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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. Introduction

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

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

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

38 to purely geometrical transformations. In particular, they don't take into
39 account topology of the studied biological structures.

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

45 One particularly important simplifying hypothesis consists of focusing on
46 the result of the developmental process without considering the intermediate
47 stages of development. This leads to a whole class of approaches in which
48 models are constructed to capture plant branching architecture in a minimal
49 way. A first approach was proposed by Viennot et al. (1989) based on a
50 statistical analysis of binary tree organization. Subtrees were characterized
51 by an integer representing their amount of asymmetry (the so called Holton-
52 Strahler -HS- number (Deussen and Lintermann (2002))). Then parameters
53 were inferred from data to describe statistically the probability of finding a
54 tree with HS number n knowing the HS number m of the parent tree. A
55 similar statistical but more general approach was later used by Durand et al.
56 (2005, 2007), based on Markovian processes to describe subtrees frequencies
57 at any given node. These inference approaches are aimed to capture the
58 statistical properties of tree organization. Viewed as compression techniques,
59 they provide approximated strategies to compress trees, relying on different
60 types of *a priori* assumptions (e.g. Markovian property). An exact and
61 deterministic approach was recently introduced in Godin and Ferraro (2010).

62 In this latter work, the authors addressed the problem of recognizing sim-
63 ilar, possibly nested, patterns in plant structures, and to compress them as
64 directed acyclic graphs (*DAGs*) in a reversible way. Trees were represented
65 as unordered labeled tree structures, where no order is assumed on the chil-
66 dren of any node of the trees. Trees whose compression are linear *DAGs*
67 define the class of self-nested trees, i.e. trees that have very high compress-
68 ibility. From this, a degree of self-nestedness could be defined for any tree
69 as the normalized edit-distance of this tree to the class of self-nested trees
70 embedding the original tree, and a polynomial-time algorithm was designed
71 to compute this distance.

72 The tree compressions techniques defined in Godin and Ferraro (2010)
73 are based on the merging of isomorphic subtrees. Any two subtrees that are
74 merged therefore result in a structure with the same height. As a result, the
75 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 compression occurs in height. However, some trees such as high dichotomic trees or fish bone-like trees have a repetitive structure in height and are therefore poorly compressed by this technique. In this paper, we consider the problem of maximally compressing trees both in width and height with no loss of information. This idea is illustrated in Fig.1. In section 3, we recall the definition of tree reduction and self-nestedness introduced in Godin and Ferraro (2010). We then introduce and study in section 4 a weak extension of tree isomorphism, called quasi-isomorphism, that will make it possible to identify similar (but not identical) tree patterns at different heights of a tree. Based on these definitions and their properties, we present in section 5 an algorithm able to compress in height the reduction of any tree. This algorithm is then applied to both theoretical tree-like structures and vegetal branching systems to characterize its compression ability.

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

2. Notational conventions

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

A *directed graph*, is a pair $G = (V, E)$ where all edges correspond to ordered pairs of vertices. Let (x, y) be an edge in E , x is called a *parent*

of y and y is a *child* of x . A vertex that has no child is called a *leaf*. We will denote, in the sequel, by $child(x)$ the set of all the children of x , and $parent(x)$ the set of all parents vertices of x . In a directed graph G , the *Indegree* of a vertex v , denoted $deg^-(v)$, is the number of its parents, and its *Outdegree*, denoted $deg^+(v)$, is the number of its children. The Indegree and the Outdegree of G are respectively the maximum Indegree and the 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*
 $P_1 \cup P_2 = \{(x_1, x_2), \dots, (x_{N-1}, x_N), (y_1, y_2), \dots, (y_{M-1}, y_M)\}$
- 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
 - $P_1 \subset P_2$ if the sequence made by the subset of edges common to both P_1 and P_2 correspond to P_1 (path *inclusion*)
 - $P_1 \prec P_2 \Leftrightarrow x_N \prec y_1$
 - $P_1 \circ P_2 \Leftrightarrow x_N = y_1$ (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

137 the root, which has no parent vertex. In a tree, each vertex, different from
 138 the root, has exactly one parent vertex. In the following, a rooted tree is
 139 simply called a tree.

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

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

144 The height $h(x)$ of a vertex x in a *DAG* is the length of the longest path
 145 from x to a leaf. The height $h(D)$ of a *DAG* D is the height of its root
 146 vertex, and its width, denoted $l(D)$, is the maximum number of vertices of
 147 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".

154 Let us consider two trees, $T_1 = (V_1, E_1)$ and $T_2 = (V_2, E_2)$. A bijection ϕ
 155 from V_1 to V_2 is a *tree isomorphism* if for each $(x, y) \in E_1$, $(\phi(x), \phi(y)) \in E_2$.
 156 If there exists an isomorphism between T_1 and T_2 , the two structures are thus
 157 identical up to a relabeling of their components. In this case we say that T_1
 158 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

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

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

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

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).

Based on these definitions, the degree of self-nestedness of any tree T can be quantified by computing the distance between T and the smallest self-nested tree (noted *NEST*) that embeds it. This distance is null if the tree is in the class of self-nested tree and augments as the tree contains increasingly self-nested structures. Godin and Ferraro (2010) showed that this distance and the corresponding *NEST* of a tree T can be computed in polynomial time by a *DAG* linearization algorithm.

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

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

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

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.

4.2. Vertex and edge signatures

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

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

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: $V = \{A, B, C, D, E\}$ and $E = \{((A, B), 1), ((A, C), 2), ((A, D), 1), ((B, C), 1), ((B, E), 1), ((C, D), 1), ((C, E), 1), ((D, E), 1)\}$. The associated vertex signature matrix is given as follow:

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

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.

$\langle x, y \rangle$ 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 x and 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:

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

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

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

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

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

280 The edges signatures allow us to study the quasi-isomorphism relation in
 281 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,*
 285 *given two distinct atomic paths $\langle x, y \rangle, \langle x', y' \rangle$. We say that $\langle x, y \rangle$ and $\langle x', y' \rangle$*
 286 *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:

$$\begin{aligned}
 288 \quad \left. \begin{aligned} \sigma_{x,y} &= \sigma_{x',y'} \\ \sigma_{xy} &= \sigma_{x'y'} = 1 \end{aligned} \right\} & \xLeftrightarrow{\text{Def.4}} \sigma_{x|y} = \sigma_{x'|y'} \\
 & \xLeftrightarrow{\text{Def.5}} \langle x, y \rangle \cong \langle x', y' \rangle
 \end{aligned}$$

289 **B.** Given the set of edges signature computed in Example 3. There exist
 290 in this set, two equal edges signatures: $\sigma_{B|C} = \sigma_{C|D}$, which induce that
 291 $\langle B, C \rangle \cong \langle C, D \rangle$. This appears clearly in Fig.2.b.i in which the subtree
 292 associated with B except its child C , and the subtree associated with C
 293 except its child D are isomorphic.

294 **Property 3 (Fundamental property)** *Let e, f be two atomic paths start-*
 295 *ing from x and let e', f' be two atomic paths starting from x' , then*

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

296 In other terms, at most one pair of quasi-isomorphic atomic paths can start
 297 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 $P = \{e_1, \dots, e_n\}$, and $Q =$
 300 $\{f_1, \dots, f_m\}$.

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

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

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

307 **B.** In the DAG $\mathcal{R}(T_2)$ (Fig.5.a) they are two quasi-isomorphic paths: $P \cong P'$
 308 with $P = \{\langle B, E \rangle, \langle E, I \rangle\}$, $P' = \{\langle C, F \rangle, \langle F, H \rangle\}$ from $\sigma_{B|E} = \sigma_{C|F} =$
 309 $[000100000001]$, and $\sigma_{E|I} = \sigma_{F|H} = [000000100001]$.

310 **Notation:** A path $P = \{\langle x_1, x_2 \rangle, \dots, \langle x_{n-1}, x_n \rangle\}$ can be denoted as: $P =$
 311 $x_1 P x_n$ 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 $|P_i| = 1$
 312 we can note $P = x_1 P_1 x_i x_{i+1} P_n x_n$.

313 4.3.3. Quasi-periodic path

314 A quasi-periodic path is a path decomposable in strictly smaller sub-paths
 315 which are all quasi-isomorphic with each other, and which are not themselves
 316 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

317 **Definition 7 (quasi-periodic path QPP)** Given two vertices x, y with
 318 $x \prec y$. We say that xPy is a quasi-periodic path if there exist paths $P_1, \dots, P_n$
 319 ($P_i \neq \lambda$) and ($n \geq 2$) for which:

- 320 • $P = P_1 \cup \dots \cup P_n$
- 321 • $P_1 \cong \dots \cong P_n$
- 322 • $\forall i \leq n, P_i$ is not a quasi-periodic path.

323 P is called $QPP(x, y)$, and the sequence $P_1, \dots, P_n$ the *normal decomposition*
 324 of P .

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

326 **Example 6 A.** From the graph $\mathcal{R}(T_1)$ of Fig.2.b, the path $P = P_1 \cup P_2$ with
 327 $P_1 = \langle B, C \rangle, P_2 = \langle C, D \rangle$ is a QPP .

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

331 By definition, each QPP is a set of path chunks which are:

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

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

339 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.

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

344 Let $P_1, \dots, P_n$ be a normal decomposition of a *QPP* P , the path corre-
 345 sponding to a subsequence $P_i, \dots, P_{i+k}$ ($i < n, 1 < k < n - i$) of the normal
 346 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*
 349 *quasi-periodic path (noted *MQPP*) is a *QPP* maximum for path inclusion.*

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

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

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

357 4.3.5. Relative positions of maximal quasi-periodic paths in a reduction graph

358 Let us now study the relative positions of two *MQPPs* in a reduction
 359 graph. As for any two paths in a reduction graph, two *MQPPs* can either
 360 *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 $P = \{\langle A, B \rangle, \langle B, G \rangle, \langle G, E \rangle\}$ and $P' = \{\langle G, C \rangle, \langle C, D \rangle\}$ are intersecting 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.

C. In the reduction graph $\mathcal{R}(T_5)$ of Fig.6.c:

$$P_1 = \{\langle A, B \rangle, \langle B, C \rangle, \langle C, D \rangle\} \cup \{\langle D, E \rangle, \langle E, F \rangle, \langle F, I \rangle\}$$

$$P_2 = \{\langle A, B \rangle, \langle B, C \rangle\}$$

$$P_3 = \{\langle D, E \rangle, \langle E, F \rangle\}$$

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 $P_1^1, \dots, P_N^1$ the normal decomposition of P^1 . The motifs P_n^1 are all quasi-isomorphic 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 *MQPP*s and each *MQPP* is composed of a series of motifs.

379 can be decomposed into a list of quasi-isomorphic motifs denoted $P_{l,m}^{1,j}$ for
 380 $m = 1..M$, and so on (see Fig.7). These series of decomposition forms a tree
 381 structure called the *MQPP tree* of P^1 . Interestingly, this tree contains many
 382 identical subtrees as indicated by the following property.

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

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

386 **Definition 10 (Normal decomposition of a DAG)** *The normal decom-*
 387 *position of a DAG D is the set $\mathcal{N}$ of all its maximally nested MQPPs.*

388 Therefore, the normal decomposition of a *DAG* contains only top level nested
 389 *MQPP*s which are either simple or decomposable in to finer *MQPP*s.

390 In general, the relative position of two *MQPP*s must respect constraints
 391 as indicated by the following property.

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

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

- 395 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:*
 - 397 (a) *at least one of the MQPPs terminates through the intersection*
 - 398 *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

- 399 (b) *the intersection corresponds to a postfix of one of the two paths,*
400 (c) *or both MQPPs belong to the same path.*

401 **Example 8** Let us illustrate the different situations of two non nested inter-
402 secting *MQPPs* cited in Property 9. Fig.8.a shows some possible situations
403 of two non nested intersecting *MQPPs*, and Fig.8.b some impossible ones,
404 so in:

- 405 • Figs.8.a.i, 8.a.ii, 8.a.iii and 8.a.iv at least one of the *MQPPs* terminates
406 through the intersection point,
- 407 • Fig.8.a.v the intersection corresponds to a postfix of one of the two
408 paths,
- 409 • Fig.8.a.vi both *MQPPs* belong to the same path,
- 410 • Figs.8.b.i, 8.b.ii and 8.b.iii there is no *MQPP* which terminates through
411 the intersection point.

Figure 9: a.i. A height repeated branch, ii. the corresponding $MQPP = P_1 \cup \dots \cup P_n$ 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

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) $P_1, \dots, P_n$ (as in Fig.9.a.ii). We will make use of this repeated pattern to compress these $MQPP$ s. This will be made by allowing introduction of loops in the original DAG s, as in Fig.9.a.iii. Let us introduce the notion of DAG with return edges.

5.1. DAGs with Return Edges

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

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

- $\forall (x, y) \in E', y \prec x$ in D .

- 424 • As for any edge of D , we associate a weight $n(x, y)$ with each edge
425 $(x, y) \in E'$.
- 426 • In addition we associate with each edge $(x, y) \in E'$ one of the children
427 z of x if any, such that $n(x, y) = n(x, z)$, z is called the exit vertex of
428 the cycle induced by the return edge (x, y) and denoted $ex(x, y)$.

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

432 **Property 10** Let $B = (e_1, \dots, e_n)$ an elementary cycle in a DAG with return
433 edges $D = (V, E, E')$. There is a unique return edge $e_k \in E'$ in B denoted
434 $re(B)$. We called the path $\{e_{k+1}, \dots, e_n, e_1, \dots, e_{k-1}\}$ the principal path of B
435 denoted $pp(B)$, and the vertex $ex(e_k) = z$ the exit vertex of the cycle B ,
436 denoted $ex(B)$.

437 Furthermore, we say that $x \in pp(B)$, for each vertex x in $pp(B)$. In the
438 DAG with return edges of Fig.10.e, for the elementary cycle $B_2 = ((x, y),$
439 $(y, z), (z, w), (w, x))$: $re(B) = (w, x)$, $pp(B) = \{(x, y), (y, z), (z, w)\}$ and
440 $ex(B) = e$.

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

447 5.2. DAG with nested returns

448 **Definition 12 (DAG with nested returns)** We define a DAG with nest-
449 ed returns as a DAG with return edges in which any two elementary cycles B_1
450 and B_2 (with respective vertex sets V_1 and V_2) are either nested ($V_1 \cap V_2 = V_1$
451 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 $\mathcal{D}_{\uparrow\uparrow} \subset \mathcal{D}_\uparrow$.

454 **Example 9** Fig.11.a shows examples of patterns of vertices of DAGs $\in \mathcal{D}_{\uparrow\uparrow}$
455 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 $x \notin pp(B)$.

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

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

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 $\mathcal{B}(x) = \{B_1, \dots, B_{b(x)}\}$, $b(x) = |\mathcal{B}(x)|$ and $B_1 \subset \dots \subset B_{b(x)}$.

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 $P_1, \dots, P_{\mu_{max}+1}$ 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.

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

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

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

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

504 analyze how *MQPPs* can be folded and then apply the resulting algorithm
 505 to a maximal set of *MQPPs* in the original *DAG*.

506 5.3.1. Compression of a *MQPP*

507 Let P be a *MQPP* in a *DAG* D and $P_1, \dots, P_n$ be its normal decomposi-
 508 tion. The idea is to replace the sequence $P_1, \dots, P_n$ by a loop over P_1 repeated
 509 n times. In practice the original *DAG* must be edited in the following way.

510 **General compression algorithm of a *MQPP*:** Let $D = (V, E, E')$ be a
 511 *DAG* of $\mathcal{D}_{\uparrow\uparrow}$, and let $P = P_1 \cup \dots \cup P_n$ the *MQPP*(x, y) with $P_i = x_{i,1}P_i x_{i,m}$
 512 ($1 \leq i \leq n$, $m \geq 2$), $x = x_{1,1}$ and $y = x_{n,m}$, as represented in Fig.13.a. We
 513 denote by B_P the cycle obtained from the compression of P , and the resulting
 514 *DAG* with nested returns by $D_{/P}$. Then $D_{/P}$ is computed in the following
 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 • this edge is also augmented with the information that when the
 520 loop ends, scanning should resume on the exit vertex y of $x_{n,m-1}$,
 521 labeled by χ .
- 522 2. Redirection of entering edges of the *MQPP*, Fig.13.c:
 - 523 • Replace each entering edge $(z, x_{i,j})$ of the *MQPP* by the equiva-
 524 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 • its multiplicity list corresponds to the multiplicity list of the edge
 527 $(z, x_{i,j})$ augmented by the multiplicity of the loop B_P for this edge,
 528 that is $n - i$.
- 529 3. Suppression of the *MQPP* repeated patterns, Fig.13.d:
 - 530 • Remove edges which belong to the paths $P_2, \dots, P_n$,
 - 531 • the last edge of P_1 ($x_{1,m-1}, x_{1,m}$) is redirected on the exit vertex
 532 χ . It is thus replaced by the edge $(x_{1,m-1}, \chi)$ whose weight is also
 533 ω ,
 - 534 • Edit the sets V , E and E' .

Figure 13: a. Formal pattern of a $MQPP$ $P = P_1 \cup \dots \cup P_n$, 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 $P_2, \dots, P_n$

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. $\mathcal{R}(T_1)_P$

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

536 **Example 10** Let us compress the MQPP $P = P_1 \cup P_2$ with $P_1 = \langle B, C \rangle$
 537 and $P_2 = \langle C, D \rangle$ of Fig.2.b. The compressed graph $\mathcal{R}(T_1)_P$ of Fig.14.c is
 538 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 $((A, B), 1, \{1\})$,
 543 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).

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

548 5.3.2. Gain of MQPP compression

549 Let us consider a MQPP P in $D \in \mathcal{D}_{\uparrow\uparrow}$, with $P = P_1 \cup \dots \cup P_n$, and
 550 $P_i = x_{i,1}P_i x_{i,m}$ ($1 \leq i \leq n$, $m \geq 2$). Let us denote $|\overline{V_P}|$ and $|\overline{E_P}|$ the
 551 number of vertices and respectively the number of edges removed from P
 552 compression. Formally

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

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

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

557 **Property 14 (compression gain of two MQPPs)** Given two paths
 558 $P, P' \in MQPP(D)$. If P and P' are either disjoint or nested, then the gain

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

559 Let P and P' two MQPPs in a DAG D . $D_{/P/P'}$ denotes the DAG with
 560 nested returns obtained by successive compressions of P and P' , and the gain
 561 $g(D_{/P/P'}) = g(\{P, P'\})$.

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

564 If two MQPPs are intersecting but no nested then the compression of one
 565 of them may impair or prevent the compression of the other. Therefore as
 566 the compression of the two MQPPs is mutually exclusive, the compression
 567 of the DAG will depend on the choice of the compression of either of them.
 568 We say that the DAG compression is ambiguous.
 569 Given a DAG D , let M be a subset of $MQPP(D)$.

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

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

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

$$577 \quad P \wedge P' \Leftrightarrow P \text{ and } P' \text{ are neither disjoint or nested.}$$

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 MQPP of 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 MQPPs unambiguous 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 $G_{M'_i}$ (i.e. a complete subgraph of G_{M_i}) with maximal gain. This optimization problem is known to be NP-complete (Feige et al. (1991)), and several approaches have been developed to solve it (Battiti and Protasi (2001); Bomze et al. (1999); Feige (2004)). Here we use the *TABU algorithm* (Gendreau et al. (1993)), as described in section 15 of the Online supplementary material. Given a class M_i the algorithm returns a maximal subset M'_i unambiguous for compression with maximal gain.

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

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

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 .

Definition 17 (Unambiguous normal decomposition of a DAG)
The unambiguous normal decomposition of D , $\mathcal{N}^$, is the set of all its maximally 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.

5.3.4. DAG compression evaluation:

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

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

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:

1. Compute the set of vertex signatures and then the set of edge signatures of D (see section 4.2).
2. Given the set of edge signatures compute the set $MQPP(D)$ (see section 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 *DAG* 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 $\mathcal{N}^*(\mathcal{H}) = \{P_1, \dots, P_n\}$ the unambiguous normal de-
637 composition of $\mathcal{H}$
638 ii. $\mathcal{H} = \mathcal{H}_{/P_1/\dots/P_n}$
639 until $\mathcal{N}^*(\mathcal{H}) = \emptyset$,
640 5. Return the *DAG* 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 $\mathcal{H}(D)$
643 denotes the compression of a *DAG* D in height then the compression $\mathcal{R}^*(T)$ of
644 the tree T in *width and height* can be computed in time $O(|T|^2 \deg(T) h(T))$
645 such as:

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

646 6. Application to the compression of plant structures

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

653 6.1. Dichotomic Structures

654 Plants may build up dichotomic branching structures due to either the
655 subdivision of their apices or to sympodial branching (the main apex stop
656 to grow and two lateral apices resume the growth). Many plants show such
657 a type of organization including one of the initial form of the earth's plant
658 life, that is the *Cooksonia Caledonica* plant (Boyce (2008)) (Fig.16.a.i), and
659 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 T then 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

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

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 $\mathcal{H}(D)$

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

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 ($j_1, \dots, j_m$) and on the main axis ($i_1, \dots, i_n$), 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

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

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

6.3. Structures with nested periodicity

The structures introduced above correspond to structures with a multi-scale 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 $\mathcal{H}(D)$

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

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

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

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

$$720 \quad f(\mathcal{H}(D)) = 1 - \frac{3\alpha' + 3\beta'}{\alpha(n+1) + \beta n}.$$

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

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

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 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.

741 Overall, the height compressibility of the four families is low for small
742 noise factor (probability ≥ 0.6). Nevertheless for increasing noise (probability
743 < 0.6) height compressibility increases as the tree tends to become a linear
744 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

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

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

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

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