# On the Complexity of Inferring Rooted Evolutionary Trees

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## Abstract

We prove that the maximum inferred local consensus tree problem is NP-complete, thus resolving an open question from [3].

### 1 Introduction

An evolutionary tree is an unordered tree whose leaves are in one-to-one correspondence with a set of species. Internal nodes represent common ancestors, and the branching structure reflects the species' evolutionary relationships. In some settings, the data does not uniquely determine a root, which leads to unrooted (as opposed to rooted) trees.

Traditional methods for evolutionary tree construction can be divided into character-based, distance-based, and maximum likelihood methods [7,8,10]. Recently, methods for inferring evolutionary history from information concerning local topological relationships between the species have been proposed and analyzed. Among these are quartet methods [5,9] which first compute unrooted topologies for subsets of cardinality four of the species and then combine them to form an unrooted evolutionary tree, and rooted consensus methods [1,3,4,6].

Aho et al. [1] studied the problem of inferring a rooted tree from a set of constraints on lowest common ancestor relations. They showed how to decide whether an instance admits a solution that is consistent with all of the constraints, and if so, how to construct it, in  $O(mn \log m)$  time, where m is the number of constraints and n the number of species. An even faster implementation restricted to constraints in the form of rooted, unordered, binary trees on three species was later given by Henzinger et al. [4]. However, data obtained experimentally often contains errors, implying that there usually will not exist a tree consistent with all of the constraints. A single erroneous constraint in the input might result in these algorithms returning the null tree. Therefore, [3] introduced optimization versions of the problem, called MICT and MILCT. [3] proved that MICT is NP-complete and proposed some approximation algorithms for MICT and MILCT, but left the computational complexity of MILCT as an open problem. In this paper, we prove that MILCT is NP-complete, obtaining a shorter NP-completeness proof for MICT in the process.

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### 2 Problem Definitions and Results

Let S be a set of elements. An LCA constraint on S is a constraint of the form  $\{i,j\} < \{k,l\}$ , where  $i,j,k,l \in S$ , which specifies that the lowest common ancestor of i and j is a proper descendant of the lowest common ancestor of k and k. An LCA constraint of the form  $\{i,j\} < \{i,k\}$  is called a 3-leaf constraint on S; it uniquely determines the relative topology of i,j, and k, and is written as  $(\{i,j\},k)$  for short. A rooted tree with leaves distinctly labeled by elements in S and an LCA constraint on S which is satisfied in the tree are consistent with each other.

The maximum inferred consensus tree problem is to construct a rooted tree that is consistent with as many LCA constraints as possible from a given set. The corresponding decision problem is:

## The maximum inferred consensus tree problem (MICT)

**Instance:** Finite set U, set V of LCA constraints on U, positive integer  $L \leq |V|$ .

**Question:** Is there a rooted tree that is consistent with L of the constraints in V?

A special case of MICT occurs when each constraint is a 3-leaf constraint:

## The maximum inferred local consensus tree problem (MILCT)

**Instance:** Finite set S, set T of 3-leaf constraints on S, positive integer  $K \leq |T|$ .

**Question:** Is there a rooted tree that is consistent with K of the constraints in T?

To determine the computational complexity of MILCT, it will be useful to know that the following problem is NP-complete (problem [MS2] in [2]):

#### Cyclic Ordering

**Instance:** Finite set A, collection C of ordered triples (a, b, c) of distinct elements from A.

**Question:** Is there a one-to-one function  $f: A \to \{1, 2, ..., |A|\}$  such that, for each  $(a, b, c) \in A$ , we have either f(a) < f(b) < f(c), or f(b) < f(c) < f(a), or f(c) < f(a)?

We are now ready to prove the main result.

Theorem 1 MILCT is NP-complete.

**PROOF.** MILCT is in NP since verifying if there exists a rooted tree that is consistent with a given subset of T can be done in polynomial time with the algorithm of Aho *et al.* [1].

To show the NP-hardness of MILCT, we give a polynomial-time reduction from Cyclic Ordering. Given an instance (A, C) of Cyclic Ordering, let S =

 $A \cup \{x_0, x_1, x_2, ..., x_{|C|}\}$  and  $K = \frac{|A| \cdot (|A|-1)}{2} + 2 \cdot |C|$ . For each  $a, b \in A$  with  $a \neq b$ , include the two constraints  $(\{x_0, a\}, b)$  and  $(\{x_0, b\}, a)$  in T. Next, for every i in  $\{1, 2, ..., |C|\}$ , add to T the three constraints  $(\{x_i, a\}, b), (\{x_i, b\}, c)$ , and  $(\{x_i, c\}, a)$ , where (a, b, c) is the ith ordered triple in C. Note that at most one of  $(\{x_0, a\}, b)$  and  $(\{x_0, b\}, a)$  and at most two of  $(\{x_i, a\}, b), (\{x_i, b\}, c)$ , and  $(\{x_i, c\}, a)$  can be consistent with any rooted tree, so the number of constraints in T that can be satisfied must be  $\leq K$ .

Claim: (A, C) has a cyclic ordering if and only if there exists a rooted tree that is consistent with K of the constraints in T.

Proof of claim: Suppose the answer to the Cyclic Ordering instance is yes. Then there exists a one-to-one function  $f: A \to \{1, 2, ..., |A|\}$  such that, for each ordered triple  $(a, b, c) \in C$ , we have either f(a) < f(b) < f(c), or f(b) < f(a), or f(c) < f(a) < f(b). We can construct a rooted tree consistent with K constraints as in Fig. 1.

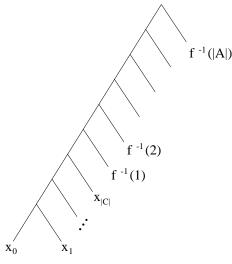


Fig. 1.

If  $f(\alpha_i) < f(\beta_i) < f(\gamma_i)$  for the *i*th ordered triple in C, then  $(\{x_i, \alpha_i\}, \beta_i)$  and  $(\{x_i, \beta_i\}, \gamma_i)$  are consistent with the tree. Also, for each pair  $a, b \in A$  with  $a \neq b$ , exactly one of  $(\{x_0, a\}, b)$  and  $(\{x_0, b\}, a)$  is consistent with the tree. Thus, the tree is consistent with  $2 \cdot |C| + \frac{|A| \cdot (|A|-1)}{2}$  of the constraints in T.

Conversely, suppose there exists a rooted tree R consistent with  $\frac{|A|\cdot(|A|-1)}{2}+2\cdot|C|$  of the constraints. At most  $\frac{|A|\cdot(|A|-1)}{2}$  constraints of type  $(\{x_0,a\},b)$  and at most  $2\cdot|C|$  constraints of type  $(\{x_i,a\},b)$  with  $i\neq 0$  can be consistent with R, so R must be consistent with precisely this many constraints of each type, respectively.  $\frac{|A|\cdot(|A|-1)}{2}$  constraints of the former type can only be satisfied if the subtree of R induced by  $A\cup\{x_0\}$  is a rooted caterpillar whose root is the parent of a leaf and an internal node, and one of the two leaves at maximum distance from the root is labeled  $x_0$  (otherwise, for some pair  $a,b\in A$ , neither

 $(\{x_0, a\}, b)$  nor  $(\{x_0, b\}, a)$  would be consistent with R). For each  $a \in A$ , let f(a) be the number of internal nodes on the path from a to  $x_0$  in the subtree of R induced by  $A \cup \{x_0\}$ . Next, because of the constraints of the second type, for every ordered triple  $(a, b, c) \in C$ , exactly two of the three corresponding constraints in T are consistent with R (if, for some ordered triple, just one constraint was consistent with R, then the number of constraints of this type consistent with R could not add up to  $2 \cdot |C|$ ). Therefore, either (1) a is closer to  $x_0$  than b is to  $x_0$  and b is closer to  $x_0$  than b is to b is closer to b than b is to b is closer to b than b is to b is closer to b than b than b than b to b than b than b than b than b to b than b to b than b t

## Corollary 2 MICT is NP-complete.

**PROOF.** MICT is in NP because the algorithm of Aho *et al.* [1] can check any given subset of the LCA constraints for consistency in polynomial time. MICT is NP-hard since it admits a direct reduction from MILCT; just replace each 3-leaf constraint  $(\{a,b\},c)$  in the given instance by  $\{a,b\} < \{a,c\}$ .

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