## **Gotoh (Local) Unit Tests**

Hint: Many test values are taken from project Algorithms for Bioninformatics of Alexander Mattheis or the lectures.

## Test 1

## Input

Sequence A: CG Sequence B: CCGA

Gap Opening: -3

 $g(k) = -3 - k \cdot 1$ 

Enlargement: -1

Match: 1
Mismatch: -1

## Output

|   |   | С  | С  | G  | Α  |
|---|---|----|----|----|----|
|   | - | -∞ | -∞ | -∞ | -∞ |
| С | - | -4 | -4 | -4 | -4 |
| G | - | -3 | -3 | -4 | -4 |

|   |   | С | С | G | Α |
|---|---|---|---|---|---|
|   | 0 | 0 | 0 | 0 | 0 |
| С | 0 | 1 | 1 | 0 | 0 |
| G | 0 | 0 | 0 | 2 | 0 |

|   |    | С  | С  | G  | Α  |
|---|----|----|----|----|----|
|   | -  | -  | -  | -  | -  |
| С | -∞ | -4 | -3 | -3 | -4 |
| G | -∞ | -4 | -4 | -4 | -2 |

Seq1 CG \*\*

Seq2 CG

**Test 2-4:** Simulation of Smith-Waterman Tests by setting gap opening to 0.