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An Algorithm for the Constrained Longest Common Subsequence and Substring Problem

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Subsequences and Substrings

-Let \sum be an alphabet and **S** a string over \sum . A subsequence of a string **S** is obtained by deleting zero or more letters from **S**.

If S ="ACGTU", then "ATU" is a subsequence of S.

-A substring of a string S is a subsequence of S consists of consecutive letters in S.

If S ="ACGTU", then "CGT" is a substring of S, "ATU" is not a substring of S,

-Every substring of **S** is also a subsequence of **S**.

-The empty string is a subsequence and a substring of any string.

The Longest Common Subsequence Problem for Two Strings

-The longest common subsequence problem for two strings X and Y is to

find a longest string, denoted LCSSeq(X, Y), which is a subsequence

of both X and Y.

-Obviously, the set of LCSSeq(X, Y) and the set LCSSeq(Y, X) are the same.

|LCSSeq(X, Y)| = |LCSSeq(Y, X)|.

The Longest Common Substring Problem for Two Strings

-The longest common substring problem for two strings X and Y is to

find a longest string, denoted LCSStr(X, Y), which is a substring

of both X and Y.

-Obviously, the set of LCSStr(X, Y) and the set LCSStr(Y, X) are the same.

|LCSStr(X, Y)| = |LCSStr(Y, X)|.

-In [1], Li, Deka, and Deka introduced the longest common subsequence and

substring problem for two strings X and Y which is to find a longest string,

denoted LCSSeqStr(X, Y), that is a subsequence of X and a substring Y.

-[1] R. Li, J. Deka, and K. Deka, An algorithm for the longest common

subsequence and substring problem, Journal of Math and Informatics

25 (2023) 77-81.

-In general, the set of LCSSeqStr(X, Y) and the set LCSSeqStr(Y, X)

are not the same. $|LCSSeqStr(X, Y)| \neq |LCSSeqStr(Y, X)|$.

-In [1], Li, Deka, and Deka designed an algorithm for LCSSeqStr(X, Y). The time

and space complexities of the algorithm are O(|X| |Y|), where |X| and |Y| are the

lengths of strings X and Y, respectively.

Examples

- Suppose $\mathbf{X} =$ "GAAAAACCCT" and $\mathbf{Y} =$ "GACACACT".
- -"AC" is a longest common substring for X (resp. Y) and Y (resp. X). Thus |LCSStr(X, Y)| = |LCSStr(Y, X)| = 2.
- -"ACT" is a longest common subsequence and substring for **X** and **Y**. Thus $|LCSSeqStr(\mathbf{X}, \mathbf{Y})| = 3.$

-"ACCCT" is a longest common subsequence and substring for **Y** and **X**. Thus $|LCSSeqStr(\mathbf{Y}, \mathbf{X})| = 5$.

-"GAAACT" is a longest common subsequence for **X** (resp. **Y**) and **Y** (resp. **X**). Thus LCSSeq(**X**, **Y**)| = |LCSSeq(**Y**, **X**)| = 6.

The Three Problems

 $-|LCSStr(X, Y)| = |LCSStr(Y, X)| \le |LCSSeqStr(X, Y)| \le$

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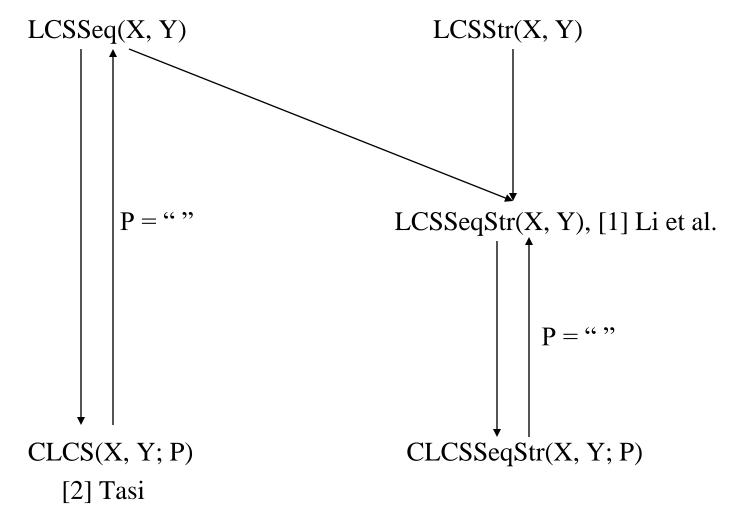
|LCSSeq(X, Y)| = |LCSSeq(Y, X)|.

 $-|LCSStr(X, Y)| = |LCSStr(Y, X)| \le \min\{|LCSSeqStr(X, Y)|, |LCSSeqStr(Y, X)|\}$

 $\leq \max\{|LCSSeqStr(X, Y)|, |LCSSeqStr(Y, X)|\}$

 $\leq |LCSSeq(X, Y)| = |LCSSeq(Y, X)|.$

The Problems



-Tsai [2] extended the longest common subsequence problem for two strings to the constrained longest common subsequence problem for two strings and a constrained string P.

-[2] Y. T. Tsai, The constrained longest common subsequence problem,

Information Processing Letters 88 (2003) 173-176.

-For two strings X, Y, and a constrained string P, the constrained longest common subsequence problem for two strings X and Y with respect to P is to find a longest string Z: = CLCSSeq(X, Y; P) such that Z is a subsequence of both X and Y and P is a subsequence of Z.

-Clearly, if P is an empty string, then CLCSSeq(X, Y; P) = LCSSeq(X, Y).

-"Such a problem could arise in computing the homology of two biological sequences which have a specific or putative structure in common" quoted from [2].

-Tsai [2] designed an $O(|X^2|Y|^2|P|)$ time algorithm for CLCSSeq(X, Y; P).

-Motivated by Tasi's work, we introduced the constrained longest common

subsequence and substring problem for two strings and a constrained string.

-For two strings X, Y, and a constrained string P, the constrained longest common

subsequence and substring problem for two strings X and Y with respect to P,

is to find a longest string Z: = CLCSSeqStr(X, Y; P) such that Z is a

subsequence of X, a substring of Y, and P is a subsequence of Z.

-Clearly, if P is an empty string, then CLCSSeqStr(X, Y; P) = LCSSeqStr(X, Y) in [1].

-Suppose $\mathbf{X} =$ "GAAAAACCCT", $\mathbf{Y} =$ "GACACACT", $\mathbf{P} =$ "AC".

-"ACT" is a constrained longest common subsequence and substring for X and Y.

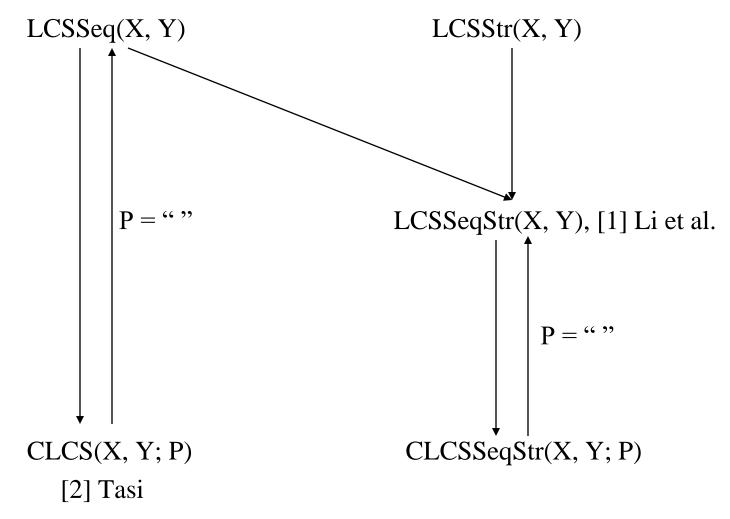
Thus $|CLCSSeqStr(\mathbf{X}, \mathbf{Y}; \mathbf{P})| = 3.$

-"GAAACT" is a constrained longest common subsequence for Y (resp. X) and X

(resp. Y). Thus |CLCSSeq(X, Y; P)| = |LCSSeq(Y, X; P)| = 6.

-In general, $|CLCSSeqStr(\mathbf{X}, \mathbf{Y}; \mathbf{P})| \leq |CLCSSeq(\mathbf{X}, \mathbf{Y}; \mathbf{P})|$.

The Problems



-Using dynamic programing (DP), we designed an algorithm for finding CLCSSeqStr(X, Y; P). Both time complexity and space complexity of our algorithm are O(|X| |Y| |P|).

-Let $S = s_1 s_2 \dots s_r$ be a string over an alphabet Σ , The r prefixes of

S are defined as $S_1 = s_1$, $S_2 = s_1s_2$, $S_3 = s_1s_2s_3$, ..., and $S_r = s_1s_2...s_r$.

 S_0 is defined as an empty string.

-The r suffixes of S are defined as $T_1 = s_1 s_2 \dots s_r$, $T_2 = s_2 s_3 \dots s_r$, \dots , $T_{r-1} = s_{r-1} s_r$,

and $T_r = s_r$,

-Let $X = x_1 x_2 \dots x_m$, $Y = y_1 y_2 \dots y_n$, and $P = p_1 p_2 \dots p_r$. Define Z[i, j, k] as a string

satisfying the following conditions.

(1) it is a subsequence of $X_i = x_1 x_2 \dots x_{i}$,

(2) it is a suffix of $Y_j = y_1 y_2 \dots y_{j}$

(3) it has $P_k = p_1 p_2 \dots p_k$ as a subsequence,

(4) under (1), (2) and (3), its length is as large as possible,

where $1 \le i \le m$, $1 \le j \le n$, and $1 \le k \le r$.

-We will use a 3-dimensional array M[m + 1][n + 1][r + 1] to store |Z[i, j, k]|.

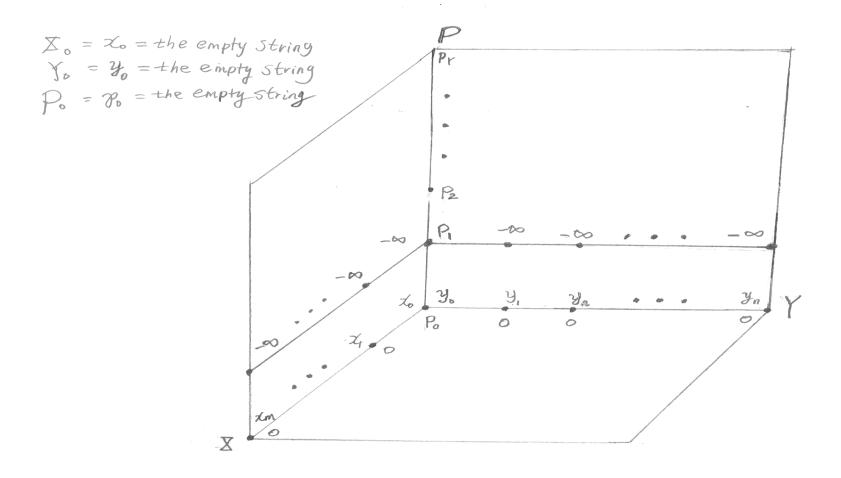
Namely, M[i][j][k] = |Z[i, j, k]|, where $0 \le i \le m$, $0 \le j \le n$, $0 \le k \le r$.

-We will recursively fill in the cells in M.

-Firstly, we fill in the boundary cells in array M.

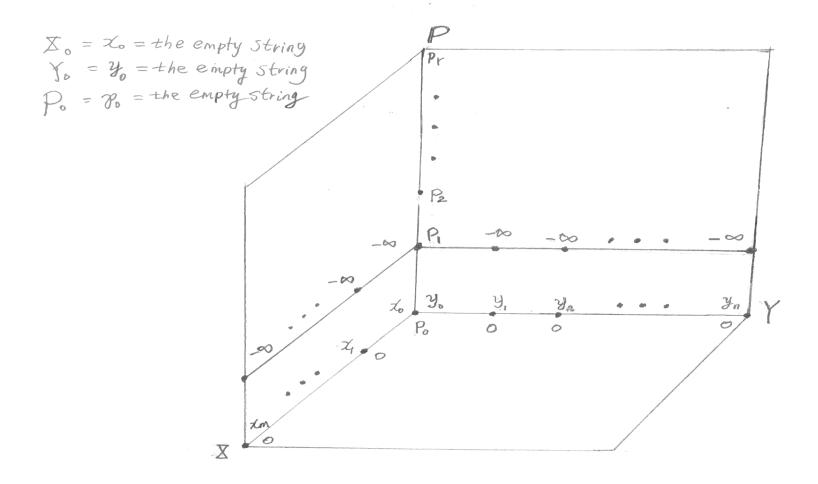
-Filling in the boundary cells.

[1] If i = 0 and k = 0 the length of a string which is a subsequence of X_i , a suffix of Y_j , and has P_k as a subsequence is zero. Thus M[0][j][0] = 0, where $0 \le j \le n$.



-Filling in the boundary cells.

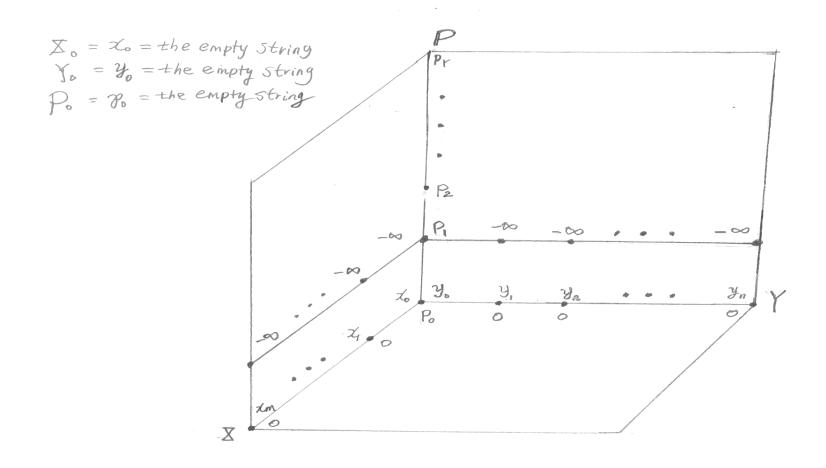
[2] If j = 0 and k = 0, the length of a string which is a subsequence of X_i , a suffix of Y_j , and has P_k as a subsequence is zero. Thus M[i][0][0] = 0, where $0 \le i \le m$.



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-Filling in the boundary cells.

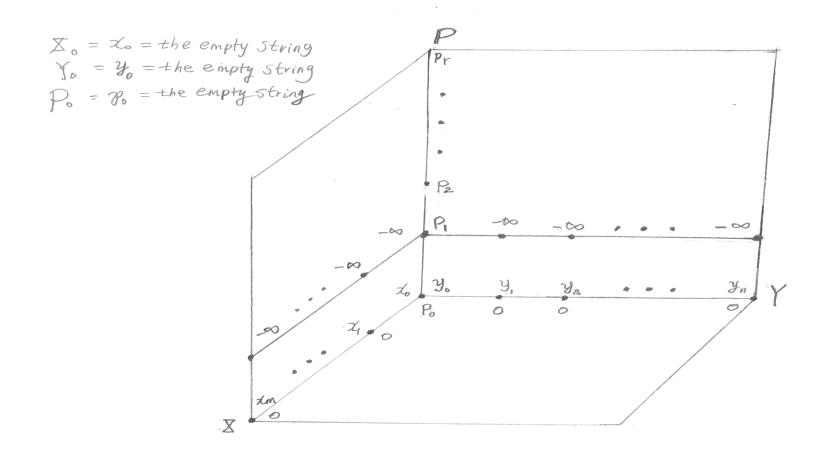
[3] If i = 0 and k \ge 1, there is not a string which is a subsequence of X_i, a suffix of Y_i, and has P_k as a subsequence. Thus M[0][j][k] = - ∞ , where $0 \le j \le n$ and $1 \le k \le r$.



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-Filling in the boundary cells.

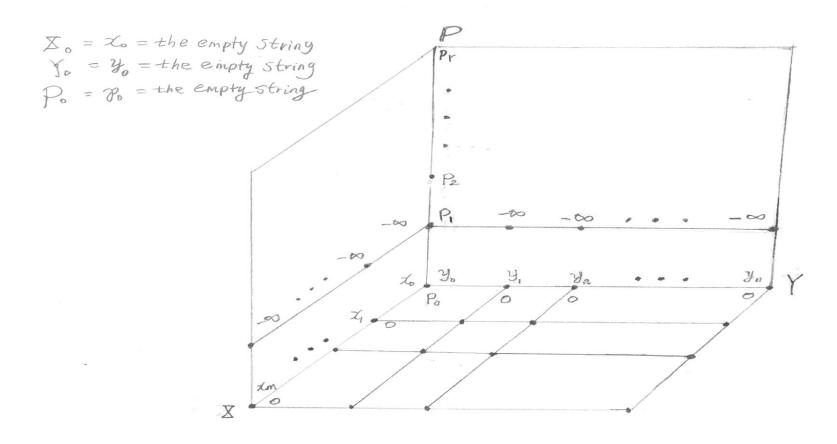
[4] If j = 0 and $k \ge 1$, there is not a string which is a subsequence of X_i , a suffix of Y_j , and has P_k as a subsequence. Thus $M[i][0][k] = -\infty$, where $0 \le i \le m$ and $1 \le k \le r$.



-Filling in the boundary cells.

[5] If k = 0 or P_k is an empty string. Then CLCSSeqStr(X, Y; P_k) = LCSSeqStr(X, Y) in [1]. The cells of M[i][j][0], where $1 \le i \le m$ and $1 \le j \le n$, can be filled in by the following rules.

If $x_i = y_j$, then M[i][j][0] = M[i - 1][j - 1][0] + 1. If $x_i \neq y_j$, then M[i][j][0] = M[i - 1][j][0].



-Filling in other cells.

-<u>Claim 1.</u> Suppose that $X_i = x_1 x_2 \dots x_i$, $Y_j = y_1 y_2 \dots y_j$, and $P_k = p_1 p_2 \dots p_k$, where $1 \le i \le m$, $1 \le j \le n$, $1 \le k \le r$. If $Z[i, j, k] = z_1 z_2 \dots z_a$ is a string satisfying conditions

(1) it is a subsequence of $X_i = x_1 x_2 \dots x_{i,j}$

(2) it is a suffix of $Y_j = y_1 y_2 \dots y_{j_j}$

(3) it has $P_k = p_1 p_2 \dots p_k$ as a subsequence,

(4) under (1), (2) and (3), its length is as large as possible,

where $1 \le i \le m$, $1 \le j \le n$, and $1 \le k \le r$.

Then we have only the following possible cases and the statement in each case is true.

-Filling in other cells.

-<u>Claim 1.</u>

Case 1. $x_i = y_j = p_k$. We have |Z[i, j, k]| = |Z[i - 1, j - 1, k - 1]| + 1 in this case.

Case 2. $x_i = y_j \neq p_k$. We have |Z[i, j, k]| = |Z[i - 1, j - 1, k]| + 1 in this case.

Case 3. $x_i \neq y_j$, $x_i \neq p_k$, and $y_j = p_k$. We have |Z[i, j, k]| = |Z[i - 1, j, k]| in this case.

Case 4. $x_i \neq y_j$, $x_i \neq p_k$, and $y_j \neq p_k$. We have |Z[i, j, k]| = |Z[i - 1, j, k]| in this case.

Case 5. $x_i \neq y_j$, $x_i = p_k$, and $y_j \neq p_k$. This case cannot happen.

-Filling in other cells.

-<u>Claim 2.</u> Suppose there is not a string which is a subsequence of $X_i = x_1 x_2 \dots x_i$,

a suffice of $Y_i = y_1 y_2 \dots y_k$ and has $P_k = p_1 p_2 \dots p_k$, as a subsequence, where

 $1 \le i \le m, 1 \le j \le n, 1 \le k \le r$. Namely, Z[i, j, k] doesn't exist. Then

[1]. If $x_i = y_j = p_k$, then there is not a string which is a subsequence of X_{i-1}

 $= x_1 x_2 \dots x_{i-1}$, a suffix of $Y_{j-1} = y_1 y_2 \dots y_{j-1}$, and has $P_{k-1} = p_1 p_2 \dots p_{k-1}$

as a subsequence. Namely, if Z[i, j, k] does not exist, then Z[i - 1, j - 1, k - 1]

does not exist either.

-Filling in other cells.

-<u>Claim 2.</u>

[2]. If $x_i = y_j \neq p_k$, then there is not a string which is a subsequence of

$$X_{i-1} = x_1 x_2 \dots x_{i-1}$$
, a suffix of $Y_{j-1} = y_1 y_2 \dots y_{j-1}$, and has $P_k = p_1 p_2 \dots p_k$

as a subsequence. Namely, if Z[i, j, k] does not exist, then Z[i - 1, j - 1, k]

does not exist either.

-Filling in other cells.

-<u>Claim 2.</u>

[3]. If $x_i \neq y_j$, $x_i \neq p_k$, and $y_j = p_k$, then there is not a string which is a

subsequence of $X_{i-1} = x_1 x_2 \dots x_{i-1}$, a suffix of $Y_j = y_1 y_2 \dots y_j$, and has

 $P_k = p_1 p_2 \dots p_k$ as a subsequence. Namely, if Z[i, j, k] does not exist, then

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-Filling in other cells.

-<u>Claim 2.</u>

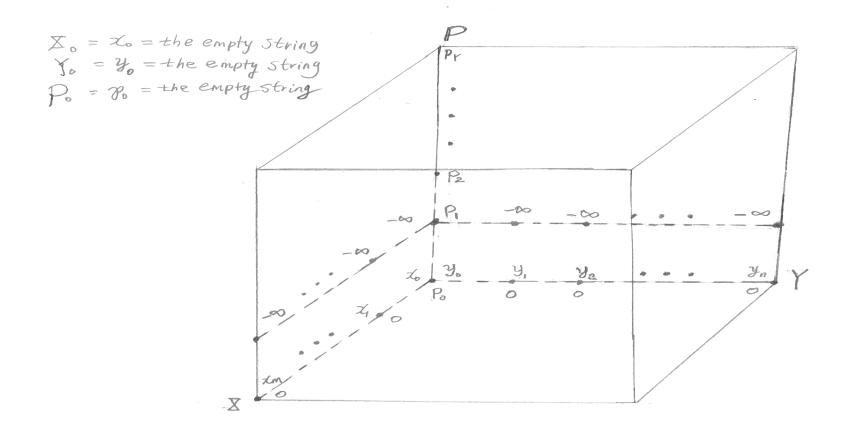
[4]. If $x_i \neq y_j$, $x_i \neq p_k$, and $y_j \neq p_k$, then there is not a string which is a

subsequence for $X_{i-1} = x_1 x_2 \dots x_{i-1}$, a suffix of $Y_j = y_1 y_2 \dots y_j$, and has

 $P_k = p_1 p_2 \dots p_k$ as a subsequence. Namely, if Z[i, j, k] does not exist,

then Z[i - 1, j, k] does not exist either.

-<u>Claim 3.</u> Let U_k be a longest string which is a subsequence of X, a substring of Y, and has P_k as a subsequence. Then $|U_k| = \max\{|Z[i, j, k]| : 1 \le i \le m, 1 \le j \le n, 1 \le k \le r\}$. Thus $|U_r| = \max\{|Z[i, j, r]| : 1 \le i \le m, 1 \le j \le n\} = |CLCSSubStr(X, Y; P)|$.



 $-|CLCSSeqStr(X, Y; P)| = \max\{M[i][j][r]: 0 \le i \le m, 0 \le j \le n\}.$

-We can also find the CLCSSeqStr(X, Y; P) when we write a program.

-The time complexity of our algorithm is

 $O((|X| + 1)(|Y| + 1)(|P| + 1)) \sim O(|X| |Y| |P|).$

-The space complexity of our algorithm also is

 $O((|X| + 1)(|Y| + 1)(|P| + 1)) \sim O(|X| |Y| |P|).$

Thanks