DAA Practical 5

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TASK 1:

Aim: Implement a dynamic algorithm for Longest Common Subsequence (LCS) to find the length and LCS for DNA sequences.

Problem Statement:

(i) DNA sequences can be viewed as strings of A, C, G, and T characters, which represent nucleotides. Finding the similarities between two DNA sequences are an important computation performed in bioinformatics.

[Note that a subsequence might not include consecutive elements of the original sequence.]

TASK 1: Find the similarity between the given X and Y sequence.

X=AGCCCTAAGGGCTACCTAGCTT

Y= GACAGCCTACAAGCGTTAGCTTG

Output: Cost matrix with all costs and direction, final cost of LCS and the LCS.

Length of LCS=16

CODE:

#include <stdio.h>

#include <string.h>

```
void LCS(char X[], char Y[]) {
  int m = strlen(X);
  int n = strlen(Y);
  int C[MAX][MAX];
  char LCS[MAX];
  int i, j;
  for (i = 0; i <= m; i++)
     C[i][0] = 0;
  for (j = 0; j \le n; j++)
     C[0][j] = 0;
  for (i = 1; i \le m; i++) {
     for (j = 1; j \le n; j++) {
       if (X[i-1] == Y[j-1])
          C[i][j] = C[i-1][j-1] + 1;
       else if (C[i - 1][j] >= C[i][j - 1])
          C[i][j] = C[i - 1][j];
       else
          C[i][j] = C[i][j - 1];
     }
```

```
}
                   int index = C[m][n];
                   LCS[index] = '\0';
                  i = m;
                  j = n;
                  while (i > 0 \&\& j > 0) {
                                       if (X[i-1] == Y[j-1]) {
                                                           LCS[index - 1] = X[i - 1];
                                                          i--;
                                                         j--;
                                                           index--;
                                      ext{ } ext{ | } ext{ 
                                                           i--;
                                       else
                                                         j--;
                  }
                   printf("Length of LCS = %d\n", C[m][n]);
                   printf("LCS = %s\n", LCS);
int main() {
```

}

```
char X[] = "AGCCCTAAGGGCTACCTAGCTT";
char Y[] = "GACAGCCTACAAGCGTTAGCTTG";

LCS(X, Y);
return 0;
}
```

Output:



TASK-2: Find the longest repeating subsequence (LRS). Consider it as a variation of the longest common subsequence (LCS) problem.

Let the given string be S. You need to find the LRS within S. To use the LCS framework, you effectively compare S with itself. So, consider string1 = S and string2 = S.

Example:

AABCBDC

LRS= ABC or ABD

CODE:

#include <stdio.h>

#include <string.h>

```
int max(int a, int b) { return a > b ? a : b; }
void LRS(char *str) {
  int n = strlen(str);
  int dp[n+1][n+1];
  for (int i = 0; i \le n; i++)
     for (int j = 0; j <= n; j++)
       if (i == 0 | | i == 0)
          dp[i][j] = 0;
       else if (str[i-1] == str[j-1] && i != j)
          dp[i][j] = dp[i-1][j-1] + 1;
       else
          dp[i][j] = max(dp[i-1][j], dp[i][j-1]);
  int i = n, j = n, k = dp[n][n];
  char lrs[k+1];
  Irs[k] = '\0';
  while (i > 0 \&\& j > 0) {
     if (str[i-1] == str[j-1] \&\& i != j) {
       Irs[--k] = str[i-1];
       i--; j--;
     } else if (dp[i-1][j] > dp[i][j-1])
       i--;
     else
```

```
j--;
}

printf("LRS Length: %d\n", dp[n][n]);
printf("LRS: %s\n", Irs);
}

int main() {
    char S[] = "AABCBDC";
    LRS(S);
    return 0;
}
```

Output:

```
Output

LRS Length: 3
LRS: ABC

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

LEET CODE:

https://leetcode.com/problems/longest-common-subsequence/description/

CODE:

```
int longestCommonSubsequence(char *text1, char *text2) {
   int m = strlen(text1);
   int n = strlen(text2);
   int dp[m + 1][n + 1];

   for (int i = 0; i <= m; i++)
        for (int j = 0; j <= n; j++)</pre>
```

Output:

