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# **Department of Computer Engineering**

Batch:-B-2

**Roll No:-**16010122151

**Experiment No:-10** 

Grade: AA / AB / BB / BC / CC / CD /DD

Signature of the Staff In-charge with date

Title: Implementation of Longest Common Subsequence String Matching Algorithm

**Objective:** To compute longest common subsequence for the given two strings.

#### CO to be achieved:

CO 2	Analyze and solve problems for divide and conquer strategy, greedy method, dynamic programming approach and backtracking and branch & bound policies.
CO 3	Analyze and solve problems for different string matching algorithms.

#### **Books/ Journals/ Websites referred:**

- 1. Ellis horowitz, Sarataj Sahni, S.Rajsekaran," Fundamentals of computer algorithm", University Press
- 2. T.H.Cormen ,C.E.Leiserson,R.L.Rivest and C.Stein," Introduction to algorithms",2nd Edition ,MIT press/McGraw Hill,2001
- 3. http://www.math.utah.edu/~alfeld/queens/queens.

# **Pre Lab/ Prior Concepts:**

Data structures, Concepts of algorithm analysis

#### **Historical Profile:**

Given 2 sequences, X = x1, ..., xm and Y = y1, ..., yn, find a subsequence common to both whose length is longest. A subsequence doesn't have to be consecutive, but it has to be in order.

### **New Concepts to be learned:**

String matching algorithm, Dynamic programming approach for LCS, Applications of LCS.

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#### **Recursive Formulation:**

Define c[i, j] = length of LCS of Xi and Yj. Final answer will be computed with c[m, n].

$$c[i, j] = 0$$
  
if  $i=0$  or  $j=0$ .  
 $c[i, j] = c[i - 1, j - 1] + 1$   
if  $i,j>0$  and  $xi=yj$   
 $c[i, j] = max(c[i - 1, j], c[i, j - 1])$   
if  $i, j > 0$  and  $x_i <> y_j$ 

#### **Algorithm: Longest Common Subsequence**

# Compute length of optimal solution-

```
LCS-LENGTH (X, Y, m, n)

for i \in 1 to m

do c[i, 0] \in 0

for j \in 0 to n

do c[0, j] \in 0

for i \in 1 to m

do if xi = yj

then c[i, j] \in c[i - 1, j - 1] + 1

b[i, j] \in "\approx"

else if c[i - 1, j] \ge c[i, j - 1]

then c[i, j] \in c[i - 1, j]

b[i, j] \in "\uparrow"

else c[i, j] \in c[i, j - 1]

b[i, j] \in "\uparrow"
```

**return** c and b

```
Print the solution-

PRINT-LCS(b, X, i, j)

if i = 0 or j = 0

then return

if b[i, j] = "\approx"

then PRINT-LCS(b, X, i = 1, j = 1)

print xi

elseif b[i, j] = "\uparrow"

then PRINT-LCS(b, X, i = 1, j)

else PRINT-LCS(b, X, i, j = 1)
```

Initial call is PRINT-LCS(b, X, m, n). b[i, j] points to table entry whose subproblem we used in solving LCS of Xi and Yj.



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When  $b[i, j] = \infty$ , we have extended LCS by one character. So longest common subsequence = entries with  $\approx$  in them.

# **Analysis of LCS computation:-**

```
#include <iostream>
#include <vector>
#include <algorithm>
std::string lcs(const std::string& X, const std::string& Y) {
    int m = X.size();
    int n = Y.size();
    // Create a table to store lengths of LCS of substrings
    std::vector<std::vector<int>> dp(m + 1, std::vector<int>(n + 1, 0));
    for (int i = 1; i <= m; ++i) {
        for (int j = 1; j <= n; ++j) {</pre>
            if (X[i - 1] == Y[j - 1]) {
                dp[i][j] = dp[i - 1][j - 1] + 1;
                dp[i][j] = std::max(dp[i - 1][j], dp[i][j - 1]);
        }
    }
    std::string lcs_string;
    int i = m, j = n;
    while (i > 0 \&\& j > 0) {
        if (X[i-1] == Y[j-1]) {
            lcs_string = X[i - 1] + lcs_string;
            --i:
        } else if (dp[i - 1][j] > dp[i][j - 1]) {
            --i:
        } else {
            --j;
        }
    return lcs_string;
```



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```
int main() {
    std::string X, Y;
    std::cout << "Enter the first string: ";
    std::cin >> X;
    std::cout << "Enter the second string: ";
    std::cin >> Y;
    std::string result = lcs(X, Y);
    std::cout << "Longest Common Subsequence: " << result << std::endl;
    return 0;
}</pre>
```

### **Output:-**

```
Enter the first string: iambtechstudent
Enter the second string: btech
Longest Common Subsequence: btech

=== Code Execution Successful ===
```

### Algorithm:-



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### **Conclusion:-**

The LCS algorithm efficiently determines the longest common subsequence between two sequences, aiding in tasks like string comparison, plagiarism detection, and bioinformatics analysis. By employing dynamic programming techniques, LCS achieves optimal substructure and overlapping subproblems resolution, resulting in a time complexity of O(mn) for sequences of lengths m and n. Its versatility and effectiveness make it a fundamental tool in various fields requiring sequence comparison and pattern recognition.