

BLG 336E – Analysis of Algorithms II
Homework 3

Local Sequence Alignment with Smith-Waterman

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Part 2. Report

2.1.a

```
// Local Alignment Runthrough

M = length of sequence 1 // lengthOfs1
N = length of sequence 2 // lengthOfs2
S = dynamic programming matrix of size M x N

// smith_waterman function

S[0,0] = 0
for i = 1 to M do
    S[i,0] = 0
for i = 1 to N do
    S[i,0] = 0
    for i = 1 to M
        0
        S[i-1,j-1] + match or mismatch
S[i,j] = max
        S[i-1,j] + gap
        S[i,j-1] + gap

return S[M,N]

// Local Alignment Traceback

// Traceback_part1
maxNumber = maximum value in S initialized with 0
for i = 0 to M do
    for i = 0 to N do
        if (S[i,j] > maxNumber)
            maxNumber = S[i,j]
return maxNumber

// Traceback_part2
targetNumber = maximum value number in S initialized with 0
for i = 0 to M do
    for i = 0 to N do
        if (S[i,j] == maxNumber)
            targetNumber++
return targetNumber++

// Traceback_part3
targetPosition[targetNumber] keeps every maxNumber row number.
targetLength[targetNumber] keeps each maxNumber diagonally from 0.
offset = offset counter for these arrays.

for i = 0 to M do
    for i = 0 to N do
        if (S[i,j] == maxNumber)
            tempi = i
            tempj = j
```

```

        while ( S[tempi, tempj] != 0)
            tempi = tempi -1
            tempj = tempj - 1
            targetLength[offset] = targetLength[offset] + 1
        targetLength[offset] = i
offset ++
return targetNumber and targetLength

```

For the desired output forms and file reading operations, the following operations have been made.

// Assignment

File reading line by line.

```

        push back for each line to vector
        sort the vector
for i = 0 to vector.size do
    for j = i + 1 to vector.size do
        s1 equal to vectors ith string
        s2 equal to vectors jth string
        call smith_waterman function
        call traceback_part1
        print Score = maxNumber
        call traceback_part2
        call_traceback_part3
        After detect target positions and length create substrings
        Create a set structure for keep substring without repetition.
        Print substrings

```

2.1.b.

Optimal alignment score for aligning M and N Time Complexity: $O(M*N)$

2.2.a

Calculations on brute force $4^{(m+n)}$ on the other hand, Smith-Waterman is processing as much as $m * n$.

2.2.b

Since Brute Force compares all possible substructures with each other, the smallest word substring will be maximum kept for it. This means that they are **single-length substrates, or letters**. Maximum kept equals **the number of letters of the lowest number of letters** among 2 sequences.

Smith-Waterman, on the other hand, has to keep all values in a matrix structure in order to reach the result. The size of the matrix is $m * n$.

2.2.c

Since the Bf algorithm is not applied in the study, this question can be answered according to the data in 1a. Time comparisons are also considered to be $4^{(m+n)}$ and $m * n$.