The difficulty of constructing a leaf-labelled tree including or avoiding given subtrees

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Abstract

Given a set of trees with leaves labelled from a set L, is there a tree T with leaves labelled by L such that each of the given trees is homeomorphic to a subtree of T? This question is known to be NP-complete in general, but solvable in polynomial time if all the given trees have one label in common (equivalently, if the given trees are rooted). Here we show that this problem is NP-complete even if there are two labels x and y such that each given tree contains x or y. On the other hand, we show that the question of whether a fully resolved (binary) tree exists which has *no* subtree homeomorphic to one of the given ones is NP-complete, even when the given trees are rooted. This sheds some light on the complexity of determining whether a probability assignment to trees is coherent.

Keywords: phylogenetic trees, compatibility, NP-complete, probability, reconstruction.

1 Introduction and definitions

A *phylogenetic tree* on a label set L is a tree with no vertices of degree 2 and exactly $|L|$ leaves, each of which is labelled with a distinct element of L. Such trees are used to represent evolutionary relationships in biology.

Suppose that T is a phylogenetic tree on L and A is a subset of L. Consider the minimal subtree of T that connects leaves from A, and suppress all vertices of degree 2 (i.e. make the tree homeomorphically irreducible) to obtain a phylogenetic tree on A, denoted $T|_A$. If T' is a binary tree, we say T is *compatible* with T' if $T|_A = T'$ for some subset A of L. A set S of binary trees is said to be *consistent* if there exists a phylogenetic tree T'' that is compatible with all the trees in S. We then say T'' *realises* S.

A general problem considered in recent literature (Aho et al. [1], Ng and Wormald [3], Steel [6] and Constantinescu and Sankoff [2]) is to determine whether there exists a tree that realises S. This general problem has been shown to be NP-complete (see Steel [6]). On the other hand, it has been shown that the problem of determining whether a set of rooted trees is consistent can be solved in polynomial time (see Aho et al. [1]; Ng and Wormald [3] and Hensinger et al. [5] consider similar questions).

Suppose L_0 is a subset of L such that every input tree has at least one label in L_0. The question we consider in this paper is: what is the complexity of the consistency problem if $|L_0|$ is fixed, that is, independent of $n = |L|$? This question was posed by Steel [6]. If $|L_0| = 1$, then the input trees can be considered as rooted trees, and so consistency can be determined in polynomial time. We shall show that the problem is NP-complete for $|L_0| = 2$. It then follows trivially that the problem is also NP-complete for any fixed value of $|L_0| \geq 2$. The proof used in [6] for the general case does not extend to the case when $|L_0| = 2$. It is interesting to note that the proof here is simpler, even though the result is stronger.

In Section 3 we consider a related problem: whether a given probabilistic distribution of subtrees can be generated in a natural way by a model of a random tree.

We finish this section with some additional definitions. A vertex of a phylogenetic tree that has degree greater than 1 is said to be *internal.* A *binary (phylogenetic) tree* is one with all internal vertices having degree 3. A *quartet* is a binary phylogenetic tree on a label set of size 4. We denote a quartet on the label set $\{a, b, c, d\}$ by $ab|cd$, if a and b are the labels of two closest leaves, as shown in Figure 1(a). A *caterpillar* is a binary tree that has at most two vertices that are each adjacent to precisely two leaves. If x and y label two leaves that are maximally far apart on a caterpillar, we shall call it an xy-caterpillar. We write $xa_1|a_2 \ldots a_{m-1}|a_m y$ to denote the xy-caterpillar shown in Figure 1(b). We note that, for each pair, x, y, there is a one-to-one correspondence between the xy-caterpillars on the label set L and linear orderings on the set $L \setminus \{x, y\}$. Note that, if T is an xy-caterpillar, and $A \subseteq L$ with $x, y \in A$, then $T|_A$ is also an xy-caterpillar.

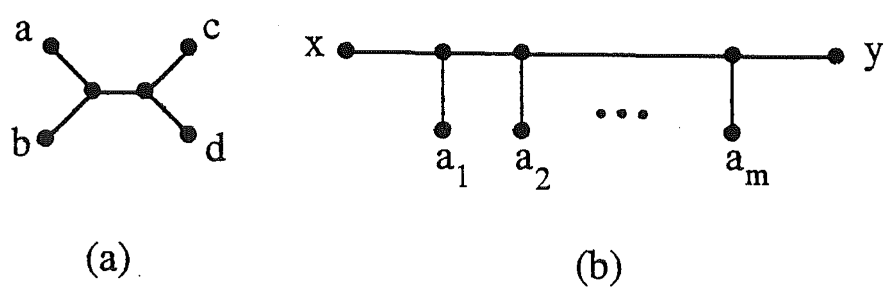

Figure 1. A quartet and a caterpillar.

2 Complexity of the problem for two roots

We henceforth consider the case where the input trees are all quartets. Suppose Q is a set of quartets and L_0 is a set of labels such that each quartet in Q has at least one label in L_0. We shall show that, if $|L_0| = 2$, then the problem of deciding whether Q is consistent is NP-complete.

Lemma 1. *Let $Q = \{x\theta|by, xb|\psi y, xa|c\theta, yc|a\psi\}$. Then the only trees on the label set $\{x, y, a, b, c, \theta, \psi\}$ that realise Q are the two caterpillars $xa|\theta b\psi|cy$ and $c\theta|xby|a\psi$ shown in Figure 2.*

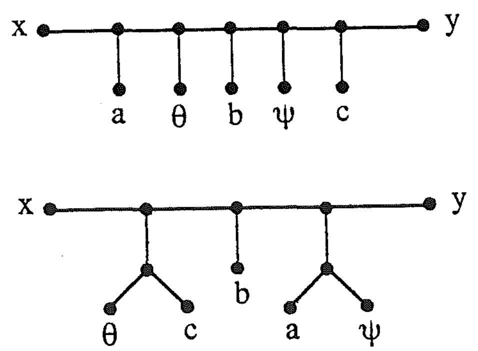

Figure 2.

Proof. There is only one tree on the label set $\{x, y, b, \theta, \psi\}$ that is compatible with the quartets $x\theta|by$ and $xb|\psi y$, namely, the caterpillar $x\theta|b|\psi y$. If the leaves a and c are added to this tree in such a way as to be compatible with $xa|c\theta$, then since x and θ are adjacent leaves, either a must be added right next to x, or c next to θ. Similarly, considering $yc|a\psi$, we need c next to y or a next to ψ. Hence there are exactly two possibilities: c can be added next to y and a next to x (giving the first tree), or c next to θ and a next to ψ (giving the second). ∎

¿From [6, Section 4], we have the following result.

Lemma 2. If a set of xy-caterpillars is consistent, then there exists an xy-caterpillar that realises the set.

The topic of this section is the following decision problem.

TWO-ROOTED QUARTET CONSISTENCY

INSTANCE: A set Q of quartets each of which includes a leaf labelled by x or by y.

QUESTION: Is Q consistent?

Theorem 1. *Two-rooted quartet consistency is NP-complete.*

Proof. The problem is clearly in NP, for, given a tree T that realises Q this consistency can be verified by checking each quartet in Q against T, and this checking can be done in polynomial time. We next describe a transformation from the following problem, which is NP-complete (Garey and Johnson [4]).

BETWEENNESS

INSTANCE: A finite set A and a collection I of ordered triples (a, b, c) of distinct elements from A (we may assume that each element of A occurs in at least one triple from I).

QUESTION: Is there a betweenness ordering f of A for I, that is, a one-to-one function $f : A \longrightarrow \{1, 2, \ldots, |A|\}$ such that for each $(a, b, c) \in I$, either $f(a) < f(b) < f(c)$ or $f(c) < f(b) < f(a)$?

Given an instance $I = \{(a_i, b_i, c_i); i = 1, \ldots, k\}$ of betweenness, we let a_i, b_i, c_i, θ_i and ψ_i $(i = 1, \ldots, k)$ be $5k$ labels, x and y two other labels, $Q_i = \{x\theta_i|b_iy, xb_i|\psi_iy, xa_i|c_i\theta_i, yc_i|a_i\psi_i\}$ and $Q(I) = \cup_{i=1}^k Q_i$. We note that each quartet in $Q(I)$ has a leaf labelled by x or by y. Clearly, the transformation can be done in polynomial time. We shall now show that $Q(I)$ is consistent if and only if I allows a betweenness ordering on the set $A = \cup_{i=1}^k \{a_i, b_i, c_i\}$.

Suppose that $Q(I)$ is consistent and T is a tree that realises $Q(I)$. Consider, for each $i, t_i := T|_{\{x,y,a_i,b_i,c_i\}}$. By Lemma 1, t_i is $xa_i|b_i|c_iy$ or $xc_i|b_i|a_iy$. Now the set $S = \{t_i; i = 1, \ldots, k\}$ is consistent since it is realised by T. By Lemma 2, there exists an xy-caterpillar T' which realises S. Then the order of the labels in A along T' provides the required betweenness ordering of A for I, since the label set of T' is $A \cup \{x, y\}$.

Conversely, suppose I allows a betweenness ordering on A. Let T' be one of the associated xy-caterpillars, obtained by ordering the labels in A along T' according to the betweenness ordering. We need to attach $2k$ additional labels $\{\theta_i, \psi_i; i = 1, \ldots, k\}$ to T' to obtain a tree compatible with $Q(I)$. For $i = 1, \ldots, k$, proceed as follows: If $T'|_{\{x,y,a_i,b_i,c_i\}} = xa_i|b_i|c_iy$, then attach θ_i and ψ_i to the xy-path of the tree so far constructed so that θ_i is between a_i and b_i, and ψ_i is between b_i and c_i. The resultant tree restricted to $\{x, y, a_i, b_i, c_i, \theta_i, \psi_i\}$ will be the caterpillar $xa_i|\theta_i b_i \psi_i|c_iy$. On the other hand, if $T'|_{\{x,y,a_i,b_i,c_i\}} = xc_i|b_i|a_iy$, then attach θ_i to the edge incident with the leaf labelled c_i, and attach ψ_i to the edge incident with the leaf labelled a_i. The resultant tree when restricted to $\{x, y, a_i, b_i, c_i, \theta_i, \psi_i\}$ will be the second tree specified in Lemma 1. In this way, we obtain a tree T which realises Q_i, for $i = 1, \ldots, k$ and hence realises $Q(I)$. ∎

Comments:

1. In the above proof, half of the quartets in $Q(I)$ have both labels x and y. One may ask the question: what is the complexity of the two-rooted quartet consistency problem, if no quartet in Q has both labels x and y? The answer is that it is still NP-complete, as we can replace each $x\theta_i|b_iy$ by two quartets $x\theta_i|b_i\alpha_i$ and $b_i\theta_i|\alpha_iy$ where α_i is a new label. These quartets imply the quartet $x\theta_i|b_iy$. A similar replacement can be done for $xb_i|\psi_iy$.
2. In defining the concepts of compatibility and consistency, we have confined them to the case when all the input trees are binary, as we intend to apply them only to input quartets. In general when the input trees are phylogentic trees, two different types of compatibility can be defined. Suppose T and T' are phylogenetic trees, we say T is *compatible* with T' if $T|_A = T'$ for some subset A of the label set of T. This definition is used by Ng and Wormald [3]. We say T is *weakly compatible* with T' if T' can be obtained from $T|_A$ by contracting certain edges. This definition was used by Steel [6]. These two definitions coincide when T' is a binary tree. The above theorem shows that the general consistency problem is NP-complete for both types of compatibility.

3 Forbidding subtrees

In this section, we consider the complexity of constructing a tree that is not compatible with any of a given set of subtrees. It will be shown that the problem is NP-complete even for rooted trees.

We consider the following decision problem.

FORBIDDEN SUBTREES

INSTANCE: A collection S of rooted binary trees whose leaf sets are subsets of a label set L.

QUESTION: Is there a leaf-labelled rooted binary tree T with label set L having no subtree containing the root homeomorphic to a tree in S?

Our main result in this section is the following.

Theorem 2. *The decision problem FORBIDDEN SUBTREES is NP-complete.*

Comments:

1. If we drop the word "binary" from both the instance and the question, then the resulting problem is still NP-complete, by polynomial transformation from FORBIDDEN SUBTREES. This is because the output tree in FORBIDDEN SUBTREES can be forced to be binary by including appropriate trees in the input which forbid all vertices of degree at least 4.

2. A more general problem can be formulated as follows. Suppose we are given a function $f : S \to [0,1]$. Then f may be considered as a measure of "confidence" or "probability" of the subtrees in S. We wish to know whether f "lifts" to a probability distribution on the set $R(L)$ of all rooted binary trees with leaf set L. That is, is there a function $\hat{f} : R(L) \to [0,1]$ with $\sum_{T \in R(L)} \hat{f}(T) = 1$ and such that, for all $t \in S$, $f(t)$ is the sum of $\hat{f}(T)$ over all T in $R(L)$ that are compatible with t. If $\hat{f}$ exists, then our beliefs represented by f are "coherent"; otherwise, not. The special case that $f(t) = 0$ for all $t \in S$ has answer yes if and only if FORBIDDEN SUBTREES has answer yes and is therefore NP-hard. On the other hand, the special case that $f(t) = 1$ for all $t \in S$ has answer yes if and only if S is consistent, so there is a polynomial time algorithm for this special case (Aho et al. [1], Ng and Wormald [3], Henzinger et al. [5]).

Proof of Theorem 2 Consider an instance I of BETWEENNESS and let A be the set of all labels i, j, k with $(i,j,k) \in I$. Let $L = A \cup \{z\}$

where $z \notin A$, and construct S as the union of the following four sets. Here $a(bc)$ denotes the rooted tree with b and c on one branch at the root and a on the other, and $a(b(cd))$ denotes the rooted tree with $b(cd)$ on one branch at the root and a on the other.

$$\begin{aligned} S_1 &= \{z(xy) : x, y \in A\} \\ S_2 &= \{j(ik) : (i,j,k) \in I\} \\ S_3 &= \{i(k(jz)) : (i,j,k) \in I\} \\ S_4 &= \{k(i(jz)) : (i,j,k) \in I\} \end{aligned}$$

Suppose that there is a leaf-labelled tree T as required in FORBIDDEN SUBTREES. Then the absence of the subtrees in S_1 forces T to be a caterpillar, with the leaves having labels from A to be attached along the path from the root to z in some linear order. Next, forbidding the rest of S forces the ordering to be a betweenness ordering for I. Conversely, if I has a betweenness ordering, then that ordering gives a permissible ordering of those leaves along the path from the root to z. The transformation implicitly described here takes polynomial time, and therefore FORBIDDEN SUBTREES is NP-complete. ∎

Open problem

In view of the second comment after Theorem 2, we ask for the complexity of the following problem, where c denotes a pre-chosen "confidence level", $0 < c \leq 1$:

c-EXPECTED SUBTREES

INSTANCE: A collection S of rooted binary trees whose leaf sets are subsets of a label set L and a function $f : S \to [0,1]$ with $f(t) \geq c$ for all $t \in S$.

QUESTION: Does f "lift" to a probability distribution on the set $R(L)$ of all rooted trees labelled from L?

Of course, for $c = 1$ this has a polynomial time algorithm.

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