# Pairwise Sequence Alignment

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## 1 Part 1

$$B(m,n) = 1$$
, if  $m = 0 \lor n = 0$ .  
 $B(m,n) = B(m-1,n)$ ,  $B(m,n-1)$ ,  $B(m-1,n-1)$  if  $m \ge 1$ ,  $n \ge 1$ 

Base case: When one of m and n have zero length, there would be only one alignment. When both of them has zero length, it would be trivial to prove.

Inductive step: When a dash is added to the end of the string of length m, the number of alignments is B(m-1,n). When a dash is added to the end of the string of length n, the number of alignments is B(m,n-1). When a dash is not added at the end of either string, the number of alignments is B(m-1,n-1). B(m,n) equals the sum of these three.

## 2 Part 2

$$S[i,j] = \begin{cases} S[i,j-1] + M_{-,Y_j}, \text{ for } i = 0 \lor j > 0 \\ S[i-1,j] + M_{X_i,-} \text{ for } i > 0 \lor j = 0 \\ \max(S[i-1,j-1] + M_{X_i,Y_j}, S[i,j-1] + M_{-,Y_j}), S[i-1,j] + M_{X_i,-}) \text{ for } i > 0 \lor j > 0 \end{cases}$$

When i is at the boundary of the matrix, it means that the length of the first string is 0, then the value at [i,j] equals the value at [0,j-1] plus  $M_{-,X_i}$ .

When When j is at the boundary of the matrix, it means that the length of the first string is 0, then the value at [i,j] equals the value at [i-1,0] plus  $M_{X_i,j}$ .

When neither i or j is at the boundary, the value at [i,j] equals the maximum of the following: 1) value at [i-1, j-1], which is the optimal value for string lengths of i-1 and j-1, plus  $M_{X_i,Y_j}$ . 2) value at [i-1, j-1] plus  $M_{-,Y_j}$  3) value at [i-1, j-1] plus  $M_{X_i,Y_j}$ 

## 3 Part 3

#### **Algorithm 1:** ComputeGlobalAlignmentScores

**Input:** Sequences X and Y, and scoring matrix M **Output:** A optimal score matrix S

## **4** Part 4

#### **Algorithm 2:** ComputeAlignment

**Input:** Sequences X,Y, scoring matrix M, and an optimal matrix S **Output:** A sequence string representing the optimal alignment

```
1 X' \leftarrowan empty sequence;
2 Y' \leftarrowan empty sequence;
i \leftarrow |X|;
4 j \leftarrow |Y|;
5 while i \geq 0 and j \geq 0 do
       if S[i, j] = S[i - 1, j - 1] + M_{X_i, Y_i} then
            add X_i to the front of X';
            add Y_j to the front of Y';
8
           i \leftarrow i - 1;
9
          j \leftarrow j - 1;
10
       else if S[i, j-1] + M_{-,Y_j} then
11
            add – to the front of X';
12
           add Y_i to the front of Y';
13
          j \leftarrow j - 1;
14
       else if S[i-1,j] + M_{X_{i,-}} then
15
            add X_i to the front of X';
16
            add – to the front of Y';
17
           i \leftarrow i - 1;
18
  if i = 0 and j > 0 then
19
       add – to the front of X';
20
       add Y_i to the front of Y';
22 else if i = 0 and j > 0 then
       add X_i to the front of X';
       add – to the front of Y';
25 return (X',Y')
```

## **5** Part **5**

We can assume that sequences X and Y each have input sizes of m and n.

Running time of ComputeGlobalAlignmentScores:

The first two lines each take 1 operation. Within the innermost loop, each line takes 1 opera-

tion. So the two loops form O(m\*n) operation.

## Running time of ComputeAlignment:

The first four lines each take 1 operation. The while loop takes  $O(\min(m,n))$  operations because each line within takes constant operations, and the rest takes constant operations.

In total, the complexity of Global Alignment is O(mn).