.

In the resulting DAG of Fig.12.d, we observe that the obtained expanded path forms a MQPP.

500 5.3. Compression of a DAG as a DAG with nested returns

We have just seen that DAGs with nested returns can be unfolded into DAGs by expending the cycles. We now consider the reciprocal question of compressing a DAG as a DAG with nested returns. To this end we will first analyze how MQPPs can be folded and then apply the resulting algorithm to a maximal set of MQPPs in the original DAG.

506 5.3.1. Compression of a MQPP

Let P be a MQPP in a DAG D and $P_1, ..., P_n$ be its normal decomposi-507 tion. The idea is to replace the sequence $P_1, ..., P_n$ by a loop over P_1 repeated 508 n times. In practice the original DAG must be edited in the following way. 509 General compression algorithm of a MQPP: Let D = (V, E, E') be a 510 DAG of $\mathcal{D}_{\uparrow\uparrow}$, and let $P = P_1 \cup \ldots \cup P_n$ the MQPP(x, y) with $P_i = x_{i,1}P_i x_{i,m}$ 511 $(1 \leq i \leq n, m \geq 2), x = x_{1,1}$ and $y = x_{n,m}$, as represented in Fig.13.a. We 512 denote by B_p the cycle obtained from the compression of P, and the resulting 513 DAG with nested returns by $D_{/P}$. Then $D_{/P}$ is computed in the following 514 515 steps:

516	1.	Creation of a return edge, Fig.13.b:
517		• add a return edge to E' from $x_{1,m-1}$ to x ,
518		• this edge has a weight $\omega = n(x_{i,m-1}, x_{i,m}),$
519 520 521		 this edge is also augmented with the information that when the loop ends, scanning should resume on the exit vertex y of x_{n,m-1}, labeled by χ.
522	2.	Redirection of entering edges of the $MQPP$, Fig.13.c:
523 524		• Replace each entering edge $(z, x_{i,j})$ of the $MQPP$ by the equiva- lent entering edge in the cycle B_P , i.e. edge $(z, x_{1,j})$,
525		• this edge has a weight $n(z, x_{i,j})$,
526 527 528		• its multiplicity list corresponds to the multiplicity list of the edge $(z, x_{i,j})$ augmented by the multiplicity of the loop B_P for this edge, that is $n - i$.
529	3.	Suppression of the $MQPP$ repeated patterns, Fig.13.d:
530		• Remove edges which belong to the paths $P_2,, P_n$,
531 532 533		• the last edge of P_1 $(x_{1,m-1}, x_{1,m})$ is redirected on the exit vertex χ . It is thus replaced by the edge $(x_{1,m-1}, \chi)$ whose weight is also ω ,
534		• Edit the sets V, E and E' .



Figure 13: a. Formal pattern of a $MQPP \ P = P_1 \cup ... \cup P_n$, b. creation of the return edge of P, c. redirection of an entering edge of P into its pattern in P_1 , d. Suppression of the repeated patterns $P_2, ..., P_n$



Figure 14: a. DAG $\mathcal{R}(T_1)$ b. creation of the return edge, c. redirection of the entering edges of P (colored edges), d. $\mathcal{R}(T_1)_{/P}$

Property 13 The algorithm can be applied on the MQPP in time O(|E|).

Example 10 Let us compress the $MQPP \ P = P_1 \cup P_2$ with $P_1 = \langle B, C \rangle$ and $P_2 = \langle C, D \rangle$ of Fig.2.b. The compressed graph $\mathcal{R}(T_1)_{/P}$ of Fig.14.c is computed by the following steps:

539	1.	create the return edge (Fig.14.a):
540		$re(B_P) = (B, B)$ of weight 1 and exit vertex D ,

- ⁵⁴¹ 2. redirect all entering edges of P into their patterns in B_{P_1} (Fig.14.b): ⁵⁴² edge $((A, B), 1, \{\})$ is replaced by $((A, B), 1, \{1\})$,
- edge $((A, C), 2, \{\})$ is replaced by $((A, B), 2, \{0\})$,
- ⁵⁴⁴ 3. suppression of C and all its incident edges, and redirection of edge ⁵⁴⁵ (B, C) on the exit vertex D (Fig.14.c).

Note on this example that due to the *MQPP* compression several edges can
appear between two vertices.

548 5.3.2. Gain of MQPP compression

Let us consider a $MQPP \ P$ in $D \in \mathcal{D}_{\uparrow\uparrow}$, with $P = P_1 \cup ... \cup P_n$, and $P_i = x_{i,1}P_ix_{i,m}$ ($1 \leq i \leq n, m \geq 2$). Let us denote $|\overline{V_P}|$ and $|\overline{E_P}|$ the number of vertices and respectively the number of edges removed from Pcompression. Formally

$$|\overline{V_P}| = |V_P| - |V_{P_1}|, \ |\overline{E_P}| = \left(\sum_{i=2}^n \left(\sum_{j=1}^{m-1} deg^+(x_{i,j})\right)\right) - 1.$$

Let α be the size of the computational representation of a vertex in V, and let β be the size of the computational representation of an edge in E.

Definition 15 (compression gain of a MQPP) We define the P compression gain, noted g(P), by $g(P) = \alpha |\overline{V_P}| + \beta |\overline{E_P}|$.

Property 14 (compression gain of two MQPPs) Given two paths $P, P' \in MQPP(D)$. If P and P' are either disjoint or nested, then the gain

$$g(\{P, P'\}) = g(P) + g(P').$$

Let P and P' two MQPPs in a DAG D. $D_{/P/P'}$ denotes the DAG with nested returns obtained by successive compressions of P and P', and the gain $g(D_{/P/P'}) = g(\{P, P'\}).$

562 5.3.3. Selection of a maximal set of non intersecting MQPPs with maximal 563 compression gain

If two MQPPs are intersecting but no nested then the compression of one of them may impair or prevent the compression of the other. Therefore as the compression of the two MQPPs is mutually exclusive, the compression of the DAG will depend on the choice of the compression of either of them. We say that the DAG compression is ambiguous.

Given a DAG D, let M be a subset of MQPP(D).

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Definition 16 M is unambiguous for compression if $\forall P, P' \in M, P$ and P'are either nested or disjoint.

In other terms, M is unambiguous for compression if all its MQPPs can be compressed independently. Therefore, if M is ambiguous for compression one needs to make choices to eliminate ambiguity in compression while maximizing the compression gain.

⁵⁷⁶ For this, consider the equivalence relation \wedge on M such that:

 $P \wedge P' \Leftrightarrow P$ and P' are neither disjoint or nested.

⁵⁷⁸ Let $\{M_i\}_i$ be the set of equivalence classes of \wedge (of Fig.15.a). In a class M_i ⁵⁷⁹ MQPP's are pairwise intersecting with MQPP's of the same class. Note ⁵⁸⁰ that classes themselves are disjoint.

⁵⁸¹ Our problem is thus to select a subset of MQPPs in M_i such that this ⁵⁸² subset is unambiguous for compression, and the total gain of these MQPP's ⁵⁸³ is maximal, Fig 15.b.



Figure 15: a. Pattern of a DAG with M_1, M_2, M_3 the equivalence classes of \wedge , i.e.maximal MQPP sets ambiguous for compression (paths of the same color belongs to the same set M_i), b. the corresponding unambiguous subsets M'_1, M'_2, M'_3

To solve this problem, let us model the intersection map in class M_i as a graph: $G_{M_i} = (V_{M_i}, E_{M_i})$ such that each vertex of V_{M_i} represents a MQPPof M_i , and there is an edge between two vertices P and P' of V_{M_i} if P and P'are either disjoint or nested. With this graph definition we can characterize sets of MQPP sunambiguous for compression:

Property 15 M_i is unambiguous for compression if and only if G_{M_i} is a complete graph.

Therefore, our problem is reduced to finding in the graph G_{M_i} a clique 591 $G_{M'_i}$ (i.e. a complete subgraph of G_{M_i}) with maximal gain. This optimization 592 problem is known to be NP-complete (Feige et al. (1991)), and several ap-593 proaches have been developed to solve it (Battiti and Protasi (2001); Bomze 594 et al. (1999); Feige (2004)). Here we use the TABU algorithm (Gendreau 595 et al. (1993)), as described in section 15 of the Online supplementary ma-596 terial. Given a class M_i the algorithm returns a maximal subset M'_i unam-597 biguous for compression with maximal gain. 598

For a *DAG* D, let $M = \bigcup_i M_i$ and $M' = \bigcup_i M'_i$, and let us denote by $\overline{MQPP}(D)$ the maximal set of MQPPs of D unambiguous for compression, with:

$$\overline{MQPP}(D) = MQPP(D) \setminus (M \setminus M')$$

 $_{602}$ Where \setminus denotes the sets difference.

Let $\mathcal{N}(D)$ be the normal decomposition of a $DAG \ D, \overline{MQPP}(D)$ allows us to eliminate the compression ambiguity on the full $DAG \ D$.

⁶⁰⁵ Definition 17 (Unambiguous normal decomposition of a *DAG*)

The unambiguous normal decomposition of D, \mathcal{N}^* , is the set of all its maxi-

mally nested MQPPs unambiguous for compression, i.e. the set

$$\mathcal{N}^* = \mathcal{N}(D) \cap \overline{MQPP}(D).$$

Section 16 of the Online supplementary material provides examples of compressed graphs obtained from graphs of Fig.6.

610 5.3.4. DAG compression evaluation:

Given a DAG D = (V, E). Let R be the corresponding DAG with nested returns with gain g(R).

⁶¹³ Definition 18 (Compression factor of a *DAG* with nested returns)

⁶¹⁴ The compression factor of a DAG with nested returns R is defined as:

$$f(R) = \frac{g(R)}{\alpha |V| + \beta |E|}$$

where α , β are the size of the computational representations of respectively a vertex in V and of an edge in E. This measure represents the relative gain in size of the compressed graph compared to the original *DAG*.

⁶¹⁸ 5.4. Algorithm for the height compression of a finite tree

Based on the definition of MQPPs and on their compressibility property, it is possible to design an optimization algorithm to maximally compress a DAG in height.

Property 16 Let D = (V, E) be a DAG, the compression in height, denoted $\mathcal{H}(D)$, can be computed in time $O(|E|^2 h(D) l(D))$.

The main steps of the computation of $\mathcal{H}(D)$ are described as follows:

- ⁶²⁵ 1. Compute the set of vertex signatures and then the set of edge signatures ⁶²⁶ of D (see section 4.2).
- ⁶²⁷ 2. Given the set of edge signatures compute the set MQPP(D) (see sec-⁶²⁸ tion 4.3).

629	3. From $MQPP(D)$ determine the sets M then M',
630	and deduce $\overline{MQPP}(D)$ (see section 5.3.3).
631	4. Iterative computation of $\mathcal{H}(D)$:
632	(a) Initialization: Let $\mathcal{H} = (V, E, E')$ be the <i>DAG</i> with nested returns
633	for which the vertex and the edge set are equal to the vertex and
634	the edge set of D , and the set of return edges is an empty set.
635	(b) Repeat
636	i. compute $\mathcal{N}^*(\mathcal{H}) = \{P_1,, P_n\}$ the unambiguous normal de-
637	composition of \mathcal{H}
638	ii. $\mathcal{H} = \mathcal{H}_{/P_1//P_n}$
639	$\mathrm{until}\;\mathcal{N}^*(\mathcal{H})=\emptyset,$
640	5. Return the <i>DAG</i> with nested returns $\mathcal{H}(D) = \mathcal{H}$ of compression factor
641	$f(\mathcal{H}(D)) = \frac{g(\mathcal{H}(D))}{\alpha V +\beta E }.$
642	Finally, if $\mathcal{R}(T)$ denotes the reduction of the tree T in width, if $\mathcal{H}(D)$

Finally, if $\mathcal{R}(T)$ denotes the reduction of the tree T in width, if $\mathcal{H}(D)$ denotes the compression of a DAG D in height then the compression $\mathcal{R}^{*}(T)$ of the tree T in width and height can be computed in time $O(|T|^2 \deg(T) h(T))$ such as:

$$\mathcal{R}^{*}(T) = \mathcal{H} \circ \mathcal{R}(T).$$

646 6. Application to the compression of plant structures

Most plants have a modular branching structure made up of the repetition of basic modules such as leaves, stem portion between two nodes, shoots, etc. (Bell (1991), Godin and Caraglio (1998)). These highly repetitive structures are therefore well suited to assess our compression schemes. In this section, we will investigate how different types of plant structures can give rise to different types of compression in height.

653 6.1. Dichotomic Structures

Plants may build up dichotomic branching structures due to either the subdivision of their apices or to sympodial branching (the main apex stop to grow and two lateral apices resume the growth). Many plants show such a type of organization including one of the initial form of the earth's plant life, that is the *Cooksonia Caledonica* plant (Boyce (2008)) (Fig.16.a.i), and the *Horneophyton lignieri* (Eggert (1974)) (Fig.16.a.ii).



Figure 16: a. Examples of vegetal dichotomic structures: i. The *Cooksonia Caledonica* shape (Tandart et al. (2014)), ii. *Horneophyton* tree (Tandart et al. (2014)), b.i. a binary tree graph T, ii. the *DAG* D of its width reduction, iii. its height reduction graph $\mathcal{H}(D)$

Formally, the general pattern of a dichotomic structure may be represented by a binary tree graph T (Fig.16.b.i). If n denotes the height of Tthen the $DAG \ D = \mathcal{R}(T)$ represents its width reduction (Fig.16.b.ii). Using our algorithm (see Property 16) this DAG can be further reduced in height in time $O(n^2)$ leading to a DAG with nested returns $\mathcal{H}(D)$ (Fig.16.b.iii) with a reduction factor of

$$f(\mathcal{H}(D)) = 1 - \frac{2\alpha' + 2\beta'}{\alpha n + \beta(n-1)}$$

where α , β are the size of the computational representations of respectively a vertex and of an edge in a *DAG D*, and α' , β' are the size of the computational representations of respectively a vertex and of an edge in $\mathcal{H}(D)$. These notations will be used all through this section.

6.2. Structures with multi-scale periodicity

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Fish bone-like structures present a typical example of structure of multi-672 scale periodicity. They can be noticed, for example, in axial trees whose 673 growth is *monopodial* (i.e. trees having a growing main stem that keeps on 674 growing while producing lateral branches). Such a structure is thus composed 675 of a principal axis on which several elements are repeated (branches, leaves, 676 fruits,...). The *Date Palm* illustrates the wide variety of plants or plant parts 677 showing a fish bone-like structure (Fig.17.a.i). It can be abstracted as the 678 graph T represented in Fig.17.a.ii, where the main axis represents the trunk of 679 the palm tree and the lateral branches represents the composed leaves. This 680 structure can be compressed in width leading to the graph D of Fig.17.a.iii 681 that can itself be further compressed in height as $\mathcal{H}(D)$ (Fig.17.a.iv) with 682 height compression factor of 683



Figure 17: a.i. A model of the date palm (*Phoenix Dactylifera*) (Rhouma (1994)) and its lateral leaf modeled using the L-studio/Vlab (Prusinkiewicz et al. (2000); Abelson and DiSessa (1981)), ii. graphical representation T of the date palm. iii. the corresponding DAG D, iv. its height reduced graph $\mathcal{H}(D)$, b.i. Pattern of a fish bone-like structure T, ii. its width reduction DAG D, iii. its height reduced $DAG \mathcal{H}(D)$

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$$f(\mathcal{H}(D)) = 1 - \frac{6\alpha' + 9\beta'}{\alpha(n+m+2) + \beta(2(n+m)-1)}$$

where 2n is the number of leaves on each lateral branch, and m the number of all lateral branches.

More generally, a fish bone-like structure can be compressed in height both on the lateral branches $(j_1, ..., j_m)$ and on the main axis $(i_1, ... i_n)$, Fig.17.b.i. The width reduction of T produces the *DAG* D (Fig.17.b.ii). Then the height reduction $\mathcal{H}(D)$ of D (Fig.17.b.iii) is computed in time $O((n+m)^3)$ with a height compression factor

$$f(\mathcal{H}(D)) = 1 - \frac{5\alpha' + 7\beta'}{\alpha(n+m+1) + \beta(2n+m-1)}.$$

⁶⁹³ Note the two cycles of $\mathcal{H}(D)$ express respectively the repetition of motifs ⁶⁹⁴ on lateral branches and the repetition of motifs on the main axis.

695 6.3. Structures with nested periodicity

The structures introduced above correspond to structures with a multiscale periodicity, which is characterized by two levels of repeated patterns. In more complex multi-scale structures there may be several types of repeated



Figure 18: a.i. *Callistemon* branch (Rosser (2014); Nicholson (2014)), ii. Pattern of the corresponding DAG D, iii. its height reduced $DAG \mathcal{H}(D)$. Vertex in green represents a tree leaf, and pink vertex a fruit, b.i. A real fern of *Filicopsida Class* in the *Pteridophyta division* (Smith et al. (2006); Esculier and Ferreol (2014)), ii. Fractal fern illustrated using the OpenAlea Lab environment (Pradal et al. (2007)), iii. the associated DAG D, iv. its height reduced $DAG \mathcal{H}(D)$

patterns (leaves, flowers, twigs,...) that may alternate leading to nested peri-699 odicity. The structure of the *Callistemon* branch (Stead and Butler (1983)) 700 (Fig.18.a.i), in which groups of n lateral branches alternate with groups of 701 m fruits (k times), illustrates such a phenomenon. The DAG correspond-702 ing to this structure can be represented in Fig.18.a.ii where the vertex in 703 green represents the class of tree leaves, and the pink vertex the class of the 704 fruits. The height reduction of the Callistemon branch $\mathcal{H}(D)$ (Fig.18.a.iii) is 705 computed in time $O(k^3(n+m)^3)$ with a height reduction factor 706

707
$$f(\mathcal{H}(D)) = 1 - \frac{7\alpha' + 11\beta'}{\alpha(k(n+m)+3) + 2\beta(k(n+m)+2)}.$$

⁷⁰⁸ Note that the nested returns in $\mathcal{H}(D)$ express the nested periodicity, ⁷⁰⁹ where the outer loop corresponds to the repeated sequence (leaves, fruits), ⁷¹⁰ and the inner loops represent respectively the leaves and the fruits repetitions.

711 6.4. Self-similar structures

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Self-similarity is a conspicuous feature of many plants (Prusinkiewicz 712 (2004)), that can be exploited to compress the plant structures. Fern pro-713 vides a typical example of such a self-similar structure (Fig.18.b.i). A model 714 of this fern can be constructed in L-system (Prusinkiewicz et al. (1995)) -715 using a unique production rule (see section 17 of the Online supplementary 716 material)- as represented in Fig.18.b.ii. The width reduction of a fern model 717 of height n produces the DAG of Fig.18.b.iii, and the height reduction of D 718 produces $\mathcal{H}(D)$ (Fig.18.b.iv) in time $O(n^3)$, with a height reduction factor 719

$$f(\mathcal{H}(D)) = 1 - \frac{3\alpha' + 3\beta'}{\alpha(n+1) + \beta n}$$

For n = 30 the reduction factor is close to 93% showing the remarkable height compressibility of self-similar structures.

A general model of self-similar branching structures has been proposed 723 by Godin and Ferraro (2010), called self-nested trees. To study and assess 724 the quality of width compression algorithm the authors defined a database 725 of self-nested branching structures made up by four families of purely self-726 nested trees (Fig.19.a and .b) together with versions of this trees altered with 727 different levels of noise, Fig. 19.c. In each family M_i (i = 0..3), starting from 728 a self-nested template plant T_i , several noisy versions were created by varying 729 the lateral branching probability (Fig.19.c). 730

⁷³¹ Width and height compression were then applied on each tree T_i and all ⁷³² there noisy versions. Fig.20.a illustrates the width reduction graphs of some ⁷³³ specimens of the M_3 family, and for which the height compression provides ⁷³⁴ the graphs of Fig.20.b. For the four families, the obtained height compression ⁷³⁵ factors depending on the branching probability (for α and β fixed to one unit) ⁷³⁶ is shown in Fig.21.

⁷³⁷ One can notice that the *monopodial* template plants $(T_0, T_1 \text{ and } T_2)$ ⁷³⁸ are weakly height compressible structures, whereas the *sympodial* tree T_3 is ⁷³⁹ remarkably height compressible. However as soon as noise is introduced in ⁷⁴⁰ this tree, its height compressibility becomes weaker.

Overall, the height compressibility of the four families is low for small noise factor (probability ≥ 0.6). Nevertheless for increasing noise (probability < 0.6) height compressibility increases as the tree tends to become a linear structure.



Figure 19: a. (i) The differentiation graph of a non-branching plant structure. (ii) The resulting axis structure, where component colors correspond to the differentiation graph states in which these components were created. The numbers attached to each loop indicate the number of steps a meristem stays in the corresponding state. (iii) Differentiation graphs used for the definition of the theoretical plants T_i (i = 0..3). Solid arrows correspond to possible transitions of the apical meristem states. Dashed arrows correspond to possible transitions from the apical meristem state to the axillary meristem states. b. the self-nested template plants T_i (i = 0..3), c. the noisy versions of the template plant T_1 with probability 0.8, 0.6, 0.4 and 0.0



Figure 20: a. The width reduction graphs of the template plant T_3 and its noisy versions $T_{3,8}$, $T_{3,6}$, $T_{3,4}$ and $T_{3,2}$ ($T_{3,i}$ is the noisy tree of branching probability 0, i). b. the corresponding height reduced graphs



Figure 21: Height compression factor $f(\mathcal{H}(\mathcal{R}(T_i)))$ depending on the branching probability

745 7. Conclusion

This paper has considered the problem of height reduction from tree struc-746 tures. This study was designed to complement the work previously done in 747 the context of lossless compression of unordered trees (Godin and Ferraro 748 (2010)), where the trees were compressed in width and not in height. Us-749 ing the property that some tree structures exhibit height regularities, we 750 proposed to represent differently these special structures, characterized by 751 a height repeated pattern. These repetitions are associated with quasi-752 isomorphic sub-trees which are regularly organized along one main branch. 753

Starting from a reduction graph of a given tree, the height reduction consists of searching of all collections of regularly height repeated patterns. Each repeated pattern is replaced by a unique instance of this pattern augmented with a loop in the compressed tree which is called a Reduction Graph with Nested Returns. The efficiency of the compression algorithm is quantified by a compression factor reflecting the ratio between the size of the *DAG* with nested returns and the size of the initial *DAG*.

The method was then illustrated on different type of plant structures and showed that in many cases height compression can add a significant compression factor to the width compression, in particular on self-similar plants.

765 Acknowledgments

The authors are grateful to the University Djillali Liabes of Sidi-Bel-Abbes (Algeria) and to the INRIA-CIRAD-INRA Virtual Plants projectteam of Montpellier (France) who funded this research.

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