

Dynamic programming

CMPSC 465 - Yana Safonova

Edit distance

Edit distance

Definition

The **edit distance** between x and y , denoted by $d(x, y)$, is the minimum number of insertions, deletions, and substitutions needed to transform x to y

$X = \text{PLACE}$
..|||

$Y = \text{SPACE}$

$\text{dist} = 2$ (two mismatches)

$X = \text{TOAD-}$
|.||

$Y = \text{TRADE}$

$\text{dist} = 2$ (1 mismatch, 1 insertion)

Edit distance and alignment

Alignment is a way of arranging DNA sequences to identify regions of similarity that show functional, structural, or evolutionary relationships between the sequences

distance = 3	0123456789	16	Mutations:
	ACGAGCGAGT	CGCGAG-	7: (A, C)
	.		9: (T, -)
	ACGAGCGC	G-CGAGC	16: (-, C)

Edit distance and alignment

Alignment is a way of arranging DNA sequences to identify regions of similarity that show functional, structural, or evolutionary relationships between the sequences

distance = 3	0123456789	16	Mutations:
	ACGAGCGAGTCGCGAG-		7: (A, C)
	.		9: (T, -)
	ACGAGCG CG -CGCGAG C		16: (-, C)

distance = 8	ACGAGCGAGTCGCGAG	
	
	ACGAGCG CGCGCGAGC	

Edit distance and alignment

Alignment is a way of arranging DNA sequences to identify regions of similarity that show functional, structural, or evolutionary relationships between the sequences

0123456789	16	Mutations:
ACGAGCGAGT	CGCGAG-	7: (A, C)
.		9: (T, -)
ACGAGCGC	G-CGAGC	16: (-, C)

Edit distance can be viewed as the number of non-matching operations in the best alignment

Edit distance can be computed using DP

Consider two strings

$$x = x_1 x_2 \cdots x_m \quad \text{and} \quad y = y_1 y_2 \cdots y_n$$

Edit distance can be computed using DP

Consider two strings

$$x = x_1 x_2 \cdots x_m \quad \text{and} \quad y = y_1 y_2 \cdots y_n$$

Subproblem: consider prefix $x_1 \cdots x_i$ and $y_1 \cdots y_j$ ($i \leq m, j \leq n$)

Define

$$E(i, j) = d(x_1 \cdots x_i, y_1 \cdots y_j)$$

Edit distance can be computed using DP

Consider two strings

$$x = x_1 x_2 \cdots x_m \quad \text{and} \quad y = y_1 y_2 \cdots y_n$$

Subproblem: consider prefix $x_1 \cdots x_i$ and $y_1 \cdots y_j$ ($i \leq m, j \leq n$)

Define

$$E(i, j) = d(x_1 \cdots x_i, y_1 \cdots y_j)$$

Optimal solution: $E(m, n)$

How to use the solution to the subproblems to solve $E(i, j)$?

Edit distance recurrence

Look at the rightmost column:

Edit distance recurrence

Look at the rightmost column:

Case 1	x_1	\cdots	x_{i-1}	x_i
	y_1	\cdots	y_j	-

Contributes 1 to the cost plus the cost of alignment

x_1	\cdots	x_{i-1}
y_1	\cdots	y_j

$$E(i, j) = 1 + E(i - 1, j)$$

Edit distance recurrence

Look at the rightmost column:

Case 1	x_1	\cdots	x_{i-1}	x_i
	y_1	\cdots	y_j	-

Contributes 1 to the cost plus the cost of alignment

x_1	\cdots	x_{i-1}
y_1	\cdots	y_j

$$E(i, j) = 1 + E(i - 1, j)$$

Case 2	x_1	\cdots	x_i	-
	y_1	\cdots	y_{j-1}	y_j

Contributes 1 to the cost plus the cost of alignment

x_1	\cdots	x_i
y_1	\cdots	y_{j-1}

$$E(i, j) = 1 + E(i, j - 1)$$

Edit distance recurrence

Look at the rightmost column:

Case 1	x_1	\cdots	x_{i-1}	$\textcolor{red}{x}_i$	
	y_1	\cdots	y_j	-	

Contributes 1 to the cost plus the cost of alignment

$$E(i, j) = 1 + E(i - 1, j)$$

Case 2	x_1	\cdots	x_i	-	
	y_1	\cdots	y_{j-1}	$\textcolor{red}{y}_j$	

Contributes 1 to the cost plus the cost of alignment

$$E(i, j) = 1 + E(i, j - 1)$$

Case 3	x_1	\cdots	x_{i-1}	$\textcolor{red}{x}_i$	
	y_1	\cdots	y_{j-1}	$\textcolor{red}{y}_j$	

$$E(i, j) = \begin{cases} E(i - 1, j - 1) & \text{if } x_i = y_j \\ 1 + E(i - 1, j - 1) & \text{otherwise} \end{cases}$$

Edit distance recurrence

The recurrence:

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

where

$$\text{diff}(i, j) = \begin{cases} 1 & \text{if } x_i \neq y_j \\ 0 & \text{otherwise} \end{cases}$$

Edit distance recurrence

The recurrence:

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

where

$$\text{diff}(i, j) = \begin{cases} 1 & \text{if } x_i \neq y_j \\ 0 & \text{otherwise} \end{cases}$$

Optimal solution: $E(m, n)$

Base case: $E(0, 0) = 0$, $E(i, 0) = i$, $E(0, j) = j$

Edit distance table

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

Edit distance table: filling base cases

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

$E(0, 0)$

$E(0, 1)$

\dots

$E(0, n - 1)$

$E(0, n)$

$E(1, 0)$

\vdots

$E(m - 1, 0)$

$E(m, 0)$

$m + 1$ rows

$n + 1$ columns

Edit distance table: computing $E(1, 1)$

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

$E(0, 0)$	$E(0, 1)$	\dots	$E(0, n - 1)$	$E(0, n)$
	\downarrow			
$E(1, 0)$	$E(1, 1)$			
⋮				
$E(m - 1, 0)$			$E(1, 1) = \min\{$ $1 + E(0, 1),$	
$E(m, 0)$				}

Edit distance table: computing $E(1, 1)$

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

$E(0, 0)$	$E(0, 1)$	\dots	$E(0, n - 1)$	$E(0, n)$
	\downarrow			
$E(1, 0)$	\rightarrow	$E(1, 1)$		
⋮				
$E(m - 1, 0)$			$E(1, 1) = \min\{$ $1 + E(0, 1),$ $1 + E(1, 0),$	
$E(m, 0)$				}

Edit distance table: computing $E(1, 1)$

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$

$E(0, 0)$	$E(0, 1)$	\dots	$E(0, n - 1)$	$E(0, n)$
	\searrow	\downarrow		
$E(1, 0)$	\rightarrow	$E(1, 1)$		
\vdots				
$E(m - 1, 0)$			$E(1, 1) = \min\{$	
$E(m, 0)$			$1 + E(0, 1),$	
			$1 + E(1, 0),$	
			$\text{diff}(1, 1) + E(0, 0)$	
			$\}$	

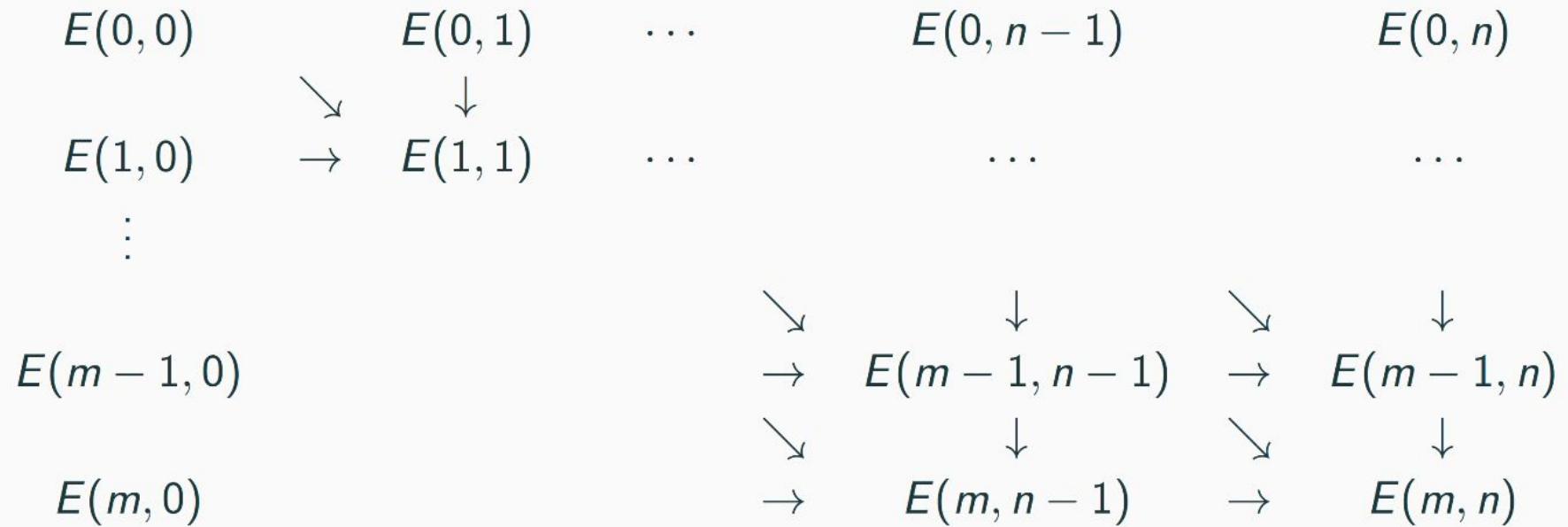
Edit distance table: completing row 1

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$



Edit distance table: the complete version

$$E(i, j) = \min\{1 + E(i - 1, j), 1 + E(i, j - 1), \text{diff}(i, j) + E(i - 1, j - 1)\},$$



Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

- - insertion to AGGT
- ↓ - insertion to ACGTA
- ↘ - match / mismatch

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

- - insertion to AGGT
- ↓ - insertion to ACGTA
- ↖ - match / mismatch

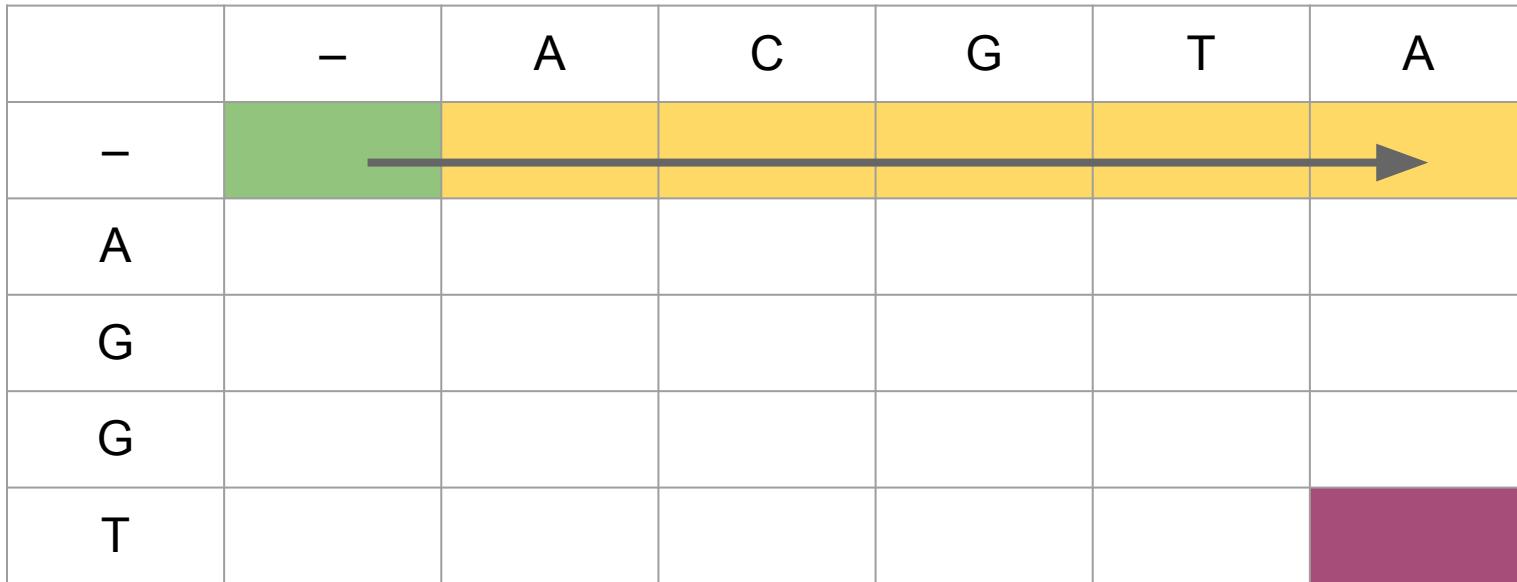
Each path from the green cell to the purple cell corresponds to an alignment

AC

--

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						



→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

ACGTA

Each path from the green cell to the purple cell corresponds to an alignment

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

```
graph LR; S1[ ] --> S2[ - ]; S2 --> S3[A]; S3 --> S4[C]; S4 --> S5[G]; S5 --> S6[T]; S6 --> S7[A]; S7 --> S8[G]; S8 --> S9[G]; S9 --> S10[T]; S10 --> S11[Purple];
```

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

ACGTA-
-----A

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

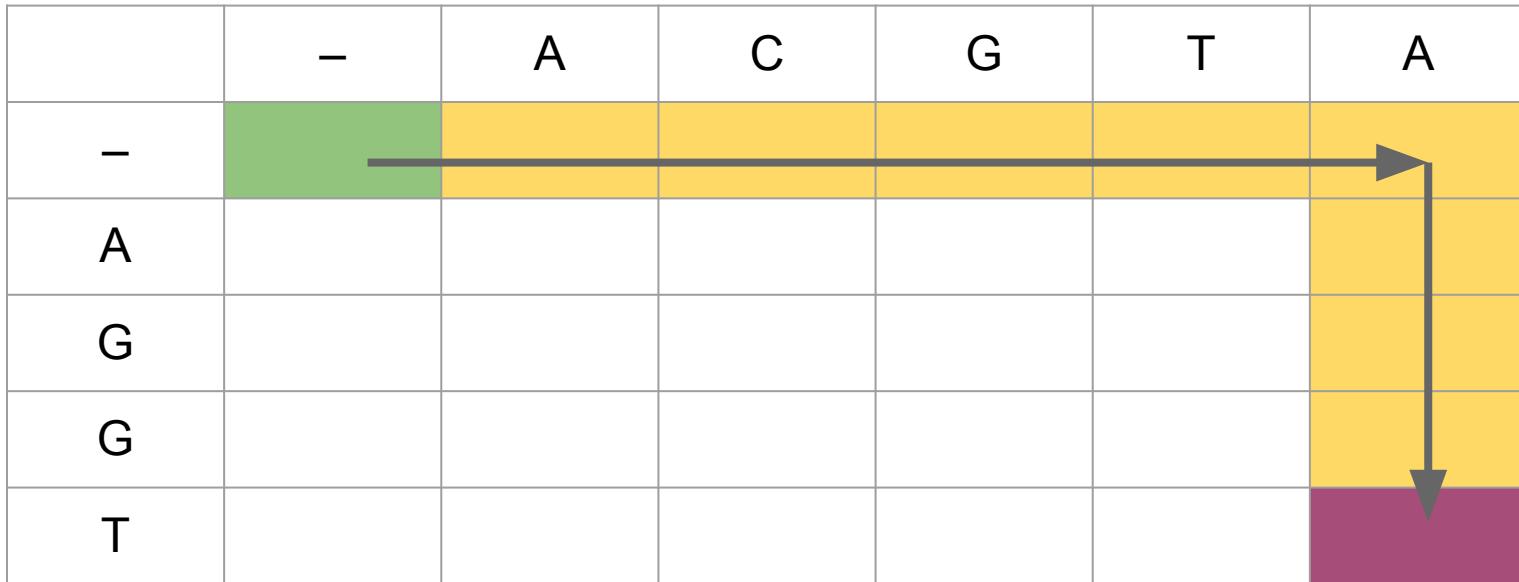
The diagram shows a sequence alignment between two strings. The top row contains the sequence: -, A, C, G, T, A. The bottom row contains the sequence: -, A, G, G, T, A. A path is drawn from the first cell (green) to the last cell (purple). The path starts at the green cell, moves horizontally to the right, and then vertically downwards to the purple cell. This path represents a specific alignment between the two sequences.

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

ACGTA--
-----AG

Example



→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Edit distance = 9

ACGTA - - -
- - - - AGGT

Each path from the green cell to the purple cell corresponds to an alignment

One of two worst alignments in terms of the edit distance

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

The diagram shows a 6x7 grid representing a dynamic programming table for sequence alignment. The columns are labeled with characters A, C, G, T, A and a blank space. The rows are labeled with characters - (indel), A, G, G, T. A green cell is at (-,-). A yellow path starts from it, moves right through (-,A), (A,A), (G,A), (G,A), and ends at (T,A) which is purple. A horizontal arrow points right from the end of the yellow path.

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Edit distance = 9

-----ACGTA
AGGT-----

Each path from the green cell to the purple cell corresponds to an alignment

Example

	-	A	C	G	T	A
-						
A						
G						
G						
T						

→ - insertion to AGGT
↓ - insertion to ACGTA
↘ - match / mismatch

Edit distance = 2

ACGTA
AGGT-

Each path from the green cell to the purple cell corresponds to an alignment

Is it the best alignment?

Example: base cases

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1					
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

Example

Indices / letters		0	1	2	3	4	5
0	-	0	1	2	3	4	5
1	A	1					
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{E(1, 0) + 1, E(0, 0) + d(A, A), E(0, 1) + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-	0	1	2	3	4	5
1	A	1					
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{1 + 1, 0 + 0, 1 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-	-	A	C	G	T	A
1	A	1	0	1	2	3	4
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{1 + 1, \textcolor{blue}{0 + 0}, 1 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	0	1	2	3	4	5
2	G	1	0				
3	G	2					
4	T	3					
5		4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

-A
A-

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	0	1	2	3	4	5
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

-A
A-

A
A

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	0	1	2	3	4	5
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
 ↓ - insertion to ACGTA
 ↘ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

-A
 A-

A
 A

A-
 -A

Example

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1	0				
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

A
A

The best choice for aligning prefixes of lengths 1 and 1

Example

Indices / letters		0	1	2	3	4	5
0	-	A	C	G	T	A	
1	A	1	0				
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
 ↓ - insertion to ACGTA
 ↘ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{E(1, 1) + 1, E(0, 1) + d(A, C), E(0, 2) + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-	0	1	2	3	4	5
1	A	1	0				
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{0 + 1, 1 + 1, 2 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	1	0	2	3	4	5
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{0 + 1, 1 + 1, 2 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	1	0	2	3	4	5
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

AC
A-

Example

Indices / letters		0	1	2	3	4	5
0	-		A	C	G	T	A
1	A	1	0	2	3	4	5
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

AC
A-

AC
-A

Example

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1	0	1			
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
 ↓ - insertion to ACGTA
 ↘ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

AC
A-

AC
-A

AC-
--A

Example

Indices / letters		0	1	2	3	4	5
0	-	-	A	C	G	T	A
1	A	1	0	1			
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

AC
A-

The best choice for aligning prefixes of lengths 2 and 1

Example

Indices / letters		0	1	2	3	4	5
0	-	A	C	G	T	A	
1	A	1	0	1			
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{E(1, 2) + 1, E(2, 0) + d(A, G), E(0, 3) + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-	A	C	G	T	A	
1	A	1	0	1			
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{1 + 1, 2 + 1, 3 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
0	-	A	C	G	T	A	
1	A	1	0	1	2	3	
2	G	2					
3	G	3					
4	T	4					

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

$$\min\{1 + 1, 2 + 1, 3 + 1\}$$

Example

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1	0	1	2	3	4
2	G	2	1	1	1	2	3
3	G	3	2	2	1	2	3
4	T	4	3	3	2	1	2

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

Example

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1	0	1	2	3	4
2	G	2	1	1	1	2	3
3	G	3	2	2	1	2	3
4	T	4	3	3	2	1	2

→ - insertion to AGGT
↓ - insertion to ACGTA
↖ - match / mismatch

Each path from the green cell to the purple cell corresponds to an alignment

edit distance between AGGT and ACGTA = 2

Pseudocode

```
def EDIT_DISTANCE(x, y):
    for i = 0, . . . , m:
        E(i, 0) = i;
    for j = 0, . . . , n:
        E(0, j) = j;
    for i = 1, . . . , m:
        for j = 1, . . . , n:
            E(i, j) =
                min{1 + E(i - 1, j), 1 + E(i, j - 1), diff(i, j) + E(i - 1, j - 1)};
    return E(m, n);
```

Running time: $O(mn)$
Memory usage: $O(mn)$

Example: how to reconstruct the best alignment?

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0	1	2	3	4	5
1	A	1	0	1	2	3	4
2	G	2	1	1	1	2	3
3	G	3	2	2	1	2	3
4	T	4	3	3	2	1	2

Edit distance modification

We use an extra table prev to record where each entry of $E(i, j)$ was coming from:

$$\text{prev}(i, j) = \begin{cases} (i - 1, j) & \text{if } E(i, j) = 1 + E(i - 1, j) \\ (i, j - 1) & \text{if } E(i, j) = 1 + E(i, j - 1) \\ (i - 1, j - 1) & \text{if } E(i, j) = \text{diff}(i, j) + E(i - 1, j - 1) \end{cases}$$

backtracking

Example: how to reconstruct the best alignment?

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0 (-, -)	1 (0, 0)	2 (0, 1)	3 (0, 2)	4 (0, 3)	5 (0, 4)
1	A	1 (0, 0)	0 (0, 0)	1 (1, 1)	2 (1, 2)	3 (1, 3)	4 (1, 4)
2	G	2 (1, 0)	1 (1, 1)	1 (1, 1)	1 (1, 2)	2 (2, 3)	3 (2, 4)
3	G	3 (2, 0)	2 (2, 1)	2 (2, 2)	1 (2, 2)	2 (2, 3)	3 (3, 4)
4	T	4 (3, 0)	3 (3, 1)	3 (3, 1)	2 (3, 3)	1 (3, 3)	2 (4, 4)

Example: how to reconstruct the best alignment?

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0 (-, -)	1 (0, 0)	2 (0, 1)	3 (0, 2)	4 (0, 3)	5 (0, 4)
1	A	1 (0, 0)	0 (0, 0)	1 (1, 1)	2 (1, 2)	3 (1, 3)	4 (1, 4)
2	G	2 (1, 0)	1 (1, 1)	1 (1, 1)	1 (1, 2)	2 (2, 3)	3 (2, 4)
3	G	3 (2, 0)	2 (2, 1)	2 (2, 2)	1 (2, 2)	2 (2, 3)	3 (3, 4)
4	T	4 (3, 0)	3 (3, 1)	3 (3, 1)	2 (3, 3)	1 (3, 3)	2 (4, 4)

Edit distance modification

We use an extra table prev to record where each entry of $E(i, j)$ was coming from:

$$\text{prev}(i, j) = \begin{cases} (i - 1, j) & \text{if } E(i, j) = 1 + E(i - 1, j) \\ (i, j - 1) & \text{if } E(i, j) = 1 + E(i, j - 1) \\ (i - 1, j - 1) & \text{if } E(i, j) = \text{diff}(i, j) + E(i - 1, j - 1) \end{cases}$$

def PRING_ALIGNMENT(x, y, prev):

 Set $i = m, j = n$;

while $i \geq 1$ and $j \geq 1$:

if $\text{prev}(i, j) = (i - 1, j - 1)$:

 print_back(y_i) _{x_i} ;

$i = i - 1, j = j - 1$;

if $\text{prev}(i, j) = (i - 1, j)$:

 print_back(\cdot) _{x_i} ;

$i = i - 1$;

if $\text{prev}(i, j) = (i, j - 1)$:

 print_back(y_j) _{\cdot} ;

$j = j - 1$;

Example: how to reconstruct the best alignment?

Indices / letters		0	1	2	3	4	5
		-	A	C	G	T	A
0	-	0 (-, -)	1 (0, 0)	2 (0, 1)	3 (0, 2)	4 (0, 3)	5 (0, 4)
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ACGTA
| . ||
AGGT-