```
Practical 5
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Class-A-4
Batch-B3
Aim-
(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
#include <iostream>
#include <string>
using namespace std;
#include <iostream>
#include <string>
using namespace std;
string printLCS(string x, string y) {
  int m = x.length();
  int n = y.length();
  int dp[100][100] = \{0\};
  // Fill dp table
  for (int i = 1; i <= m; i++) {
    for (int j = 1; j \le n; j++) {
```

```
if (x[i-1] == y[j-1]) {
        dp[i][j] = dp[i - 1][j - 1] + 1;
      } else {
         dp[i][j] = max(dp[i - 1][j], dp[i][j - 1]);
      }
    }
  }
  // Reconstruct LCS from dp table
  int i = m, j = n;
  string lcsStr = "";
  while (i > 0 \&\& j > 0) {
    if (x[i-1] == y[j-1]) {
      lcsStr = x[i - 1] + lcsStr;
      i--;
      j--;
    i--;
    } else {
      j--;
    }
  }
  return lcsStr;
int main() {
  string x = "AGCCCTAAGGGCTACCTAGCTT";
  string y = "GACAGCCTACAAGCGTTAGCTTG";
```

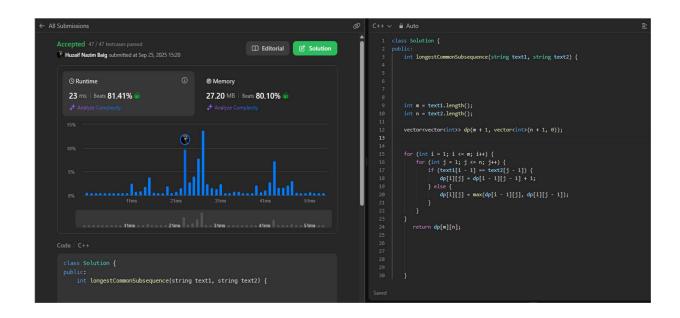
}

```
string lcsStr = printLCS(x, y);
cout << "The length of the LCS is " << lcsStr.length() << endl;
cout << "The LCS is: " << lcsStr << endl;
return 0;
}</pre>
```

## Output

```
The length of the LCS is 16
The LCS is: GCCCTAAGCTTAGCTT
```

## Leetcode Problem-



## LRS Part 2

Code-

```
#include <iostream>
#include <vector>
#include <string>
using namespace std;
int main() {
  string s1, s2;
  cin >> s1 >> s2;
  int n = s1.length();
  vector<vector<int>> dp(n + 1, vector < int > (n + 1, 0));
  vector < vector < char >> result(n + 1, vector < char > (n + 1, ' '));
  for (int i = 1; i <= n; i++) {
     for (int j = 1; j \le n; j++) {
       if (s1[i-1] == s2[j-1] \&\& i != j) {
          dp[i][j] = 1 + dp[i-1][j-1];
          result[i][j] = s1[i-1];
       } else {
          dp[i][j] = max(dp[i-1][j], dp[i][j-1]);
          result[i][j] = ' ';
       }
    }
  }
  cout << "DP Matrix:" << endl;</pre>
  for (int i = 0; i \le n; i++) {
     for (int j = 0; j \le n; j++) {
       cout << dp[i][j] << " ";
```

```
}
cout << endl;
}

cout << "Repeated Characters Matrix:" << endl;
for (int i = 0; i <= n; i++) {
    for (int j = 0; j <= n; j++) {
        cout << result[i][j] << " ";
    }
    cout << endl;
}

cout << "LRS Length: " << dp[n][n] << endl;
return 0;
}

Output-</pre>
```

```
0 0 0 0 0 0
            0 0
       1
         1
           1
 1 1 1 1 2
          2 2 2
 1 1 1 1 2 2 2 2
 1 1 2 2 2 2 2 2
 1 1 2 2 2 2
            2 2
Repeated Characters Matrix:
         В
     В
              D
LRS Length: 3
```