



An Efficient Algorithm for Enumerating Longest Common Increasing Subsequences

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Abstract. The longest common increasing subsequence (LCIS) problem is the combination of two classic problems in algorithms: the longest increasing subsequence (LIS) problem and the longest common subsequence (LCS) problem. In this paper, we propose an algorithm that finds every LCIS of two sequences a, b of length n in $O(n + \sigma + I_a)$ time and space, where σ denotes the size of the alphabet set and I_a the total number of increasing subsequences contained in a (thus, the running time is output-sensitive). Our algorithm employs the trie and some simple data structures, and thus is implementation-wise simple. In addition, it can be proved that our algorithm is optimal in time complexity when $\sigma \leq \log_2 n$.

Keywords: LCIS · Trie · Data structure

1 Introduction

The longest common increasing subsequence (LCIS) problem can be formulated as follows: Given a sequence $a = a_1, a_2, \dots, a_n$, a sequence $a_{i_1}, a_{i_2}, \dots, a_{i_k}$ is a subsequence of a if $1 \leq i_j < i_{j+1} \leq n$ for all $1 \leq j < k$. And, given two sequences a, b of length n , the LCIS problem asks for a longest common subsequence of a, b that is strictly increasing.

This problem can be seen as a combination of the longest increasing subsequence (LIS) problem and the longest common subsequence (LCS) problem, and was first introduced by Yang et al. [6] and then applied to the whole genome alignment by Chan et al. [1] in 2005. Yang et al. and Chan et al. proposed algorithms of $O(n^2)$ and $O(\min(r \log \sigma, n\sigma + r) \log \log n + \text{Sort}_n)$ time, respectively, where Sort_n denotes the time required to sort input sequences a, b , and r the number of ordered pairs (i, j) such that $a_i = b_j$. In 2006, Sakai presented a linear-space and $O(n^2)$ -time algorithm using a divide-and-conquer approach [5]. In 2011, Kutz et al. designed an algorithm of $O(n)$ space and $O(nl \log \log \sigma + \text{Sort}_n)$ time [3], where l denotes the length of the LCIS of a, b . And, for small alphabet set, algorithms of $O(n)$ and $O(n \log \log n)$ time were proposed for $\sigma = 2$ and $\sigma = 3$, respectively. In 2016, Zhu et al. proposed an

$O(n^2)$ -time and linear-space algorithm [7]. Recently, in 2020, Lo et al. proposed an algorithm of $O(n + l(n - l) \log \log \sigma)$ time and $O(n)$ space [4], and Duraj presented the first algorithm of subquadratic time [2].

The rest of this paper is organized as follows: In Sect. 2, the proposed algorithm is presented. In Sect. 3, the correctness and complexity are analyzed. Finally, we conclude this paper in Sect. 4.

2 The Proposed Algorithm

In this section, three assumptions are first introduced. Subsequently, we outline the proposed algorithm, followed by a step-by-step explanation along with the pseudocode. Finally, an example is given.

2.1 Assumptions

Input Format. Given the size of the alphabet set σ , we assume the alphabet set consists of integers $0, 1, \dots, \sigma - 1$, i.e., each integer in the input sequences a, b is in $\{0, 1, \dots, \sigma - 1\}$.

Fast Computation. We assume that the bitwise shift (or bitwise OR) on one (or two) binary encoded data of no more than σ bits can be done in $O(1)$ time.

Constant Space. We assume that a bitstring of length up to σ takes $O(1)$ space.

Remark that when the desired input format is not satisfied, one can map integers in a, b to the integers in $[0, \sigma - 1]$ without changing the order of integers in $O(n \log \sigma)$ time using a balanced binary search tree.

2.2 Algorithm Overview

The main procedure of the proposed algorithm (Algorithm 1) involves building a trie T containing the information of every increasing subsequence in a . Let IS_u denote the *increasing subsequence* with *binary encoding* u , i.e., the i -th bit in u is 1 if and only if i is contained in IS_u . For example, $IS_{10100100}$ denotes the increasing subsequence $[2, 5, 7]$ as $\sigma = 8$. Then, T has the following properties:

1. A node T_u associated with the length l_u of IS_u exists in T to denote an *increasing subsequence* IS_u if and only if IS_u is found in a . Also, the binary encoding u of IS_u is stored in T_u to help retrieval of sequence information.
2. A directed edge associated with $x \in \{0, 1, \dots, \sigma - 1\}$ from T_u to T_v , denoted by the tuple (T_u, T_v, x) , exists in T if and only if IS_v is the concatenation of IS_u and x .

After building T , a similar trie-building process is run for sequence b ; but instead of building a new trie for b , we walk along the nodes of T that denote the common increasing subsequences of a, b , and meanwhile record all the found longest common increasing subsequences of a, b .

For the complexity of Algorithm 1, the initialization step takes $O(\sigma+n)$ time. Building T takes $O(n+I_a)$ time by using additional data structures that take $O(\sigma+I_a)$ space. And, walking on T takes $O(n+I_a)$ time. To sum up, Algorithm 1 has space and time complexity of $O(n+\sigma+I_a)$.

2.3 Detailed Description

See Algorithm 1 for the pseudocode. Algorithm 1 consists of 5 parts as follows.

Input/Output. Algorithm 1 takes two sequences a, b , the length n of a, b , and the size of the alphabet set σ as the inputs, and outputs a list L containing the binary encoding of every LCIS of a, b .

Initialization for First Loop (Lines 2–19). Firstly, build an array Cnt , where $Cnt[i]$ is the frequency of i in a for all $i \in \{0, 1, \dots, \sigma - 1\}$. This can be done in $O(n)$ time by simply scanning a once. Secondly, build a doubly linked list K of nodes to store every integer i with $Cnt[i] > 0$ in an increasing order (from $i = 0$ to $i = \sigma - 1$). Then, a pointer array M of size σ is created. And, for each integer i stored in K , the pointer to the node containing i in K is stored in $M[i]$ so that the node can be removed from K , if necessary, in $O(1)$ time. Thirdly, build the root node T_0 of T , which denotes an empty sequence, and set l_0 to 0. Then, for every i with $Cnt[i] > 0$, create a trie node T_{2^i} containing the binary encoding of the sequence $[i]$, set $l_{2^i} = 1$, and add an edge (T_0, T_{2^i}, i) from T_0 to T_{2^i} . Finally, build σ queues $Next_0, \dots, Next_{\sigma-1}$, where each queue supports $O(1)$ push and pop (for our purpose, one can also use different data structures such as stacks or dynamic arrays, as long as they support push and pop in $O(1)$ time). Let A_u be the address of T_u . Then, for all i , queue $Next_i$ initially contains A_{2^i} if $Cnt[i] > 0$, and is left empty otherwise.

First Loop (Lines 20–31). Algorithm 1 iterates the following two steps when sequence a is scanned one by one from left to right. Firstly, for the i -th integer a_i in a , we decrease $Cnt[a_i]$ by 1. Secondly, in the inner loop, Algorithm 1 iterates the following two substeps until queue $Next_{a_i}$ is empty. First pop A_u from queue $Next_{a_i}$ and get u from T_u . Then, in the (yet deeper) inner loop, for each integer x with $a_i < x < \sigma$ and $Cnt[x] > 0$ (every such x can be found efficiently using K), first create a new node T_v of T , set l_v to $l_u + 1$, add an edge (T_u, T_v, x) from T_u to T_v , where $v = u + 2^x$, and then push A_v into queue $Next_x$. At last, at the end of the i -th iteration, remove the node containing a_i from K if $Cnt[a_i]$ has become 0.

Algorithm 1: LCIS

Input: (n, σ, a, b) : the length of each sequence, the size of the alphabet set, the two sequences

Output: L : a list containing the binary code of every LCIS of (a, b)

```

1 begin
2    $Cnt \leftarrow$  new 1D integer array of size  $\sigma$ ;
3    $M \leftarrow$  new 1D pointer array of size  $\sigma$ ;
4   for  $i \leftarrow 0$  to  $\sigma - 1$  do
5      $Cnt[i] \leftarrow 0$ ;
6   for  $i \leftarrow 1$  to  $n$  do
7      $Cnt[a_i] \leftarrow Cnt[a_i] + 1$ ;
8    $K \leftarrow$  new doubly linked list;
9   create trie node  $T_0$ ;
10   $l_0 \leftarrow 0$ ;
11  for  $i \leftarrow 0$  to  $\sigma - 1$  do
12     $Next_i \leftarrow$  new queue;
13    if  $Cnt[i] > 0$  then
14      add the node  $K_i$  containing  $i$  to  $K$ ;
15       $M[i] \leftarrow$  address of  $K_i$ ;
16      create trie node  $T_{2^i}$ ;
17       $l_{2^i} \leftarrow 1$ ;
18      add edge  $(T_0, T_{2^i}, i)$ ;
19      push  $A_{2^i}$  into  $Next_i$ ;
20  for  $i \leftarrow 1$  to  $n$  do
21     $Cnt[a_i] \leftarrow Cnt[a_i] - 1$ ;
22    for  $A_u \in Next_{a_i}$  do
23      pop  $A_u$  from  $Next_{a_i}$  and get  $u$  from  $T_u$ ;
24      for  $x \leftarrow a_i + 1$  to  $\sigma - 1$  in  $K$  do
25         $v \leftarrow u + 2^x$ ;
26        create trie node  $T_v$ ;
27         $l_v \leftarrow l_u + 1$ ;
28        add edge  $(T_u, T_v, x)$ ;
29        push  $A_v$  into  $Next_x$ ;
30    if  $Cnt[a_i] = 0$  then
31      remove the node containing  $a_i$  from  $K$ ;
32   $len \leftarrow 0$ ;
33   $L \leftarrow$  new list;
34  insert 0 into  $L$ ;
35  for  $i \leftarrow 0$  to  $\sigma - 1$  do
36    if trie node  $T_{2^i}$  exists then
37      push  $A_{2^i}$  into  $Next_i$ ;
38  ... (continued in next page)

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37
38   for  $i \leftarrow 1$  to  $n$  do
39     for  $A_u$  in  $Next_{b_i}$  do
40       pop  $A_u$  from  $Next_{b_i}$  and get  $u$  and  $l_u$  from  $T_u$ ;
41       if  $l_u > len$  then
42          $len \leftarrow l_u$ ;
43         empty  $L$ ;
44       if  $len = l_u$  then
45         insert  $u$  into  $L$ ;
46       for every edge  $(T_u, T_v, x)$  from  $T_u$  do
47         push  $A_v$  into  $Next_x$ ;
48   return the list  $L$ ;

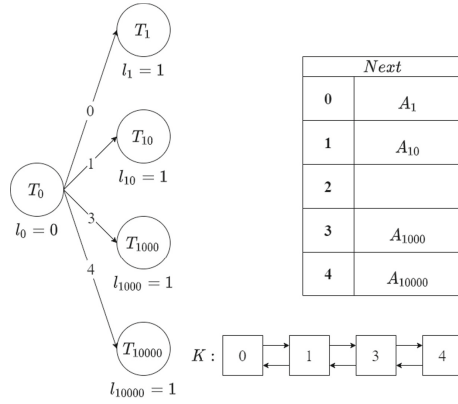
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Initialization for Second Loop (Lines 32–37). Firstly, set len , denoting the length of LCIS of a, b currently found, to 0. Secondly, build a list L to store the binary encoding of every common increasing subsequence (CIS) of length len of a, b , where L contains only 0 (the binary encoding of the empty sequence) initially. Thirdly, reuse $Next$ queues and for each queue $Next_i$, push A_{2^i} into queue $Next_i$ if T_{2^i} exists in T .

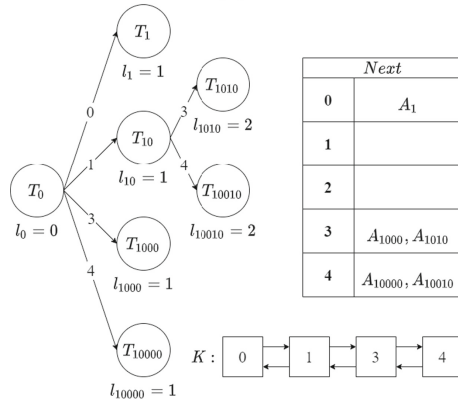
Second Loop (Lines 38–47). Algorithm 1 iterates the following step when sequence b is scanned one by one from left to right. For the i -th integer b_i in b , Algorithm 1 iterates the following substeps until queue $Next_{b_i}$ is empty in the inner loop. First pop one A_u from queue $Next_{b_i}$. Then, since IS_u is a newly found CIS of a, b , we may need to update len and L accordingly: 1) if $l_u > len$ (i.e., the length of IS_u is greater than that of any CIS of a, b currently found), empty L and update len to l_u , and 2) if $l_u = len$, add u into L . Finally, in the (yet deeper) inner loop, push A_v into queue $Next_x$ for each edge (T_u, T_v, x) from T_u .

2.4 Example

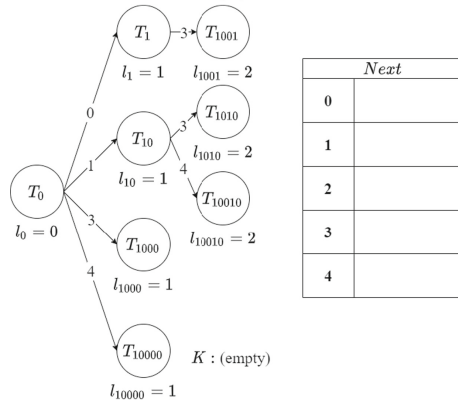
Figures 1a and 1b show the statuses of K , $Next$, and T on the termination of the initialization and iteration 1, respectively, of the first loop of Algorithm 1 for $a = [1, 4, 1, 0, 3]$ and $\sigma = 5$. During the execution of the initialization, $Cnt[0] = Cnt[3] = Cnt[4] = 1$, $Cnt[1] = 2$, and $Cnt[2] = 0$ since the frequencies of integers 0, 1, 2, 3, 4 are 1, 2, 0, 1, 1, respectively. And, since $Cnt[i] > 0$ for $i = 0, 1, 3, 4$, the nodes storing integers 0, 1, 3, 4 are doubly linked in sequence in K , the nodes T_1 , T_{10} , T_{1000} , and T_{10000} (which contains the binary encodings of integers 0, 1, 3, 4, respectively) are created in T , and A_1 , A_{10} , A_{1000} , and A_{10000} (which are the addresses of T_1 , T_{10} , T_{1000} , and T_{10000} , respectively) are contained in $Next_0$, $Next_1$, $Next_3$, and $Next_4$, respectively. In iteration 1, $a_1 = 1$. Thus, $Cnt[1]$ is



(a)

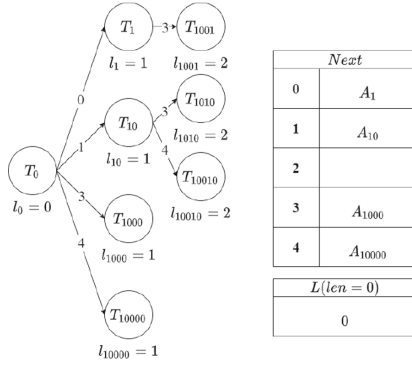


(b)



(c)

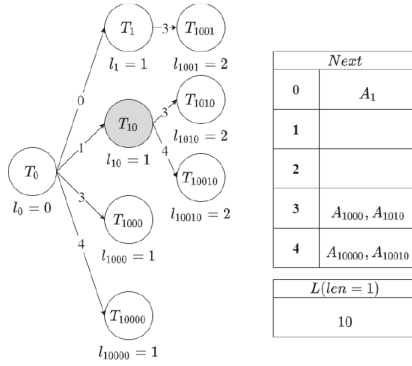
Fig. 1. The statuses of *K*, *Next*, and *T* on the termination of (a) the initialization, (b) iteration 1, (c) the last iteration of the first loop of Algorithm 1 as the input sequence a is $[1, 4, 1, 0, 3]$.



<i>Next</i>	
0	A_1
1	A_{10}
2	
3	A_{1000}
4	A_{10000}

$L(icn = 0)$	
0	

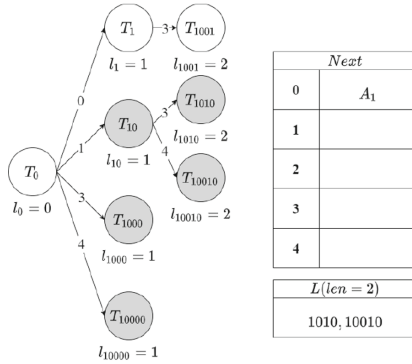
(a)



<i>Next</i>	
0	A_1
1	
2	
3	A_{1000}, A_{1010}
4	A_{10000}, A_{10010}

$L(icn = 1)$	
10	

(b)



<i>Next</i>	
0	A_1
1	
2	
3	
4	

$L(icn = 2)$	
1010, 10010	

(c)

Fig. 2. The statuses of L , $Next$, and the encountered trie nodes in T on the termination (a) the initialization, (b) iteration 1, (c) the last iteration of the second loop of Algorithm 1 as the input sequence b is $[1, 4, 3, 1, 3]$, where the encountered trie nodes in T are shown in grey.

decreased to 1 and A_{10} is popped from $Next_1$. Due to that $a_1 = 1 < x < 5 = \sigma$ and $Cnt[x] > 0$ for $x = 3, 4$, the nodes T_{1010} and T_{10010} (which contains the binary encodings of increasing sequences $[1, 3], [1, 4]$, respectively) are added to T , and A_{1010} and A_{10010} are pushed into $Next_3$ and $Next_4$, respectively. In iteration 2, the node containing integer 4 is removed from K since $a_2 = 4$ and $Cnt[4]$ becomes 0. Similarly, the nodes containing integers 1, 0, and 3 are removed from K in iterations 3, 4, and 5, respectively. In addition, A_{10000} and A_{10010} are popped from $Next_4$ in iteration 2, A_1 is popped from $Next_0$, T_{1001} is added to T , and A_{1001} is pushed into $Next_3$ in iteration 4, and A_{1001} is popped from $Next_3$ in iteration 5. The statuses of K , $Next$, and T on the termination of the first loop is shown in Fig. 1c.

Figures 2a and 2b show the statuses of L , $Next$ and the encountered trie nodes in T on the termination of the initialization and iteration 1, respectively, of the second loop of Algorithm 1 for $b = [1, 4, 3, 1, 3]$. For the second loop, initially, A_1 , A_{10} , A_{1000} , and A_{10000} are pushed into $Next_0$, $Next_1$, $Next_3$, and $Next_4$, respectively, since trie nodes T_1 , T_{10} , T_{1000} , and T_{10000} exist in T ; also, L contains a single element 0, and len is set to 0. In iteration 1, since $b_1 = 1$, A_{10} is popped from $Next_1$, T_{10} is encountered, and L is updated to contain 10 only. Meanwhile, since edge $(T_{10}, T_{1010}, 3)$ exists in T , A_{1010} is pushed into $Next_3$. Similarly, A_{10010} is pushed into $Next_4$. In iteration 2, A_{10000} and A_{10010} is popped from $Next_4$, T_{10000} and T_{10010} are encountered, and L is updated to contain 10010. In iteration 3, A_{1000} and A_{1010} are popped from $Next_3$, T_{1000} and T_{1010} are encountered, and 1010 is inserted to L . In iteration 4 (or 5), the statuses of L and $Next$ remains unchanged since $Next_1$ ($Next_3$) is empty. Figure 2c shows the statuses of L , $Next$ and the encountered trie nodes in T on the termination of the second loop.

3 The Analysis

In this section, we first show the correctness of the proposed algorithm. Subsequently, the time and space complexity of the proposed algorithm is studied.

3.1 Correctness

Lemma 1. *In the first loop, a non-empty increasing subsequence IS_u of a exists if and only if A_u has been popped from some $Next$ queue.*

Proof. It suffices to show for each i ($1 \leq i \leq n$), on the termination of iteration i of the first loop, a non-empty increasing subsequence IS_u exists in a_1, a_2, \dots, a_i (a prefix of a) if and only if A_u has been popped from some $Next$ queue. We show it by induction on the number of iterations executed.

Clearly, on the termination of iteration 1, $[a_1]$ is the only one non-empty increasing subsequence of a . Besides, in the initialization of the first loop, Algorithm 1 pushes A_{2^u} into queue $Next_u$ once for each integer u that exists in a . Since Algorithm 1 pops all items in queue $Next_{a_1}$ in iteration 1, only $A_{2^{a_1}}$ has

been popped on the termination of iteration 1. Thus, we have a basis. We then assume the induction hypothesis: on the termination of iteration k , a non-empty increasing subsequence IS_u exists in a_1, a_2, \dots, a_k if and only if A_u has been popped from some *Next* queue. To complete the proof, we only need to show the induction step: on the termination of iteration $k + 1$, a non-empty increasing subsequence IS_u exists in a_1, a_2, \dots, a_{k+1} if and only if A_u has been popped from some *Next* queue.

For the *if* part, if A_u is popped from some *Next* queue before iteration $k + 1$, IS_u exists in a_1, a_2, \dots, a_k by induction hypothesis, and thus IS_u exists in a_1, a_2, \dots, a_{k+1} . So, we only need to consider the case where A_u is popped from some *Next* queue in iteration $k + 1$. Note that Algorithm 1 pops all items in queue $Next_{a_{k+1}}$ in iteration $k + 1$. Also note that Algorithm 1 only pushes A_u into queue $Next_{a_{k+1}}$ when IS_u ends with a_{k+1} . Let IS_u be the concatenation of IS_v and a_{k+1} . Clearly, if IS_v is an empty sequence, $IS_u = a_{k+1}$ is an increasing subsequence of a . Otherwise, let IS_v end with a_j ; then, A_v has been popped from some *Next* queue on the termination of iteration k and $a_j < a_{k+1}$ because otherwise, A_u is not in *Next* queues in iteration $k + 1$ by Algorithm 1. By induction hypothesis, IS_v is an increasing subsequence in a_1, a_2, \dots, a_k . This implies IS_u is an increasing subsequence in a_1, a_2, \dots, a_{k+1} , completing the proof of the *if* part.

For the *only if* part, if IS_u does not end with a_{k+1} , IS_u exists in a_1, a_2, \dots, a_k , and thus A_u has been popped from some *Next* queue on the termination of iteration k by the induction hypothesis. So, we only need to consider the case where IS_u ends with a_{k+1} . Since Algorithm 1 pops all items in queue $Next_{a_{k+1}}$ in iteration $k + 1$, we only need to show A_u is in queue $Next_{a_{k+1}}$ on the termination of iteration k . Let IS_u be the concatenation of IS_v and a_{k+1} . Then, if IS_v is an empty sequence, A_u is pushed into queue $Next_{a_{k+1}}$ in the initialization of the first loop. Otherwise, IS_v is a non-empty increasing subsequence in a_1, a_2, \dots, a_k . Let IS_v end with a_j . Since IS_u is an increasing subsequence, we have $a_j < a_{k+1}$. Then, on the termination of iteration k , A_v has been popped from some *Next* queue by induction hypothesis, and then A_u has been pushed into queue $Next_{a_{k+1}}$ due to $a_j < a_{k+1}$ and $Cnt[a_{k+1}] > 0$, completing the *only if* part.

Theorem 1. *In the first loop, IS_u is a non-empty increasing subsequence of a if and only if a trie node T_u is created in T .*

Proof. Note that Algorithm 1 creates a trie node T_u right before A_u is pushed into some *Next* queue in the first loop. Thus, a trie node T_u is created in T if and only if A_u has been pushed into some *Next* queue. Besides, IS_u is a non-empty increasing subsequence of a if and only if A_u has been popped from some *Next* queue by Lemma 1. Thus, to complete the proof, we only need to show A_u has been pushed into some *Next* queue if and only if A_u has been popped from some *Next* queue. Clearly, A_u has been pushed into some *Next* queue if A_u has been popped from some *Next* queue. On the other hand, suppose A_u is pushed into some *Next* queue, say $Next_x$, in iteration j . Then, $Cnt[x] > 0$ in iteration j . This implies x exists in $a_{j+1}, a_{j+2}, \dots, a_n$. Let a_k be the first integer

in $a_{j+1}, a_{j+2}, \dots, a_n$ such that $x = a_k$. Then, A_u is popped from queue $Next_x$ in iteration k . This completes the proof.

Lemma 2. *In the second loop, a non-empty common increasing subsequence IS_u of a, b exists if and only if A_u has been popped from some $Next$ queue.*

Proof. It suffices to show for each i ($1 \leq i \leq n$), on the termination of iteration i of the second loop, a non-empty common increasing subsequence IS_u of a, b exists in b_1, b_2, \dots, b_i (a prefix of b) if and only if A_u has been popped from some $Next$ queue. We show it by induction on the number of iterations executed. In iteration 1, Algorithm 1 pops all items from queue $Next_{b_1}$. Let u be the binary encoding of b_1 . Then, b_1 exists in a_1, a_2, \dots, a_n if and only if trie node T_u exists in T by Theorem 1, and thus b_1 exists in a_1, a_2, \dots, a_n if and only if A_u is pushed into queue $Next_{b_1}$ in the initialization of the second loop. This implies that a non-empty common increasing subsequence IS_u of a, b exists in b_1 if and only if A_u has been popped from queue $Next_{b_1}$ in iteration 1. Thus, we have a basis. We omit the induction hypothesis and the proof of the induction step due to their similarities to that of Lemma 1.

Theorem 2. *By Algorithm 1, the list L contains exactly the binary encoding of every LCIS of a, b .*

Proof. By Lemma 2, for every non-empty CIS IS_u of a, b , A_u has been popped from some $Next$ queue in the second loop. In Algorithm 1, when A_u is popped from some $Next$ queue, the binary encoding of IS_u is added to L if the length of IS_u is equal to that of the CIS of a, b in L , and L is updated to contain only the binary encoding of IS_u if the length of IS_u is greater than that of the CIS of a, b in L . This ensures the binary encoding of every LCIS of a, b is contained in L .

3.2 Complexity

Lemma 3. *Every A_u for a non-empty increasing subsequence IS_u of a is pushed into $Next$ queues at most once in (a) the first loop and (b) the second loop.*

Proof. The proof of (b) is omitted due to its similarity to that of (a). We show (a) by contradiction. Suppose that A_u is the first one to be pushed into $Next$ queues more than once. Then, $|IS_u|$ must be greater than 1; otherwise, A_u is pushed into $Next$ queues only once in the initialization step.

Let IS_u be the concatenation of IS_v and x (i.e., IS_u ends with x). Note that A_u is pushed into queue $Next_x$ right after A_v is popped from some queue. Thus, A_u is pushed into $Next$ queues the second time right after A_v is popped from some queue the second time. This implies A_v is pushed into some queue twice before A_u is pushed into some queue twice. This constitutes a contradiction.

Theorem 3. *The time and space complexity of Algorithm 1 is $O(n + \sigma + I_a)$.*

Proof. Apart from the input sequences a, b themselves, K and Cnt require $O(\sigma)$ space, and T and $Next$ queues require $O(I_a)$ space by Lemma 3, so the space complexity is $O(n + \sigma + I_a)$. Note that the space complexity can be reduced to $O(n + I_a)$ through removing integers that do not exist in both of a and b from the alphabet set by additional preprocessing before Algorithm 1.

For time complexity, the initialization steps of the first and second loops require $O(n + \sigma)$ time. In the first and second loops, there are $O(I_a)$ queue and trie node operations by Lemma 3. Each queue operation requires $O(1)$ time. And, since the address of T_u is pushed into $Next$ queues, each trie node operation can be achieved in $O(1)$ time. Since Algorithm 1 uses Cnt and K to avoid iterations without queue or trie node operation, the time complexity is $O(n + \sigma + I_a)$.

Remark that due to that $I_a \leq 2^\sigma$, the time complexity of Algorithm 1 is $O(n)$, which is optimal for the LCIS problem, as $\sigma \leq \log_2 n$.

4 Conclusion and Discussion

In this paper, we present an algorithm of $O(n + \sigma + I_a)$ time and space complexity to find every LCIS of sequences a, b of length n . If the proposed algorithm is run on two computers in parallel, the time complexity can be improved to $O(n + \sigma + \min(I_a, I_b))$. When the alphabet set is small, an algorithm of $O(n \log \log n)$ time complexity was proposed in the literature for $\sigma = 3$ [3]. By contrast, the proposed algorithm has $O(n)$ time and space complexity as $\sigma \leq \log_2 n$. For the LCIS problem of k ($k > 2$) sequences, an algorithm of $O(n + \sigma + kI_a)$ time complexity can be obtained through slight modification of the proposed algorithm by just running the second loop for each sequence other than a and keeping track of how many times each node in T is encountered. Whether the proposed algorithm can be modified to better adapt to the cases of more than 2 sequences may be worthy of discussion.

References

1. Chan, W.T., Zhang, Y., Fung, S.P.Y., Ye, D., Zhu, H.: Efficient algorithms for finding a longest common increasing subsequence. *J. Comb. Optim.* **13**(3), 277–288 (2006). <https://doi.org/10.1007/s10878-006-9031-7>
2. Duraj, L.: A sub-quadratic algorithm for the longest common increasing subsequence problem. [arXiv:1902.06864](https://arxiv.org/abs/1902.06864) [cs], January 2020
3. Kutz, M., Brodal, G.S., Kaligosi, K., Katriel, I.: Faster algorithms for computing longest common increasing subsequences. *J. Discret. Algorithms* **9**(4), 314–325 (2011). <https://doi.org/10.1016/j.jda.2011.03.013>
4. Lo, S.F., Tseng, K.T., Yang, C.B., Huang, K.S.: A diagonal-based algorithm for the longest common increasing subsequence problem. *Theor. Comput. Sci.* **815**, 69–78 (2020). <https://doi.org/10.1016/j.tcs.2020.02.024>. <https://www.sciencedirect.com/science/article/pii/S0304397520301158>
5. Sakai, Y.: A linear space algorithm for computing a longest common increasing subsequence. *Inf. Process. Lett.* **99**(5), 203–207 (2006). <https://doi.org/10.1016/j.ipl.2006.05.005>

6. Yang, I.H., Huang, C.P., Chao, K.M.: A fast algorithm for computing a longest common increasing subsequence. *Inf. Process. Lett.* **93**(5), 249–253 (2005). <https://doi.org/10.1016/j.ipl.2004.10.014>
7. Zhu, D., Wang, L., Wang, T., Wang, X.: A simple linear space algorithm for computing a longest common increasing subsequence. [arXiv:1608.07002](https://arxiv.org/abs/1608.07002) [cs], August 2016