Hirschberg

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

Test 1 (used Needleman-Waterman-Implementation for calculation)

Input

Sequence A: AATCG Sequence B: AACG

Deletion: 2
Insertion: 2
Match: -1
Mismatch: 1

Output

1. $Trace(A_1, A_2, T_3, C_4, G_5 | A_1, A_2, C_3, G_4)$ and i = ceil(5/2) = 3

		Α	Α	С	G
	0	2	4	6	8
Α	2	-1	1	3	5
Α	4	1	-2	0	2
T	6	3	0	-1	1
С	8	5	2	-1	0
G	10	7	4	1	-2

		Α	Α	С	G
	-2	1	4	7	10
Α	0	-1	2	5	8
Α	1	-1	0	3	6
T	2	0	-2	1	4
С	5	3	1	-1	2
G	8	6	4	2	0

Optimum: -2

	0	1	j=2	3	4
i=3	6	3	0	-1	1
	2	0	-2	1	4
Σ	8	3	-2	0	5

or: i = 5-3 = 2 in the matrix below and then reverse the line horizontally

		G	С	Α	Α
	0	2	4	6	8
G	2	-1	1	3	5
С	4	1	-2	0	2
Т	6	3	0	-1	1
Α	8	5	2	-1	-2 -2
Α	10	7	4	1	-2

	Α	Α	С	G
Α				
Α				
T				
С				
G				

1.1. $Trace(A_1, A_2|A_1, A_2)$:

and i = ceil(2/2) = 1

212177 0000 (11])112 111)112				
		Α	Α	
	0	2	4	
Α	2	-1	1	
Α	4	1	-2	

		Α	Α
	-2	1	4
Α	1	-1	2
Α	4	2	0

Optimum:



	0	j=1	2
	2	-1	1
i=1	1	-1	2
Σ	3	-2	3

	Α	Α
Α		
Α		

1.1.1. $Trace(|A_1|)$:

and	i	_	ceil	(a	/21	_	0
anu		=	сетт	ľ	/	=	٠.

1.1.1. I I acc (111).				
		Α		
	0	2		

	Α
2	0

Not shown!

1.1.2. $Trace(A_2|A_1, A_2)$:

and	i =	ceil(´1/2`) = 1 (local	for	that	matrix)

		Α	Α
	0	2	4
Α	2	-1	1
А		- 1	1

		Α	Α
	1	-1	2
Α	4	2	0

Optimum:

	0	j=1	2
	2	-1	1
i=1	4	2	0
Σ	6	1	1

(search minimum from right side, because right node?)

1.2. $Trace(C_4, G_5|A_2, C_3, G_4)$:

and i = ceil(2/2) = 1 (local for that matrix)

		Α	С	G
	0	2	4	6
С	2	1	1	3
G	4	3	2	0

		Α	С	G
	0	-2	1	4
С	3	1	-1	2
G	6	4	2	0

Optimum:

	0	2	j=3	4
i=4	2	1	1	3
	3	1	-1	2
Σ.	5	2	0	5

	Α	С	G
С			
G			

1.2.1. $Trace(|A_2, C_3)$:

	Α	С
0	2	4

and i = ceil(0/2) = 0 (local for that matrix)

Not shown!

1.2.2. $Trace(C_4|C_3, G_4)$:

and	i -	ceil(1/2)	- 1	(10031	for	that	matniv)	١.
ano	=	$(e_{11}(1/2)$) = 1	CLOCAL	TOI.	แกสเ	matr. TX)

		С	G
	0	2	4
G	2	1	1

		С	G
	1	-1	2
G	4	2	0

Optimum: 1

	0	2	j=4
i=5	2	1	1
	4	2	0
Σ	6	3	1

so:

	Α	Α	С	G
Α				
Α				
Т				
C G				
G				