# Faster Algorithms for Computing Longest Common Increasing Subsequences

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Note: letters may occur repeatedly in the subsequence

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*Important:* here, letters may not occur repeatedly (strictly increasing subsequence)



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- LIS in O(n log n) time [Fredman, 1975]
  (also as corollary of O(r log n)-time algorithm above)



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Quite recently introduced by Yang, Huang, and Chao (IPL, 2005): They compute LCIS in  $\Theta(mn)$  time and space.



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#### **New Result:**

An LCIS for a length-m and a length-n sequence can be computed in  $O\left((m+n\ell)\log\log|\Sigma|+Sort_{\Sigma}(m)\right) \text{ time, where } \ell = \text{length of LCIS.}$ 



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Even O(m) space possible using randomized data structures; then it's *expected* running time.

(uses Willard's y-fast tries)



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4-letter LCWIS remains open


# Applications



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 $L_1[1] = 8$ 

**Theorem.** An LCIS for a length-m seq. A and a length-n seq. B can be computed in  $O((m + n\ell) \log \log |\Sigma| + Sort_{\Sigma}(m))$  time.

A dynamic-programming approach, but not over the  $A \times B$  table.

1
 2
 3
 4
 5
 6
 7
 8
 9

 A:
 
$$\gamma$$
 $\alpha$ 
 $\beta$ 
 $\delta$ 
 $\alpha$ 
 $\beta$ 
 $\epsilon$ 
 $\gamma$ 

 B:
  $\beta$ 
 $\delta$ 
 $\beta$ 
 $\alpha$ 
 $\epsilon$ 
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A dynamic-programming approach, but not over the  $A \times B$  table.

Instead, evaluate arrays  $L_i[j]$ : minimal index  $\kappa$  in B such that there exists lenght-i CIS on A[1..i] and  $B[1..\kappa]$  ending on  $a_i$ .  $L_1[1] = 8$  $L_1[9] = 8$ 1 2 3 4 5 6 7 8 9



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$L_1[9] = 8$		1	2	3	4	5	6	7	8	9
$L_2[4] = 9$	A :	γ	α	α	β	δ	α	β	e	γ
	B:	β	δ	β	α	α	e	δ	γ	β



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 $L_2[5] = 2$ 

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$L_2[5] = 2$	B :	β	δ	β	α	α	e	δ	γ	β
$L_3[8] = 6$										





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combine McCreight's priority search tree with van Emde Boas trees



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**priority**: the index k (in B)

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 $\begin{array}{ll} \text{items:} & \text{length-}(i-1) \text{ CIS ending on } a_h = b_k & \text{(in A resp. B)} \\ \text{key:} & \text{the letter } a_h = b_k \\ \text{priority:} & \text{the index } k \text{ (in B)} \end{array}$ 

Each operation in  $O(\log \log |\Sigma|)$  time

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Example: want to compute  $L_3[8]$ 

• query ( $\epsilon$ ) "where does longest length-2 sequence with last letter <  $\epsilon$  end in B?"



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- query ( $\epsilon$ ) "where does longest length-2 sequence with last letter <  $\epsilon$  end in B?"
- answer: at position 2
- find next occurrence of  $\epsilon$  after position 2 in B: 6
- set new value L<sub>3</sub>[8] := 6



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- take the best result over all r

 $\longrightarrow O(m)$  time



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A hierarchical distribution of information reduces all information storage to  $O(m + n \log n)$  time.

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An LCIS or LCWIS of k length-n sequences can be computed in  $O(r \log^{k-1} \log \log r)$  time, where r = # of match vectors.



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#### 4 Multiple Sequences

In this section we consider the problem of finding an LCIS of k length-n sequences, for  $k \geq 3$ . We will denote the sequences by  $A^1 = (a_1^1, \ldots, a_n^1)$ ,  $A^2 = (a_1^2, \ldots, a_n^2), \ldots, A^k = (a_1^k, \ldots, a_n^k)$ . A match is a vector  $(i_1, i_2, \ldots, i_k)$  of indices such that  $a_{i_1}^1 = a_{i_2}^2 = \cdots = a_{i_k}^k$ . Let r be the number of matches. Chan et al. [4] showed that an LCIS can be found in  $O(\min(kr^2, kr \log \sigma \log^{k-1} r) + kSort_{\Sigma}(n))$  time (they present two algorithms, each corresponding to one of the terms in the min). We present a simpler solution which replaces the second term by  $O(r \log^{k-1} r \log \log r)$ .

We denote the *i*th coordinate of a vector v by v[i], and the alphabet symbol corresponding to the match described by a vector v will be denoted s(v). A vector v dominates a vector v' if v[i] > v'[i] for all  $1 \le i \le k$ , and we write v' < v. Clearly, an LCIS corresponds to a sequence  $v_1, \ldots, v_\ell$  of matches such that  $v_1 < v_2 < \cdots < v_\ell$  and  $s(v_1) < s(v_2) < \cdots < s(v_\ell)$ .

To find an LCIS, we use a data structure by Gabow et al. [6, Theorem 3.3], which stores a fixed set of n vectors from  $\{1, \ldots, n\}^k$ . Initially all vectors are *inactive*. The data structure supports the following two operations:

- 1. Activate a vector with an associated priority.
- 2. A query of the form "what is the maximum priority of an active vector that is dominated by a vector p?"

A query takes  $O(\log^{k-1} n \log \log n)$  time and the total time for at most n activations is  $O(n \log^{k-1} n \log \log n)$ . The data structure requires  $O(n \log^{k-1} n)$  preprocessing time and space.

Each of the r matches  $v = (v_1, \ldots, v_k)$  corresponds to a vector. The priority of v will be the length of the longest LCIS that ends at the match v. We will consider the matches by non-decreasing order of their symbols. For each symbol s of the alphabet, we first compute the priority of every match v with s(v) = s. This is equal to 1 plus the maximum priority of a vector dominated by v. Then, we activate these vectors in the data structure with the priorities we have computed; they should be there when we compute the priorities for matches v with s(v) > s.

The algorithm applies to the case of a common weakly-increasing subsequence by the following modification: The matches will be considered by non-decreasing order of s(v) as before, but within each symbol also in non-decreasing lexicographic order of v. For each match, we compute its priority and immediately activate it in the data structure (so that it is active when considering other matches with the same symbol). The lexicographic order ensures that if v > v'then v' is in the data structure when v is considered.

**Theorem 4.** An LCIS or LCWIS of k length-n sequences can be computed in  $O(r \log^{k-1} r \log \log r)$  time, where r counts the number of match vectors.

#### 5 Outlook

The central question about the LCS problems is, whether it can be solved in  $O(n^{2-\epsilon})$  time in general. It seems that with LCIS we face the same frontier. Our

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- With LCS, is the 2-letter case as hard as the general problem?