

DESIGN AND ANALYSIS OF ALGORITHM

PRACTICAL-6

NAME: VEDANT BHUTADA

ROLL: 69

BATCH: A4

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

Problem Statement: 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, final cost of LCS.

TASK-2: A subsequence of a given sequence is palindrome if it reads the same when read from left to right or right to left. Design an algorithm that take a sequence X[1...n]. Find all the possible palindrome sub-sequences for the given DNA sequence A C G T G T C A A A T C G

TASK-1

ITERATIVE

```
def print_matrix_with_direction(matrix, X, Y):
    m = len(X)
    n = len(Y)
    for i in range(m + 1):
        for j in range(n + 1):
            if i == 0 or j == 0:
                print(f"{matrix[i][j]:>2}h", end=" ")
            else:
                if X[i - 1] == Y[j - 1]:
                    direction = "d" # diagonal
                elif matrix[i - 1][j] >= matrix[i][j - 1]:
                    direction = "u" # up
                else:
                    direction = "s" # side
                print(f"{matrix[i][j]:>2}{direction}", end=" ")
        print()

def longest_common_subsequence(X, Y):
    m = len(X)
    n = len(Y)
    # Create a matrix to store the length of LCS for each subproblem
    dp = [[0] * (n + 1) for _ in range(m + 1)]

    for i in range(1, m + 1):
        for j in range(1, n + 1):
            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])

    # Retrieve the LCS length from the matrix
    lcs_length = dp[m][n]

    # Construct the LCS string
    lcs = "" * lcs_length
    i, j = m, n
    while i > 0 and j > 0:
        if X[i - 1] == Y[j - 1]:
            lcs[lcs_length - 1] = X[i - 1]
            i -= 1
            j -= 1
            lcs_length -= 1
        elif dp[i - 1][j] >= dp[i][j - 1]:
            i -= 1
        else:
            j -= 1

    return lcs, len(lcs), dp

X = "AGCCCTAAGGGCTACCTAGCTT"
Y = "GACAGCCTACAAGCGTTAGCTTG"
lcs, lcs_length, dp = longest_common_subsequence(X, Y)
```

```

LCS: AGCCCAAGGTTAGCTT
LCS Length: 16
Cost Matrix:
0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h
0h 0u 1d 1s 1d 1s 1s 1s 1s 1d 1s 1d 1d 1s 1s 1s 1s 1d 1s 1s 1s 1s 1s
0h 1d 1u 1u 1u 1u 2d 2s 2s 2s 2s 2s 2d 2s 2d 2s 2s 2s 2d 2s 2s 2s 2d
0h 1u 1u 2d 2s 2u 3d 3d 3s 3s 3d 3s 3s 3d 3s 3s 3s 3s 3d 3d 3s 3s 3d
0h 1u 1u 2d 2u 2u 3d 4d 4s 4s 4d 4s 4s 4s 4d 4s 4s 4s 4d 4s 4s 4s 4s
0h 1u 1u 2d 2u 2u 3d 4d 4u 4u 5d 5s 5s 5d 5s 5s 5s 5s 5d 5s 5s 5s
0h 1u 1u 2u 2u 2u 3u 4u 5d 5s 5u 5u 5u 5u 5u 5u 5d 6d 6s 6s 6d 6d 6s
0h 1u 2d 2u 3d 3s 3u 4u 5u 6d 6s 6d 6d 6s 6s 6s 6u 6u 7d 7s 7s 7s 7s
0h 1u 2d 2u 3d 3u 3u 4u 5u 6d 6u 7d 7d 7s 7s 7s 7s 7d 7u 7u 7u 7u
0h 1d 2u 2u 3u 4d 4s 4u 5u 6u 6u 7u 7u 8d 8s 8d 8s 8s 8s 8d 8s 8d
0h 1d 2u 2u 3u 4d 4u 4u 5u 6u 6u 7u 7u 8d 8u 9d 9s 9s 9s 9d 9s 9s
0h 1d 2u 2u 3u 4d 4u 4d 5u 6u 6u 7u 7u 8d 8u 9d 9u 9u 9u 10d 10s 10s 10d
0h 1u 2u 3d 3u 4u 5d 5d 5u 6u 7d 7u 7u 8u 9d 9u 9u 9u 10u 11d 11s 11s
0h 1u 2u 3u 3u 4u 5u 5u 6d 6u 7u 7u 7u 8u 9u 9u 10d 10d 10s 10d 12d 12s
0h 1u 2d 3u 4d 4u 5u 5u 6u 7d 7u 8d 8d 8u 9u 9u 10u 10u 11d 11s 11u 12u 12u
0h 1u 2u 3d 4u 4u 5d 6d 6u 7u 8d 8u 8u 8u 9d 9u 10u 10u 11u 11u 12u 12u
0h 1u 2u 3d 4u 4u 5d 6d 6u 7u 8d 8u 8u 8u 9d 9u 10u 10u 11u 11u 12u 12u
0h 1u 2u 3u 4u 4u 5u 6u 7d 7u 8u 8u 8u 8u 9u 9u 10d 11d 11u 11u 12u 13d 13d
0h 1u 2d 3u 4d 4u 5u 6u 7u 8d 8u 9d 9d 9s 9u 9u 10u 11u 12d 12s 12u 13u 13u
0h 1d 2u 3u 4u 5d 5u 6u 7u 8u 8u 9u 9u 10d 10s 10d 10u 11u 12u 13d 13s 13u 14d
0h 1u 2u 3d 4u 5u 6d 6d 7u 8u 9d 9u 9u 10u 11d 11s 11u 12u 13u 14s 14s 14u
0h 1u 2u 3u 4u 5u 6u 6u 7d 8u 9u 9u 9u 10u 11u 11u 12d 12d 12u 13u 14u 15d 15s
0h 1u 2u 3u 4u 5u 6u 6u 7d 8u 9u 9u 9u 10u 11u 11u 12d 13d 13s 13u 14u 15d 16d

```

```
def print_matrix_with_direction(matrix, X, Y):
    m = len(X)
    n = len(Y)
    for i in range(m + 1):
        for j in range(n + 1):
            if i == 0 or j == 0:
                print(f"{matrix[i][j]:>2}h", end=" ")
            else:
                if X[i - 1] == Y[j - 1]:
                    direction = "d" # diagonal
                elif matrix[i - 1][j] >= matrix[i][j - 1]:
                    direction = "u" # up
                else:
                    direction = "s" # side
                print(f"{matrix[i][j]:>2}{direction}", end=" ")
        print()

def longest_common_subsequence(X, Y):
    m = len(X)
    n = len(Y)

    dp = [[0] * (n + 1) for _ in range(m + 1)]

    for i in range(1, m + 1):
        for j in range(1, n + 1):
            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])

    lcs_length = dp[m][n]

    def construct_lcs(i, j):
        if i == 0 or j == 0:
            return []
        if X[i - 1] == Y[j - 1]:
            lcs = construct_lcs(i - 1, j - 1)
            lcs.append(X[i - 1])
            return lcs
        if dp[i - 1][j] >= dp[i][j - 1]:
            return construct_lcs(i - 1, j)
        else:
            return construct_lcs(i, j - 1)

    lcs = construct_lcs(m, n)
```

```

return lcs, lcs_length, dp

X = "AGCCCTAAGGGCTACCTAGCTT"
Y = "GACAGCCTACAAGCGTTAGCTTG"
lcs, lcs_length, dp = longest_common_subsequence(X, Y)

print("LCS:", "".join(lcs))
print("LCS Length:", lcs_length)
print("Cost Matrix:")
print_matrix_with_direction(dp, X, Y)
print("Final Cost of LCS:", dp[len(X)][len(Y)])

LCS: AGCCCAAGGTTAGCTT
LCS Length: 16
Cost Matrix:
0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h 0h
0h 0u 1d 1s 1d 1s 1s 1s 1s 1d 1s 1d 1d 1s 1s 1s 1s 1d 1s 1s 1s 1s
0h 1d 1u 1u 1u 2d 2s 2s 2s 2s 2s 2s 2d 2s 2d 2s 2s 2d 2s 2s 2s 2d
0h 1u 1u 2d 2s 2u 3d 3d 3s 3s 3d 3s 3s 3d 3s 3s 3s 3s 3d 3s 3s 3s
0h 1u 1u 2d 2u 2u 3d 4d 4s 4s 4d 4s 4s 4d 4s 4s 4s 4s 4d 4s 4s 4s
0h 1u 1u 2d 2u 2u 3d 4d 4u 4u 5d 5s 5s 5d 5s 5s 5s 5s 5d 5s 5s 5s
0h 1u 1u 2u 2u 2u 3u 4u 5d 5s 5u 5u 5u 5u 5u 5u 6d 6d 6s 6s 6d 6s
0h 1u 2d 2u 3d 3u 3u 4u 5u 6d 6s 6d 6d 6s 6s 6s 6u 6u 7d 7s 7s 7s
0h 1u 2d 2u 3d 3u 3u 4u 5u 6d 6u 7d 7d 7s 7s 7s 7s 7d 7u 7u 7u 7u
0h 1d 2u 2u 3u 4d 4s 4u 5u 6u 6u 7u 7u 8d 8s 8d 8s 8s 8s 8d 8s 8s 8d
0h 1d 2u 2u 3u 4d 4u 4u 5u 6u 6u 7u 7u 8d 8u 9d 9s 9s 9s 9d 9s 9s 9d
0h 1d 2u 2u 3u 4d 4u 4u 5u 6u 6u 7u 7u 8d 8u 9d 9u 9u 10d 10s 10s 10d 10d
0h 1u 2u 3d 3u 4u 5d 5d 5u 6u 7d 7u 7u 8u 9d 9u 9u 9u 10u 11d 11s 11s 11s
0h 1u 2u 3u 3u 4u 5u 5u 6d 6u 7u 7u 7u 8u 9u 9u 10d 10d 10s 10u 11u 12d 12d 12s
0h 1u 2d 3u 4d 4u 5u 6u 6u 7d 7u 8d 8d 8u 9u 9u 10u 10u 11d 11s 11u 12u 12u 12u
0h 1u 2u 3d 3u 4u 5d 6d 6u 7u 8d 8u 8u 9d 9u 10u 10u 11u 11u 12d 12u 12u 12u
0h 1u 2u 3d 4u 4u 5d 6d 6u 7u 8d 8u 8u 9d 9u 10u 10u 11u 11u 12d 12u 12u 12u
0h 1u 2u 3u 4u 4u 5u 6u 7d 7u 8u 8u 8u 9u 9u 10d 11d 11u 11u 12u 13d 13d 13s
0h 1u 2d 3u 4d 4u 5u 6u 7u 8d 8u 9d 9d 9s 9u 9u 10u 11u 12d 12s 12u 13u 13u 13u
0h 1d 2u 3u 4u 5d 5u 6u 7u 8u 8u 9u 9u 10d 10s 10d 10u 11u 12u 13d 13s 13u 14d
0h 1u 2u 3d 4u 5u 6d 6d 7u 8u 9d 9u 9u 10u 11d 11s 11s 11u 12u 13u 14d 14s 14u
0h 1u 2u 3u 4u 5u 6u 6u 7d 8u 9u 9u 9u 10u 11u 11u 12d 12d 12u 13u 14u 15d 15s
0h 1u 2u 3u 4u 5u 6u 6u 7d 8u 9u 9u 9u 10u 11u 11u 12d 13d 13s 13u 14u 15d 16d 16s
Final Cost of LCS: 16

```

TASK-2

```

def is_palindrome(subsequence):
    return subsequence == subsequence[::-1]

def find_palindrome_subsequences(sequence):
    def backtrack(start, path):
        if len(path) >= 2 and is_palindrome(path):
            palindromic_subsequences.add(path)
        if start == len(sequence):
            return
        for i in range(start, len(sequence)):
            path += sequence[i]
            backtrack(i + 1, path)
            path = path[:-1]

    palindromic_subsequences = set()
    backtrack(0, "")

    return list(palindromic_subsequences)

sequence = "ACGTGTCAAATCG"
palindrome_subsequences = find_palindrome_subsequences(sequence)
print("Palindromic Subsequences:")
for subsequence in palindrome_subsequences:
    print(subsequence)
print("Total Number of Palindromic Subsequences:", len(palindrome_subsequences))

TTT
CGC
GCG
CTATC
ATGTA
GCAAACG
TAAAT
GAG
AA
ACTCA

```

GTIG
TCT
GTCTG
TT
ACGTGCA
GTATG
GTTTG
ACGCA
AGA
CTGTC
CTTTC
CAAC
GG
GCAACG
GCCG
ATA
GCTCG
CGGC
TAT
GTG
CTC
GAAG
CGTGC
AGGA
GAAAG
AGTGA
GTAATG
ACTTCA
AAAA
GTAAATG
ATTA
CTTC
CC
CAAAC
GCACG
GGG
AAA
CTAATC
ACA
CAC
TGT
TAAT
ACCA
CCC

Total Number of Palindromic Subsequences: 59

