

The Longest Common Subsequence Problem with Crossing-Free Arc-Annotated Sequences

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Abstract. An arc-annotated sequence is a sequence, over a given alphabet, with additional structure described by a set of arcs, each arc joining a pair of positions in the sequence. As a natural extension of the longest common subsequence problem, Evans introduced the LONGEST ARC-PRESERVING COMMON SUBSEQUENCE (LAPCS) problem as a framework for studying the similarity of arc-annotated sequences. This problem has been studied extensively in the literature due to its potential application for RNA structure comparison, but also because it has a compact definition. In this paper, we focus on the nested case where no two arcs are allowed to cross because it is widely considered the most important variant in practice. Our contributions are three folds: (i) we revisit the nice NP-hardness proof of Lin et al. for LAPCS(NESTED, NESTED), (ii) we improve the running time of the FPT algorithm of Alber et al. from $O(3.31^{k_1+k_2}n)$ to $O(3^{k_1+k_2}n)$, where resp. k_1 and k_2 deletions from resp. the first and second sequence are needed to obtain an arc-preserving common subsequence, and (iii) we show that LAPCS(STEM, STEM) is NP-complete for constant alphabet size.

1 Introduction

Structure comparison for RNA has become a central computational problem bearing many computer science challenging questions. Indeed, RNA secondary structure comparison is essential for (i) identification of highly conserved structures during evolution (which cannot always be detected in the primary sequence, since it is often unpreserved) which suggest a significant common function for the studied RNA molecules, (ii) RNA classification of various species (phylogeny), (iii) RNA folding prediction by considering a set of already known secondary structures, and (iv) identification of a consensus structure and consequently of a common role for molecules. From an algorithmic point of view, RNA structure comparison was first considered in the framework of ordered trees [12] and, later on, in the one of *arc-annotated sequences* [5]. An *arc-annotated sequence* over some fixed alphabet Σ is a pair (S, P) , where S (the *sequence*) is a string of Σ^* and P (the *annotation*) is a set of arcs $\{(i, j) : 1 \leq i < j \leq |S|\}$. In the context of RNA structures, S is a sequence of RNA bases and P represents hydrogen bonds between pairs of elements of S . From a purely combinatorial point of view, arc-annotated sequences are a natural extension of simple sequences. However, using arcs for modeling non-sequential information together with restrictions on the relative positioning of

arcs allow for varying restrictions on the structure of arc-annotated sequences. Observe that a (plain) sequence without any arc can be viewed as an arc-annotated sequence with an empty arc set.

Different pattern matching and motif search problems have been considered in the context of arc-annotated sequences among which we can mention finding a *longest arc-annotated subsequence*, finding an *arc-preserving subsequence*, finding a *maximum arc-preserving common subsequence*, and computing the *edit distance for arc-annotated sequences*. Refer to [3] and [2] for overview references.

In an arc-annotated sequence (S, P) , two arcs (i_1, j_1) and (i_2, j_2) are *crossing* if $i_1 < i_2 < j_1 < j_2$ or $i_2 < i_1 < j_2 < j_1$. An arc (i_1, j_1) is *nested* into an arc (i_2, j_2) if $i_2 < i_1 < j_1 < j_2$. In her pioneering work [4], Evans has introduced a five level hierarchy¹ for arc-annotated sequences that is described as follows: UNLIMITED: no restriction at all, CROSSING: each base is incident to at most one arc, NESTED: each base is incident to at most one arc and no two arcs are crossing, STEM: each base is incident to at most one arc, and given any two arcs one is nested into the other, and PLAIN: there is no arc. This hierarchy is clearly organized according to the following chain of inclusions: PLAIN \subset STEM \subset NESTED \subset CROSSING \subset UNLIMITED.

Let (S_1, P_1) and (S_2, P_2) be two arc-annotated sequences. If $S_1[i] = S_2[j]$ for some pair of integers i and j ($1 \leq i \leq |S_1|$ and $1 \leq j \leq |S_2|$), we refer to $\langle i, j \rangle$ as a *base-match*. If $S_1[i] = S_2[j]$ and $S_1[k] = S_2[l]$ with $(i, k) \in P_1$ and $(j, l) \in P_2$, we refer to the pair $(\langle i, k \rangle, \langle j, l \rangle)$ as an *arc-match*. A common subsequence T of S_1 and S_2 can be viewed as a set of pairwise disjoint base-matches $M = \{\langle i_k, j_k \rangle : 1 \leq k \leq |T|, 1 \leq i_k \leq |S_1|, 1 \leq j_k \leq |S_2|\}$ such that $\forall 1 \leq k_1 < k_2 \leq |T|, i_{k_1} < i_{k_2}$ and $j_{k_1} < j_{k_2}$ (i.e. preserving order). The common subsequence T is said to be *arc-preserving* if the arcs induced by M are preserved, i.e., for any distinct $\langle i_{k_1}, j_{k_1} \rangle, \langle i_{k_2}, j_{k_2} \rangle \in M, (i_{k_1}, i_{k_2}) \in P_1$ if and only if $(j_{k_1}, j_{k_2}) \in P_2$. Among the many paradigms referring to arc-annotated sequences we focus here on the most natural extension of the longest common subsequence problem, the so-called LONGEST ARC-PRESERVING COMMON SUBSEQUENCE (LAPCS) problem which is defined as follows [4]: Given two arc-annotated sequences (S_1, P_1) and (S_2, P_2) , find the longest common subsequence of S_1 and S_2 that is arc-preserving. It is well-known that the LAPCS problem is **NP**-complete [4].

The LAPCS problem is traditionally parameterized by the arc-structure of the two input arc-annotated sequences. We focus on the nested case because it is widely considered the most important variant in practice [10,11,1]. We denote by LAPCS(NESTED, NESTED) (resp. LAPCS(STEM, STEM)) the LAPCS problem where both arc-annotated sequences are NESTED (resp. STEM). It has been shown in [9] that the LAPCS(NESTED, NESTED) problem is **NP**-complete, even for an unary alphabet. This result has been extended in [8] where it is shown that the LAPCS(STEM, STEM) problem is **NP**-complete. Alber et al. [1] presented two FPT algorithm for the LAPCS(NESTED,NESTED) problem. Given two arc-annotated sequences of maximum length n , their first algorithm decides in $O((3|\Sigma|)^\ell \ell n)$ time whether the two sequences have an arc-preserving common subsequence of length ℓ , and their second

¹ Our presentation actually replaces the original CHAIN level with the STEM level due to its importance for practical issues [7].

algorithm decides in $O(3.31^{k_1+k_2}n)$ time whether an arc-preserving common subsequence can be obtained by deleting k_1 letters from the first sequence and k_2 letters from the second sequence. Improving the exponential running times of the two algorithms was left as an immediate open question. Moreover, Alber et al. [1] noted that their second algorithm relies on a breadth-first search that is very space-consuming, and asked whether it can be replaced by a simple depth-first search. Our paper makes the following contributions. First, we revisit the nice **NP**-hardness proof of Lin et al. [11] for the **LAPCS(NESTED, NESTED)** problem. We point out a problem and provide a simple solution. Second, we improve the running time of the (second) FPT algorithm of Alber et al. [1] from $O(3.31^{k_1+k_2}n)$ to $O(3^{k_1+k_2}n)$. Our algorithm uses the bounded search tree technique, and can be implemented using a simple depth-first search. Third, we show that the **LAPCS(STEM, STEM)** problem is **NP**-complete for constant alphabet size. The proof is by a tricky modification of [8].

2 LAPCS(NESTED, NESTED) Is NP-complete

In this section we prove that the **LAPCS(NESTED, NESTED)** problem is **NP**-complete even if both arc-annotated sequences are unary. We actually point out a problem in a previous proof by Lin et al. [11] for the same result, and give a simple solution for the correctness of the proof. Our proof is for a large part the same as the proof of Lin et al. [11]. The only difference is that we use larger *barriers* of length $\Omega(n)$ each.

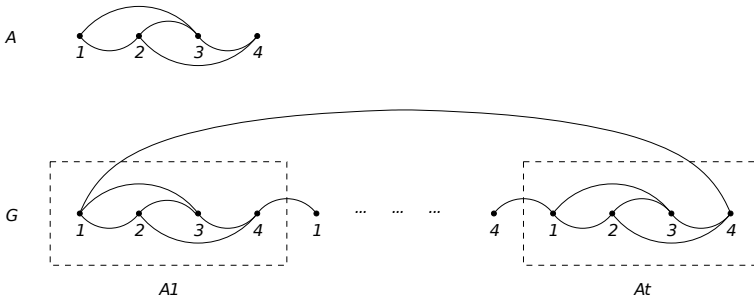


Fig. 1. The counter-example graph G .

Our counter-example graph for the proof of Lin et al. [11] is presented Figure 1. The graph A has 4 vertices v_1, v_2, v_3 and v_4 . The graph G has $n = 4t$ vertices, and consists of t copies A_1, A_2, \dots, A_t of the graph A linked into a circular “list” (for convenience let $A_0 = A_t$ and $A_{t+1} = A_1$) by one additional edge from the vertex v_1 of each A_i to the vertex v_4 of A_{i-1} . One can easily verified that G is cubic, planar, bridgeless, and connected. Moreover, G has a natural two-page book embedding such that each vertex is incident to at least 1 and at most 2 edges on each page, as illustrated in Figure 1. We have the following lemma about the graph G .

Lemma 1. *The maximum cardinality k^* of an independent set in the graph G is $\lfloor \frac{3}{8}n \rfloor$.*

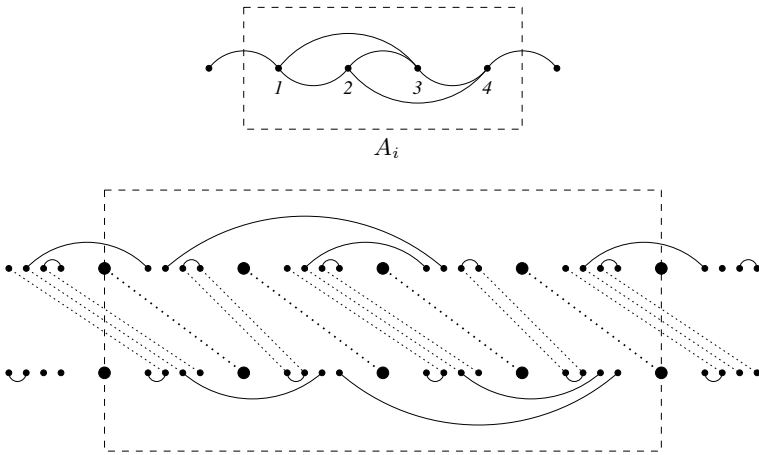


Fig. 2. The two arc-annotated sequences P_1 and P_2 for the graph G . The separating blocks, each of length 8, are illustrated by large dots.

We now turn to pointing out the problem in the proof of Lin et al. [11]. Refer to Figure 2 for the construction of the two arc-annotated sequences P_1 and P_2 based on the graph G according to the reduction of Lin et al. [11]. As illustrated by the dotted lines between the two sequences, the two arc-annotated sequences (S_1, P_1) and (S_2, P_2) has an arc-preserving common subsequence of length $\ell = 8n + \frac{3+2}{2}n - 6 = 10n + \frac{1}{2}n - 6$. Lin et al. [11] claimed that every LAPCS can be transformed into a good LAPCS (of the same length). We show that this claim is wrong. Following their proof, the graph G has an independent set of cardinality k if and only if (S_1, P_1) and (S_2, P_2) have a good LAPCS of length $8(n+1) + 2n + k = 10n + k + 8$. Then, by Lemma 1, the maximum length of a good LAPCS of (S_1, P_1) and (S_2, P_2) is at most $\ell_{\text{good}} = 10n + \lfloor \frac{3}{8}n \rfloor + 8$. Note that for $t > 28$ and correspondingly $n = 4t > 112$, we have $\ell > \ell_{\text{good}}$. This disproves their claim. For a correct proof, we increase the length of each separating block in the reduction from 8 to $s = 4n$. Then, following their proof, the length of an LAPCS is at least $s(n+1) + 2n + k^*$. If a common subsequence has a *far* match $\langle i, j \rangle$ such that $|j - i| \geq n$, then in each sequence there must be at least n unmatched bases on each side of the match. It follows that the length of the common subsequence is at most $s(n+1) + 4n - 2n$, which is less than $s(n+1) + 2n + k^*$. Therefore every match $\langle i, j \rangle$ of an LAPCS must be *near*, i.e., $|j - i| < n$. By the same argument, an LAPCS must include at least one arc from each separating block in each sequence, because otherwise a separating block with no arcs in the LAPCS would have at least $4n/2 = 2n$ unmatched bases. Since all matches must be near, any arc (i_1, i_2) in the LAPCS that comes from a separating block in P_1 must match an arc (j_1, j_2) from the corresponding separating block in P_2 such that either $i_1 \leq j_1 \leq j_2 \leq i_2$ or $j_1 \leq i_1 \leq i_2 \leq j_2$. Then a simple replacement argument shows that all separating blocks are matched completely, and consequently any LAPCS can be transformed into a good LAPCS of the same length.

3 A Faster Algorithm for the LAPCS(NESTED, NESTED) Problem

Theorem 1. *There is an $O(3^{k_1+k_2}n)$ -time algorithm for LAPCS(NESTED,NESTED) that decides whether an arc-preserving common subsequence of two arc-annotated sequences of maximum length n can be obtained by deleting k_1 letters from the first sequence and k_2 letters from the second sequence.*

We first observe that the two parameters k_1 and k_2 are not independent. Let n_1 and n_2 be the lengths of the two sequences. Then the problem admits a valid solution only if $n_1 - k_1 = n_2 - k_2$. Without loss of generality, we use a single parameter $k = k_1 + k_2$ for the total number of letters deleted from the two arc-annotated sequences. The running time of our algorithm is thus $O(3^k n)$. For an arc-annotated sequence S and an index i , define $\text{buddy}(S, i) = j$ if $S[i]$ is connected to $S[j]$ by an arc, and $\text{buddy}(S, i) = 0$ otherwise. For an arc-annotated sequence S of length n and two indices $i \leq j$, denote by $S[i, j]$ the subsequence obtained from S by deleting letters $S[1], S[2], \dots, S[i-1]$ and $S[j+1], S[j+2], \dots, S[n]$ together with the incident arcs. For an arc-annotated sequence S and three indices $i \leq j \leq k$, denote by $S[i, \bar{j}, k]$ the subsequence obtained from $S[i, k]$ by deleting $S[j]$ and its incident arc (if any).

Algorithm $\text{lapcs}(S, T, k)$

Input: Two arc-annotated sequences S and T , an integer k .

Output: returns k^* – the minimum number of letters that must be deleted from S and T to obtain an arc-preserving common subsequence – if $k^* \leq k$; ∞ otherwise.

The algorithm is recursive. For the base case, the algorithm returns 0 if $S = T$ and $k \geq 0$, and returns ∞ if $S \neq T$ and $k \leq 0$. For the inductive case, the algorithm tries all applicable following cases and returns the minimum value. Let s and t be the lengths of the two sequences S and T , respectively. Put $i = \text{buddy}(S, 1)$ and $j = \text{buddy}(T, 1)$.

Case 1. $S[1] \neq T[1]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k-1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k-1) + 1$.

Case 2.1. $S[1] = T[1]$, $i = j = 0$.

- Match $S[1] \sim T[1]$, then return $\text{lapcs}(S[2, s], T[2, t], k)$.

Case 2.2. $S[1] = T[1]$, $i > 0$ and $j = 0$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k-1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k-1) + 1$.
- Delete $S[i]$, match $S[1] \sim T[1]$, then return $\text{lapcs}(S[2, \bar{i}, s], T[2, t], k-1) + 1$.

Case 2.3. $S[1] = T[1]$, $i = 0$ and $j > 0$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k-1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k-1) + 1$.
- Delete $T[j]$, match $S[1] \sim T[1]$, then return $\text{lapcs}(S[2, s], T[2, \bar{j}, t], k-1) + 1$.

Case 2.4. $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] \neq T[j]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $S[i]$ and $T[j]$, match $S[1] \sim T[1]$, then return $\text{lapcs}(S[2, \bar{i}, s], T[2, \bar{j}, t], k - 2) + 2$.

Case 2.5.1. $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $S[2, i - 1] = T[2, j - 1]$.

- Match $S[1, i] \sim T[1, j]$, then return $\text{lapcs}(S[i + 1, s], T[j + 1, t], k)$.

Case 2.5.2. $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $S[i + 1, s] = T[j + 1, t]$.

- Match $S[1] \sim T[1]$ and $S[i, s] \sim T[j, t]$, then return $\text{lapcs}(S[2, i - 1], T[2, j - 1], k)$.

Case 2.5.3 $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $\exists a : S[2, \bar{a}, i - 1] = T[2, j - 1]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $S[a]$, match $S[1, \bar{a}, i] \sim T[1, j]$, then return $\text{lapcs}(S[i + 1, s], T[j + 1, t], k - 1) + 1$.

Case 2.5.4 $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $\exists b : S[2, i - 1] = T[2, \bar{b}, j - 1]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $T[b]$, match $S[1, i] \sim T[1, \bar{b}, j]$, then return $\text{lapcs}(S[i + 1, s], T[j + 1, t], k - 1) + 1$.

Case 2.5.5 $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $\exists a : S[i + 1, \bar{a}, s] = T[j + 1, t]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $S[a]$, match $S[1] \sim T[1]$ and $S[i, \bar{a}, s] \sim T[j, t]$, then return $\text{lapcs}(S[2, i - 1], T[2, j - 1], k - 1) + 1$.

Case 2.5.6 $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $\exists b : S[i + 1, s] = T[j + 1, \bar{b}, t]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $T[b]$, match $S[1] \sim T[1]$ and $S[i, s] \sim T[j, \bar{b}, t]$, then return $\text{lapcs}(S[2, i - 1], T[2, j - 1], k - 1) + 1$.

Case 2.5.7 $S[1] = T[1]$, $i > 0$ and $j > 0$, $S[i] = T[j]$, $S[2, i - 1] \neq T[2, j - 1]$, $S[i + 1, s] \neq T[j + 1, t]$, $\forall a : S[2, \bar{a}, i - 1] \neq T[2, j - 1]$, $\forall a : S[i + 1, \bar{a}, s] \neq T[j + 1, t]$, $\forall b : S[2, i - 1] \neq T[2, \bar{b}, j - 1]$, $\forall b : S[i + 1, s] \neq T[j + 1, \bar{b}, t]$.

- Delete $S[1]$, then return $\text{lapcs}(S[2, s], T, k - 1) + 1$.
- Delete $T[1]$, then return $\text{lapcs}(S, T[2, t], k - 1) + 1$.
- Delete $S[i]$ and $T[j]$, match $S[1] \sim T[1]$, then return $\text{lapcs}(S[2, \bar{i}, s], T[2, \bar{j}, t], k - 2) + 2$.

- Match $S[1] \sim T[1]$ and $S[i] \sim T[j]$, compute $k' = \text{lapcs}(S[2, i - 1], T[2, j - 1], k - 2) + \text{lapcs}(S[i + 1, s], T[j + 1, t], k - 2)$, then return k' if $k' \leq k$, or ∞ if $k' > k$.

The correctness of the algorithm is self-evident for the cases from 1 to 2.5.2. To justify the four cases from 2.5.3 and 2.5.6, we have the following easy lemma.

Lemma 2. *For each case from 2.5.3 to 2.5.6, if the condition of the case is met, then there is an optimal solution that corresponds to one of the three branches for that case.*

Finally, the condition for case 2.5.7 ensures that at least two deletions are necessary in each of the two subproblems for $(S[2, i - 1], T[2, j - 1])$ and $(S[i + 1, s], T[j + 1, t])$. Thus in the last branch of this case, it is sufficient to set the third parameter to $k - 2$ in the two recursions. In terms of time complexity, the seven cases 2.2, 2.3, and 2.5.3–2.5.7 are the worst cases. The six cases 2.2, 2.3, and 2.5.3–2.5.4 correspond to the characteristic polynomial equation $1 = x^{-1} + x^{-1} + x^{-1}$; the last case 2.5.7 corresponds to the characteristic polynomial equation $1 = x^{-1} + x^{-1} + x^{-2} + (x^{-2} + x^{-2})$. Both equations have a unique positive real root $x_0 = 3$.

4 LAPCS(STEM, STEM) for Constant Alphabet Size

The LAPCS(STEM, STEM) problem turns out to be of particular interest for RNA practical issues [7]. This problem has been shown to be **NP**-complete for arbitrarily large alphabets [8]. This section is devoted to investigating the LAPCS(STEM, STEM) problem for constant alphabet size. We first make the easy observation that the LAPCS(STEM, STEM) problem for an alphabet of size 1 admits a polynomial-time exact algorithm by dynamic programming. Unfortunately, this approach cannot be pushed too far. Indeed, we now show that the constant alphabet size assumption is not enough to gain tractability for the LAPCS(STEM, STEM) problem.

Theorem 2. *The LAPCS(STEM, STEM) problem is **NP**-complete for constant alphabet size.*

To prove hardness, we propose a reduction from the **NP**-complete **3-SAT** problem [6] which is defined as follows: Given a collection $C_q = \{c_1, c_2, \dots, c_q\}$ of q clauses, where each clause is the disjunction of 3 literals on a finite set of n boolean variables $V_n = \{x_1, x_2, \dots, x_n\}$, determine whether there exists a truth assignment to the variables so that each clause has at least one true literal. Let (C_q, V_n) be an arbitrary instance of the **3-SAT** problem. For convenience, let L_i^j denote the j -th literal of the i -th clause (*i.e.* c_i) of C_q . In the following, given a sequence S over an alphabet Σ , let $\text{occ}(i, c, S)$ denote the i -th occurrence of the letter c in S . We build two arc-annotated sequences (S_1, P_1) and (S_2, P_2) as follows. An illustration of a full example is given in figures 3 and 4, where $n = 4$ and $q = 3$. For readability reasons, the arc-annotated sequences resulting from the construction have been split into several parts and a schematic overview of the overall placement of each part is provided.

Let S_1 and S_2 be the two sequences defined as follows:

$$\begin{aligned} S_1 &= C_q^1 S C_{q-1}^1 \dots C_2^1 S C_1^1 S S_M^1 S P_1^1 S P_2^1 \dots P_{q-1}^1 S P_q^1 \\ S_2 &= C_q^2 S C_{q-1}^2 \dots C_2^2 S C_1^2 S S_M^2 S P_1^2 S P_2^2 \dots P_{q-1}^2 S P_q^2 \end{aligned}$$

where, for all $1 \leq i \leq q$ and $1 \leq k \leq n$,

- $S = 2^\beta$
- $C_i^1 = 9^\delta 6^\gamma 8^\delta 6^\gamma X_1^1 X_2^1 \dots X_n^1 6^\gamma 8^\delta 6^\gamma 7^\delta$ with $X_k^1 = 0$ $s_j 1 2^\alpha$ if $x_k = L_i^j$ or $\bar{x}_k = L_i^j$, with $s_1 = 3, s_2 = 4$ and $s_3 = 5$; $X_k^1 = 0 1 2^\alpha$ otherwise;
- $P_i^1 = 6^\gamma 6^\gamma 9^\delta X_n^1 \dots X_{\frac{n}{2}+1}^1 8^\delta X_{\frac{n}{2}}^1 \dots X_1^1 7^\delta 6^\gamma 6^\gamma$ s.t. $X_k^1 = 1 0 2^\alpha$;
- $C_i^2 = X_1^2 \dots X_n^2 9^\delta 6^\gamma X_1^2 \dots X_{\frac{n}{2}}^2 8^\delta X_{\frac{n}{2}+1}^2 \dots X_n^2 6^\gamma 7^\delta X_1^2 \dots X_n^2$ s.t. $\forall 1 \leq j \leq 3, \text{occ}(j, X_k^2, C_i^2) = 1 0 s_j 2^\alpha$ (resp. $s_j 1 0 2^\alpha$) if $x_k = L_i^j$ (resp. $\bar{x}_k = L_i^j$), with $s_1 = 3, s_2 = 4$ and $s_3 = 5$; $\text{occ}(j, X_k^2, C_i^2) = 1 0 2^\alpha$ otherwise;
- $P_i^2 = (0 1 2^\alpha)^n 7^\delta 6^\gamma (0 1 2^\alpha)^{\frac{n}{2}} 8^\delta (0 1 2^\alpha)^{\frac{n}{2}} 6^\gamma 9^\delta (0 1 2^\alpha)^n$.
- $S_M^1 = (0 1 2^\alpha)^n$ and $S_M^2 = (1 0 2^\alpha)^n$

Notice that, by construction, there is only one occurrence of each $\{3, 4, 5\}$ in C_i^1 and C_i^2 . Moreover, let $\alpha = 2n + 1, \beta = |S_M^1| + \sum_{1 \leq i \leq q} (|C_i^1| + |P_i^1|), \delta = \alpha(n + 1)$ and $\gamma = 5\delta + 4$. Let us now define P_1 and P_2 . Add an arc in P_1 between $\text{occ}(k, 0, S_M^1)$ (resp. $\text{occ}(k, 1, S_M^1)$) and $\text{occ}(n - k + 1, 0, P_1^1)$ (resp. $\text{occ}(n - k + 1, 1, P_1^1)$). For all $1 \leq i \leq q - 1$, (1) add an arc in P_1 between $\text{occ}(k, 0, C_i^1)$ (resp. $\text{occ}(k, 1, C_i^1)$) and $\text{occ}(n - k + 1, 0, P_{i+1}^1)$ (resp. $\text{occ}(n - k + 1, 1, P_{i+1}^1)$), $\forall 1 \leq k \leq n$ (see Fig. 3.d and 4.b); for all $1 \leq i \leq q$, (2) add an arc in P_2 between $\text{occ}(j * k, 0, C_i^2)$ (resp. $\text{occ}(j * k, 1, C_i^2)$) and $\text{occ}(3n - jk + 1, 0, P_i^2)$ (resp. $\text{occ}(3n - jk + 1, 1, P_i^2)$), $\forall 1 \leq j \leq 3, 1 \leq k \leq n$ (see Fig. 3.c, 4.a and 4.c); (3) add an arc in P_2 between $\text{occ}(k, j, C_i^2)$ and $\text{occ}(\delta - k + 1, j, P_i^2)$, $\forall j \in \{7, 8, 9\}$ and $1 \leq k \leq \delta$ (see Fig. 3.c, 4.a and 4.c). Clearly, this construction can be achieved in polynomial-time, and yields two arc-annotated sequences (S_1, P_1) and (S_2, P_2) that are both of type STEM. We now give an intuitive description of the different elements of this construction. Each clause $c_i \in C_q$ is represented by a pair (C_i^1, C_i^2) of sequences. The sequence C_i^2 is composed of three subsequences representing a selection mechanism of one of the three literals of c_i . The pair (S_M^1, S_M^2) of sequences is a control mechanism that will guarantee that a variable x_k cannot be true and false simultaneously. Finally, for each clause $c_i \in C_q$, the pair (P_i^1, P_i^2) of sequences is a propagation mechanism whose aim is to propagate the selection of the assignment (*i.e.* true or false) of any literal x_k all over C_q . Notice that all the previous intuitive notions will be detailed and clarified afterwards. In the sequel, we will refer to any such construction as a *snail-construction*. In order to complete the instance of the LAPCS(STEM, STEM) problem, we set $k' = |S_1| - \varepsilon$ with $\varepsilon = q(2(n + 2\delta + 2\gamma + 1)) + n$ where k' is the desired length of the solution. Let (S_1, P_1) and (S_2, P_2) denote the arc-annotated sequences obtained by a snail-construction. We will denote S_d the set of symbols deleted in a solution of the LAPCS problem on (S_1, P_1) and (S_2, P_2) (*i.e.* the symbols that do not belong to the common subsequence). We need some technical lemmas:

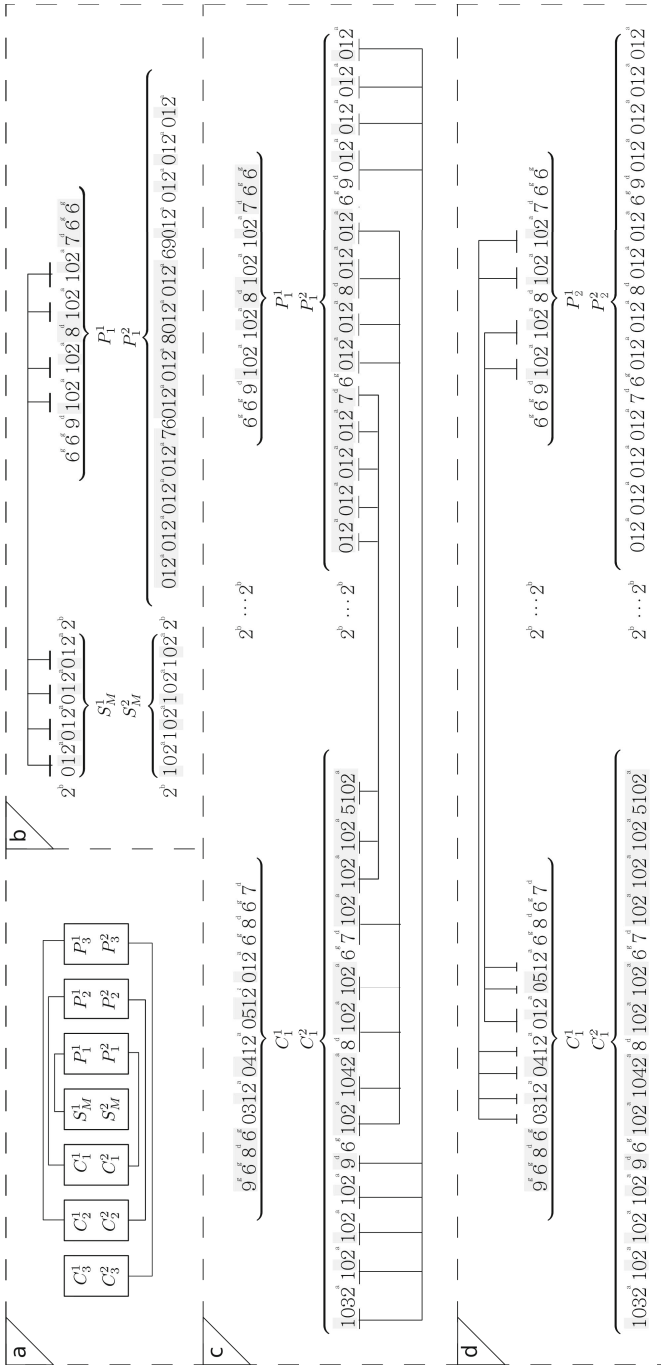


Fig. 3. Considering $C_q = (x_1 \vee x_2 \vee \overline{x_3}) \wedge (\overline{x_1} \vee \overline{x_2} \vee x_4) \wedge (x_2 \vee \overline{x_3} \vee \overline{x_4})$. For readability, all the arcs have not been drawn, consecutive arcs are representing by a unique arc with lines for endpoints. Symbols over a grey background may be deleted to obtain an optimal LAPCS. a) A schematic view of the overall arrangement of the components of the two sequences. b) Description of $S_M^1, S_M^2, P_1^1, P_1^2$ and the corresponding arcs in P_1 . c) Description of $C_1^1, C_1^2, P_1^1, P_1^2$ and the corresponding arcs in P_1 . d) Description of $C_1^1, C_1^2, P_2^1, P_2^2$ and the corresponding arcs in P_1 .

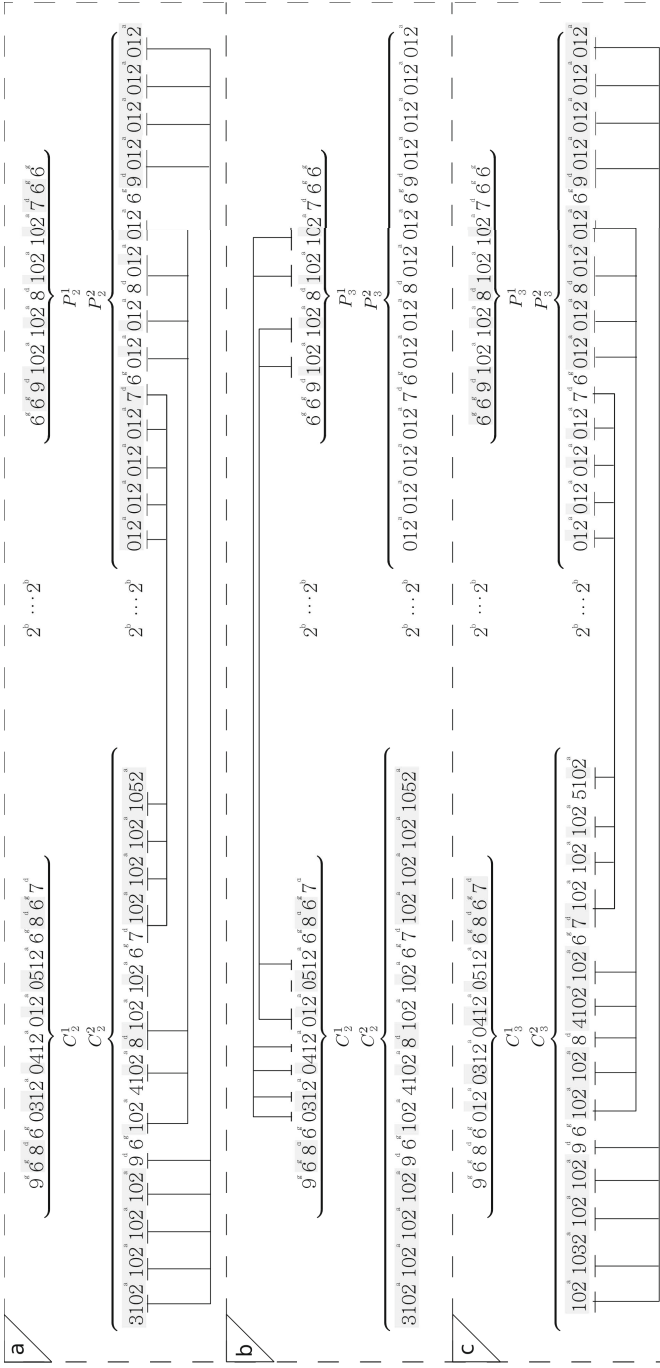


Fig. 4. Considering $C_q = (x_1 \vee x_2 \vee \overline{x_3}) \wedge (\overline{x_1} \vee \overline{x_2} \vee x_4) \wedge (x_2 \vee \overline{x_3} \vee \overline{x_4})$. For readability all the arcs have not been drawn, consecutive arcs representing by a unique arc with lines for endpoints. Symbols over a grey background may be deleted to obtain an optimal LAPCS. a) Description of $C_1^1, C_2^1, P_1^1, P_2^1$ and the corresponding arcs in P_2 . c) Description of $C_2^1, C_2^2, P_3^1, P_3^2$ and the corresponding arcs in P_1 . d) Description of $C_3^1, C_3^2, P_3^1, P_3^2$ and the corresponding arcs in P_2 .

Lemma 3. *Any optimal solution of the LAPCS(STEM, STEM) problem on (S_1, P_1) and (S_2, P_2) is of length $|S_1| - \varepsilon$.*

Lemma 4. *In any optimal solution of the LAPCS(STEM, STEM) problem on (S_1, P_1) and (S_2, P_2) , if $\text{occ}(k, 1, S_M^1)$ (resp. $\text{occ}(k, 0, S_M^1)$) for a given $1 \leq k \leq n$ is deleted then, $\forall 1 \leq j \leq q$, $\text{occ}(k, 1, C_j^1)$ (resp. $\text{occ}(k, 0, C_j^1)$) is deleted.*

The following theorem proves Theorem 2.

Theorem 3. *Given an instance of the problem 3SAT with n variables and q clauses, there exists a satisfying truth assignment if and only if the LAPCS(STEM, STEM) problem for (S_1, P_1) and (S_2, P_2) is of length $k' = |S_1| - \varepsilon$.*

Proof. (\Rightarrow) An optimal solution for $C_q = (x_1 \vee x_2 \vee \overline{x_3}) \wedge (\overline{x_1} \vee \overline{x_2} \vee x_4) \wedge (x_2 \vee \overline{x_3} \vee \overline{x_4})$ – i.e. $x_1 = x_3 = \text{true}$ and $x_2 = x_4 = \text{false}$ – is illustrated in figures 3 and 4 where any symbol over a grey background has to be deleted. Suppose we have a solution for our 3-SAT instance, that is an assignment of each variable of V_n satisfying each clause of C_q . Let us first list all the symbols to delete in S_1 . For all $1 \leq k \leq n$, if $x_k = \text{false}$ then delete, $\forall 1 \leq j \leq q$, $\{\text{occ}(k, 0, C_j^1), \text{occ}(k, 1, P_j^1)\}$ and $\text{occ}(k, 0, S_M^1)$; otherwise delete, $\forall 1 \leq j \leq q$, $\{\text{occ}(k, 1, C_j^1), \text{occ}(k, 0, P_j^1)\}$ and $\text{occ}(k, 1, S_M^1)$.

For each L_i^j satisfying c_i with the biggest index j with $1 \leq i \leq q$,

if (1) $j = 1$ then from C_i^1 , delete all the symbols 9, the two first substrings of γ symbols 6, the first substring of δ symbols 8, symbols 4 and 5. Moreover, from P_i^1 delete all the symbols 7 and 8, the two last substrings of γ symbols 6 (cf Fig. 3.c).

if (2) $j = 2$ then from C_i^1 , delete all the symbols 8, the first and the last substrings of γ symbols 6, symbols 3 and 5. Moreover, from P_i^1 delete all the symbols 7 and 9, the first and the last substrings of γ symbols 6 (cf Fig. 4.a).

if (3) $j = 3$ then from C_i^1 , delete all the symbols 7, the two last substrings of γ symbols 6, the last substring of δ symbols 8, symbols 3 and 4. Moreover, from P_i^1 delete all the symbols 8 and 9, the two first substrings of γ symbols 6.

Let us now list all the symbols in S_2 to be deleted. For all $1 \leq k \leq n$, if $x_k = \text{false}$ then delete $\text{occ}(k, 0, S_M^2)$; otherwise delete $\text{occ}(k, 1, S_M^2)$. For each L_i^j satisfying c_i with the biggest index j with $1 \leq i \leq q$,

if (1) $j = 1$ then, in C_i^2 , delete all the symbols not in $\{6, 7, 8\}$ appearing after $\text{occ}(1, 9, C_i^2)$ (included). Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(k, 0, C_i^2)$, otherwise delete $\text{occ}(k, 1, C_i^2)$ (cf Fig. 3.c). Moreover, in P_i^2 , delete all the symbols not in $\{6, 9\}$ appearing before $\text{occ}(1, 9, P_i^2)$. Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(3n - k + 1, 0, P_i^2)$, otherwise delete $\text{occ}(3n - k + 1, 1, P_i^2)$ (cf Fig. 3.c);

if (2) $j = 2$ then, in C_i^2 , delete all the symbols 8 and all the symbols appearing before $\text{occ}(1, 9, C_i^2)$ (excluded) or after $\text{occ}(\delta, 7, C_i^2)$ (excluded). Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(n + k, 0, C_i^2)$, otherwise delete $\text{occ}(n + k, 1, C_i^2)$ (cf Fig. 4.a). Moreover, in P_i^2 , delete all the symbols appearing before $\text{occ}(1, 6, P_i^2)$ (excluded) or after $\text{occ}(2\gamma, 6, P_i^2)$ (excluded). Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(2n - k + 1, 0, P_i^2)$, otherwise delete $\text{occ}(2n - k + 1, 1, P_i^2)$ (cf Fig. 4.a);

if (3) $j = 3$ then, in C_i^2 , delete all the symbols not in $\{6, 8, 9\}$ appearing before $\text{occ}(\delta, 7, C_i^2)$ (included). Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(2n + k, 0, C_i^2)$, otherwise delete $\text{occ}(2n + k, 1, C_i^2)$ (cf Fig. 4.c). Moreover, in P_i^2 , delete all the symbols not in $\{6, 7\}$ appearing after $\text{occ}(1, 7, P_i^2)$. Moreover, if $x_k = \text{false}$ with $1 \leq k \leq n$ then delete, $\text{occ}(n - k + 1, 0, P_i^2)$, otherwise delete $\text{occ}(n - k + 1, 1, P_i^2)$ (cf Fig. 4.c);

By construction, the natural order of the symbols of S_1 and S_2 allows the corresponding set of undeleted symbols to be conserved in a common arc-preserving common subsequence between (S_1, P_1) and (S_2, P_2) . Let us now prove that the length of this last is k' . One can easily check that in this solution, in S_1 , n symbols have been deleted from S_M^1 and $\forall 1 \leq i \leq q$, $2\delta + 2\gamma + n + 2$ symbols from C_i^1 and $2\delta + 2\gamma + n$ symbols from P_i^1 have been deleted. Thus, the length of the solution is $|S_1| - [q(2(n + 2\delta + 2\gamma + 1)) + n]$.

(\Leftarrow) Suppose we have an optimal solution – i.e. a set of symbols S_d to delete – for LAPCS of (S_1, P_1) and (S_2, P_2) . Let us define the truth assignment of V_n s.t., $\forall 1 \leq i \leq q$, if in C_i^1 symbol 3 is not deleted, then the first literal of clause c_i (i.e. L_i^1) is true; if in C_i^1 symbol 4 is not deleted, then the second literal of clause c_i (i.e. L_i^2) is true; if in C_i^1 symbol 5 is not deleted, then the third literal of clause c_i (i.e. L_i^3) is true. Let us prove that it is a solution for our 3-SAT instance. By construction, if $L_i^j = x_k$ (resp. $\overline{x_k}$) then in C_i^1 , symbol $2 + j$ (i.e. 3, 4 or 5) appears between $\text{occ}(k, 0, C_i^1)$ and $\text{occ}(k, 1, C_i^1)$ whereas in C_i^2 it appears after $\text{occ}(k, 1, C_i^2)$ (resp. before $\text{occ}(k, 0, C_i^2)$). Thus, if symbol $2 + j$ (i.e. 3, 4 or 5) in C_i^1 is not deleted then $\text{occ}(k, 1, C_i^1)$ (resp. $\text{occ}(k, 0, C_i^1)$) in C_i^1 is deleted if $L_i^j = x_k$ (resp. $\overline{x_k}$). Consequently, according to the proof of Lemma 4, if symbol $2 + j$ (i.e. 3, 4 or 5) in C_i^1 is not deleted then $\text{occ}(k, 1, C_{i'}^1)$ (resp. $\text{occ}(k, 0, C_{i'}^1)$) in all $C_{i'}^1$, with $1 \leq i' \leq q$ is deleted if $L_i^j = x_k$ (resp. $\overline{x_k}$). Therefore, we can ensure that one cannot obtain $\overline{L_i^j}$ and $L_{i'}^{j'}$ being true whereas $L_i^j = \overline{L_{i'}^{j'}}$ (that is a variable cannot be simultaneously true and false). By Lemma 3, we can ensure that for any $1 \leq i \leq q$ exactly one of $\{3, 4, 5\}$ is conserved in C_i^1 . \square

References

1. Alber, J., Gramm, J., Guo, J., Niedermeier, R.: Computing the similarity of two sequences with nested arc annotations. *Theoretical Computer Science* 312(2-3), 337–358 (2004)
2. Blin, G., Crochemore, M., Vialette, S.: Algorithmic Aspects of Arc-Annotated Sequences. In: *Algorithms in Computational Molecular Biology: Techniques, Approaches and Applications*. Wiley (2010) (to appear)
3. Blin, G., Denise, A., Dulucq, S., Herrbach, C., Touzet, H.: Alignment of RNA structures. *IEEE/ACM Transactions on Computational Biology and Bioinformatics* (2008) (to appear)
4. Evans, P.A.: Algorithms and Complexity for Annotated Sequences Analysis. PhD thesis, University of Victoria (1999)
5. Evans, P.A.: Finding Common Subsequences with Arcs and Pseudoknots. In: Crochemore, M., Paterson, M. (eds.) *CPM 1999*. LNCS, vol. 1645, pp. 270–280. Springer, Heidelberg (1999)
6. Garey, M.R., Johnson, D.S.: *Computers and Intractability: A guide to the theory of NP-completeness*. W.H. Freeman, San Francisco (1979)

7. Guignon, V., Chauve, C., Hamel, S.: An Edit Distance Between RNA Stem-Loops. In: Consens, M.P., Navarro, G. (eds.) SPIRE 2005. LNCS, vol. 3772, pp. 335–347. Springer, Heidelberg (2005)
8. Blin, G., Hamel, S., Vialette, S.: Comparing RNA Structures with Biologically Relevant Operations Cannot Be Done without Strong Combinatorial Restrictions. In: Rahman, M. S., Fujita, S. (eds.) WALCOM 2010. LNCS, vol. 5942, pp. 149–160. Springer, Heidelberg (2010)
9. Jiang, T., Lin, G., Ma, B., Zhang, K.: A general edit distance between RNA structures. *Journal of Computational Biology* 9(2), 371–388 (2002)
10. Jiang, T., Lin, G., Ma, B., Zhang, K.: The Longest Common Subsequence Problem for Arc-Annotated Sequences. In: Giancarlo, R., Sankoff, D. (eds.) CPM 2000. LNCS, vol. 1848, pp. 154–165. Springer, Heidelberg (2000)
11. Lin, G., Chen, Z.-Z., Jiang, T., Wen, J.: The longest common subsequence problem for sequences with nested arc annotations. *J. of Computer and System Sc.* 65, 465–480 (2002)
12. Shasha, D., Zhang, K.: Simple fast algorithms for the editing distance between trees and related problems. *SIAM Journal on Computing* 18(6), 1245–1262 (1989)