

# Approximate pattern matching

Lecture 05.01

*by Marina Barsky*

# Sequence similarity

- The biological sequences encode and reflect higher-level molecular structures and mechanisms
- **In bimolecular sequences (DNA, RNA or protein), high sequence similarity usually implies significant structural and functional similarity**
- A tractable, though partly heuristic way to infer the structure and function of an unknown protein is to search for the similar known proteins at the sequence level

# Similar but not identical!

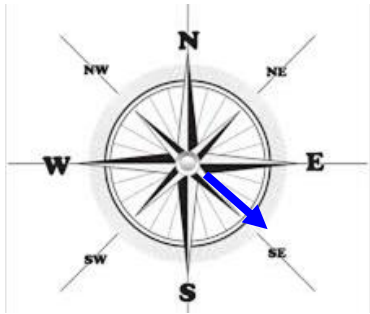
- We are looking for sequences that are similar to each other
- However they are never exactly the same due to small changes accumulated over generations
- How do we define and measure similarity?

# Approximate pattern matching

- Approximate – means some errors are allowed in valid matches
- The shift is accompanied by a shift in technique:  
*dynamic programming*

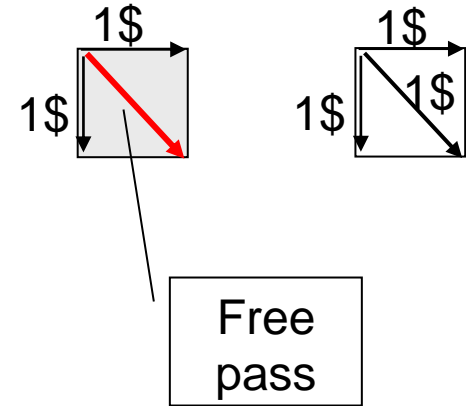
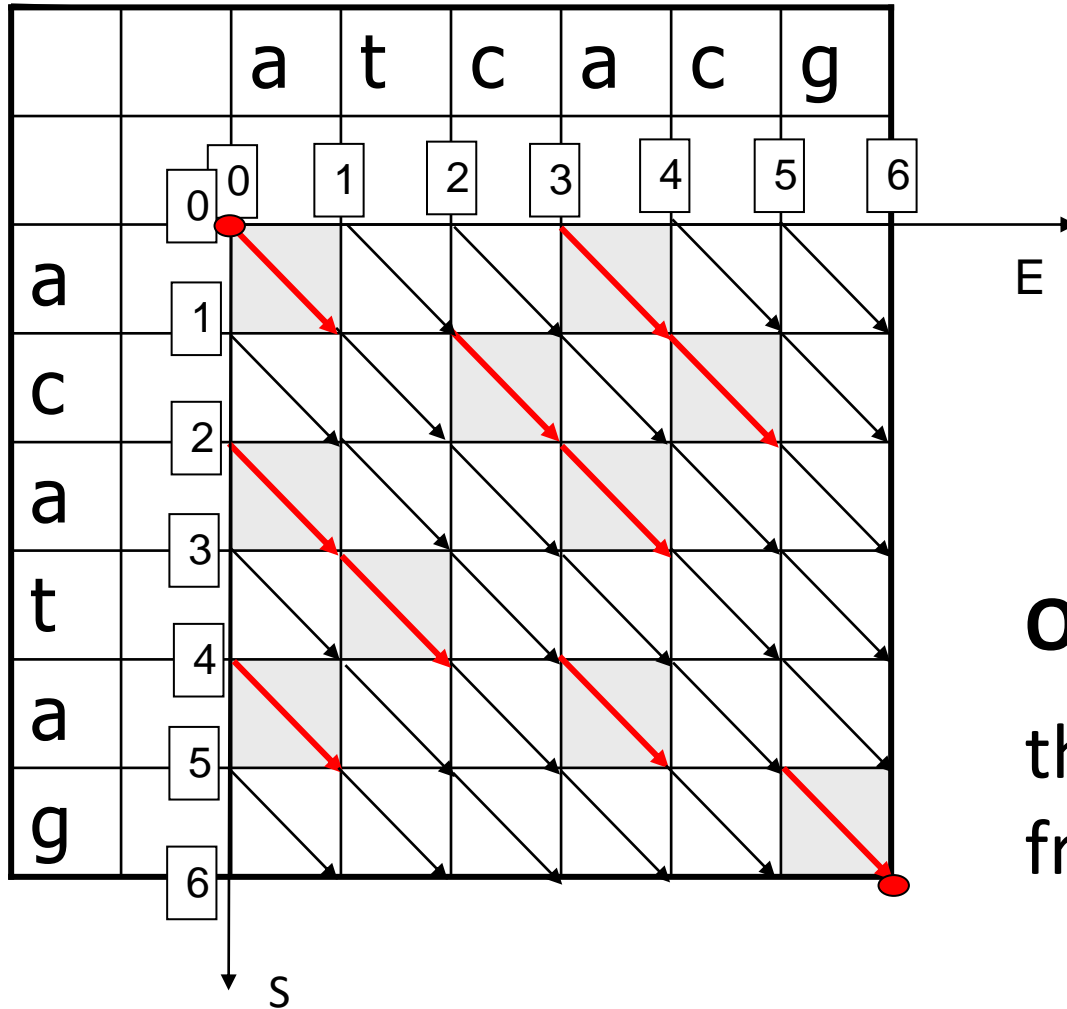
# Dynamic Programming

The main tool in approximate pattern matching



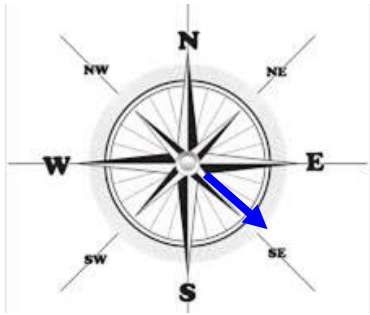
# Problem: the cheapest path in a special grid

**Input:**

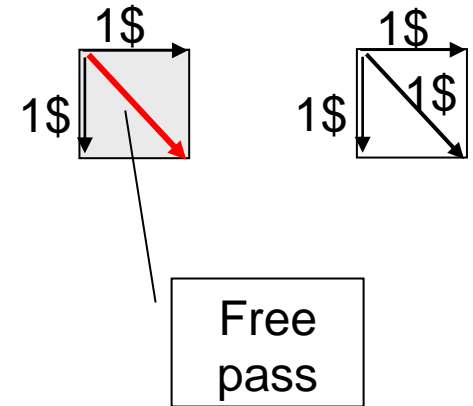
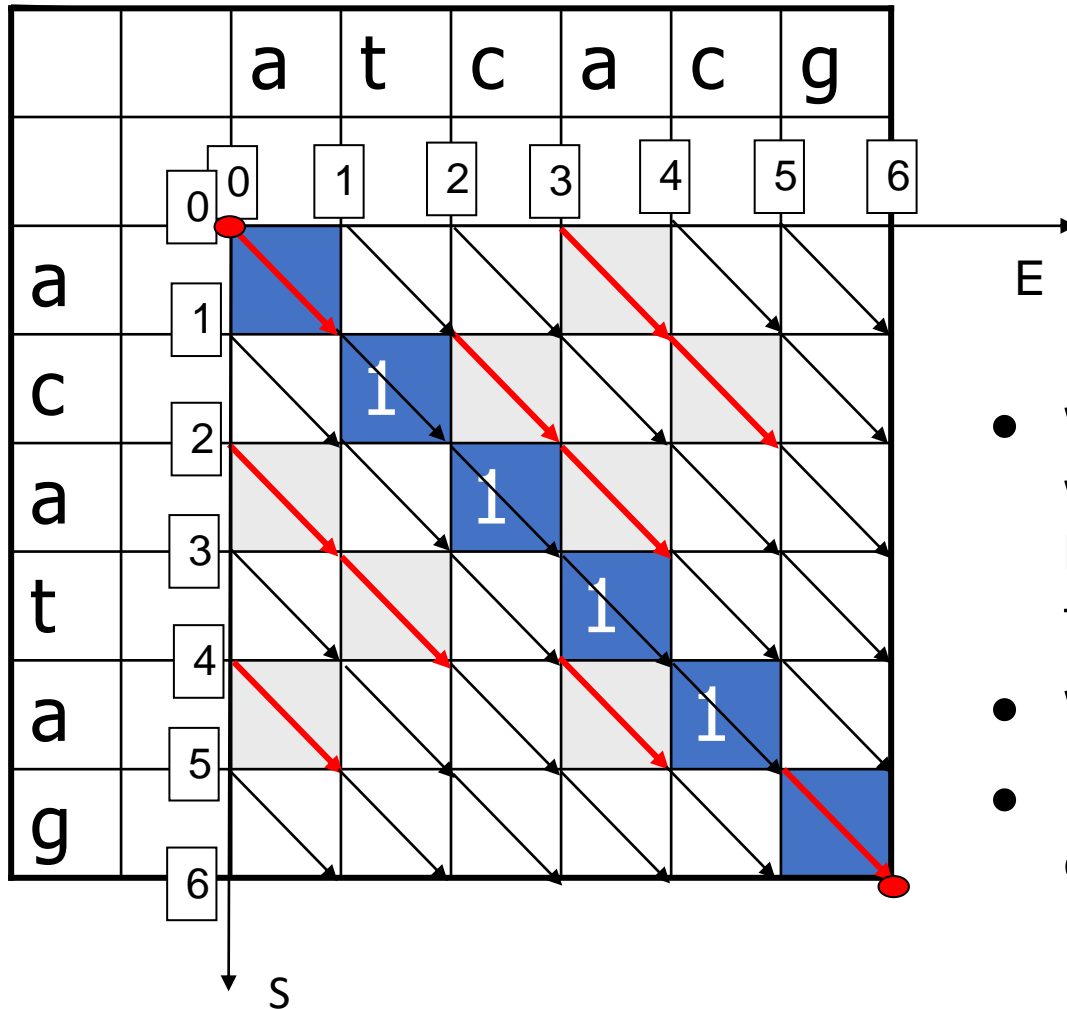


**Output:**

the cheapest path  
from (0,0) to (6,6)

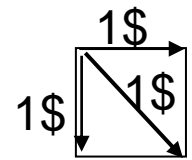
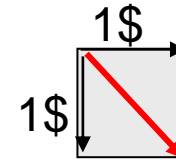
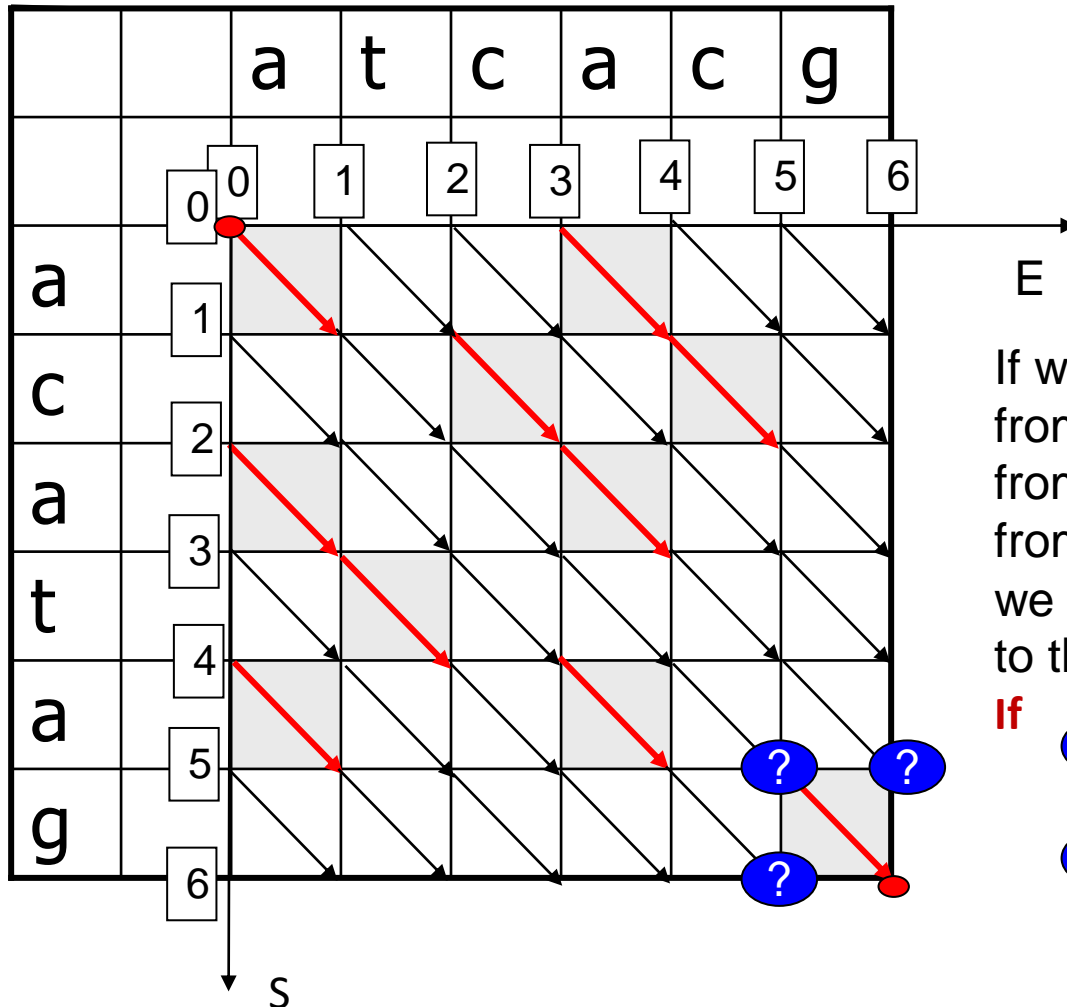
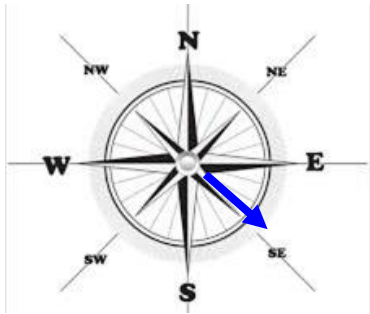


# Without the map

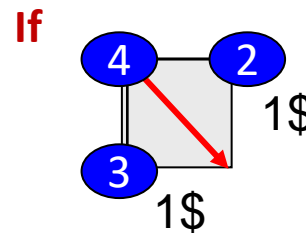


- Without additional information, we will always head South-East hoping to reach the destination faster
- We will pay 4\$
- However a better (cheaper) path exists with more free cells

# Sub-problems approach

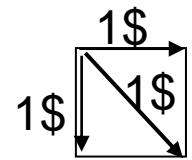
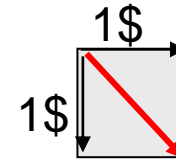
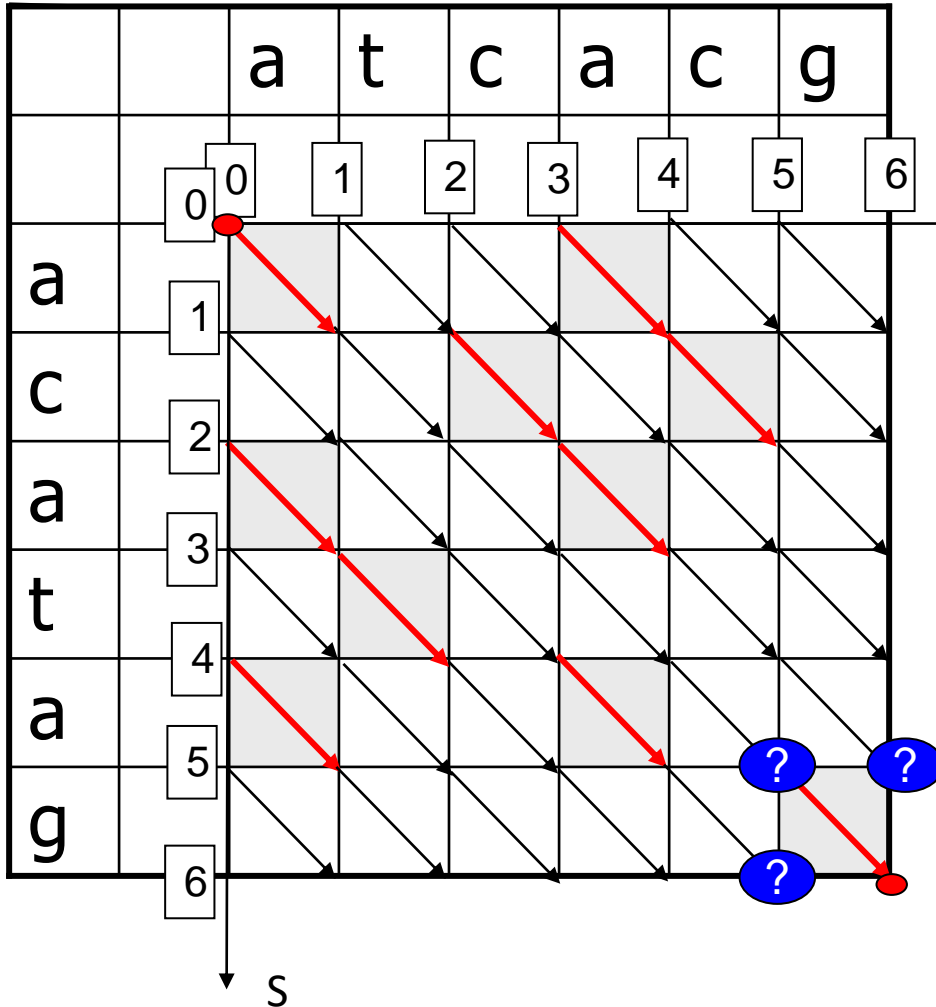
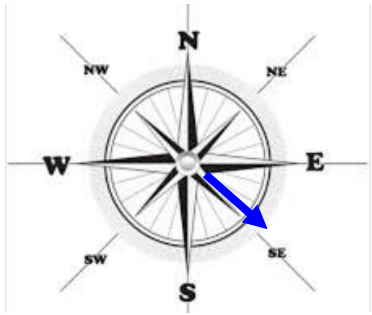


If we knew the cheapest paths  
 from (0,0) to (5,5)  
 from (0,0) to (6,5)  
 from (0,0) to (5,6)  
 we could choose the best last step  
 to the destination:

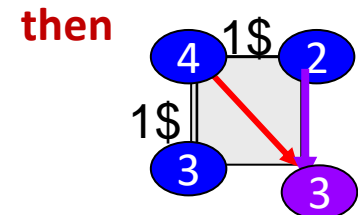
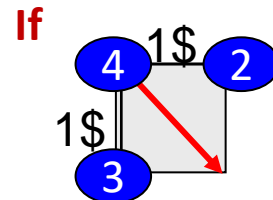




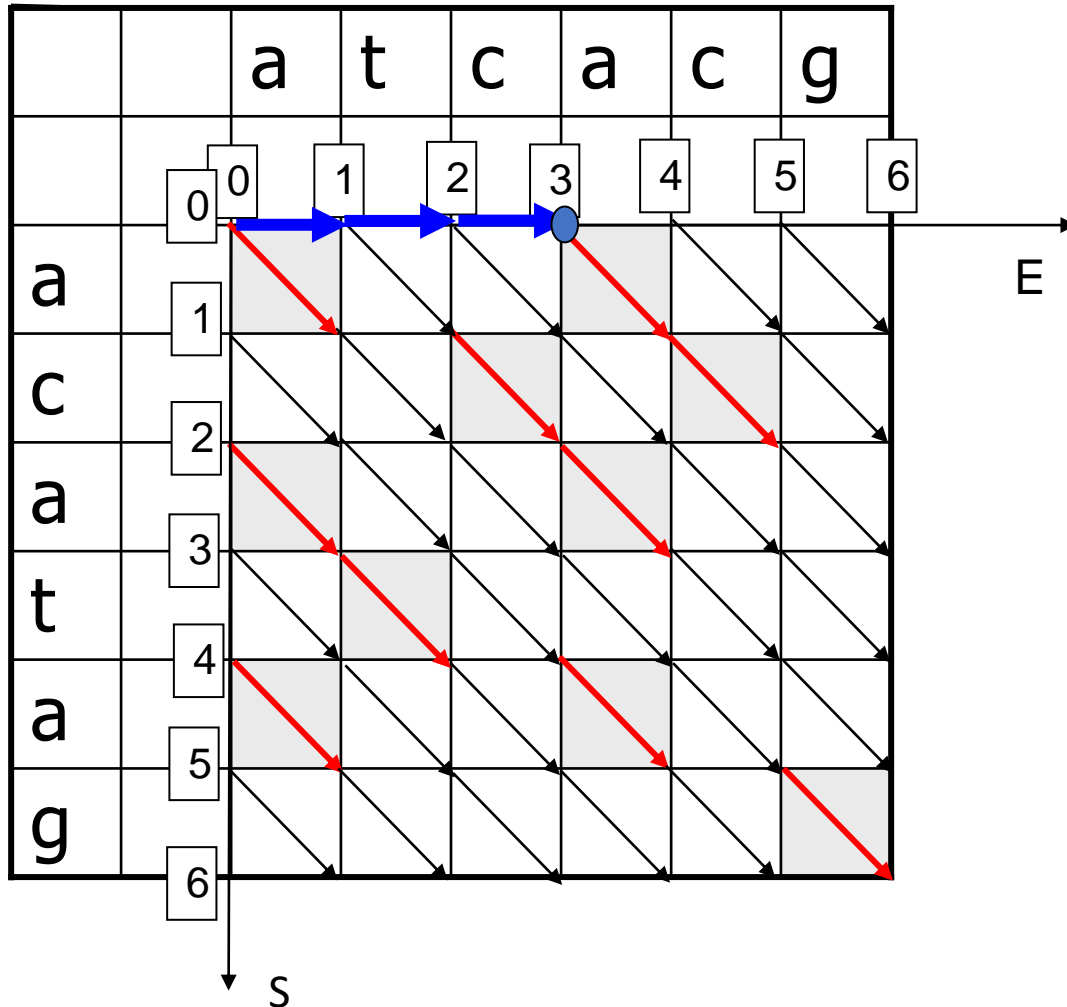
# Sub-problems approach



E  
And this is true for any cell – what path to choose depends on the cheapest paths to the left, upper, and upper-left corner.  
Since we choosing only 1 step, we can take the min of the result



# Recurrence relation – base condition



When  $i=0$ , there is no cheaper way of going from  $(0,0)$  to  $(0,j)$  than to pay  $j\$$  - heading strictly to the right (East)

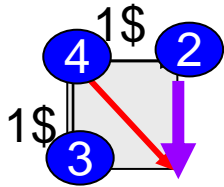
The same for  $j=0$ .

The base condition:

if  $i=0$  then  $COST(i,j)=j$

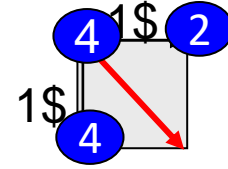
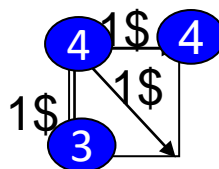
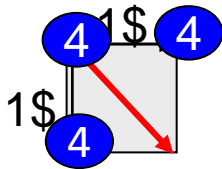
if  $j=0$  then  $COST(i,j)=i$

# Recurrence relation (for $i > 0$ and $j > 0$ )



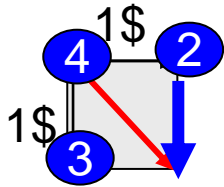
$$\text{COST}(i,j) = \min$$

$$\left\{ \begin{array}{l} \text{COST}(i-1,j)+1 \\ \text{COST}(i,j-1)+1 \\ \text{COST}(i-1,j-1)+\text{DIAGONAL}(i,j) \end{array} \right.$$



For each case, what is the best choice?

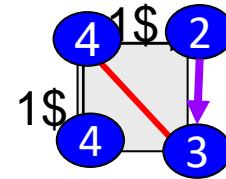
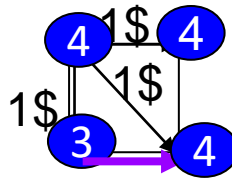
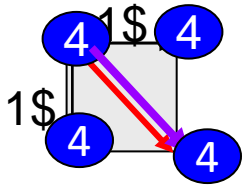
# Recurrence relation (for $i > 0$ and $j > 0$ )



$$\text{COST}(i,j) = \min$$

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For each case, what is **the best choice**?



# Recursive algorithm

$$\text{COST}(i,j) = \min \begin{cases} \text{COST}(i-1,j)+1 \\ \text{COST}(i,j-1)+1 \\ \text{COST}(i-1,j-1)+\text{DIAGONAL}(i,j) \end{cases}$$

**algorithm** *cheapestPath* ( array *diagonalCost*, *N*, *M* )

**return** *cost* ( *N*, *M*, *diagonalCost* )

**algorithm** *cost* ( *i*, *j*, *diagonalCost* )

**if** *i*=0 **then**

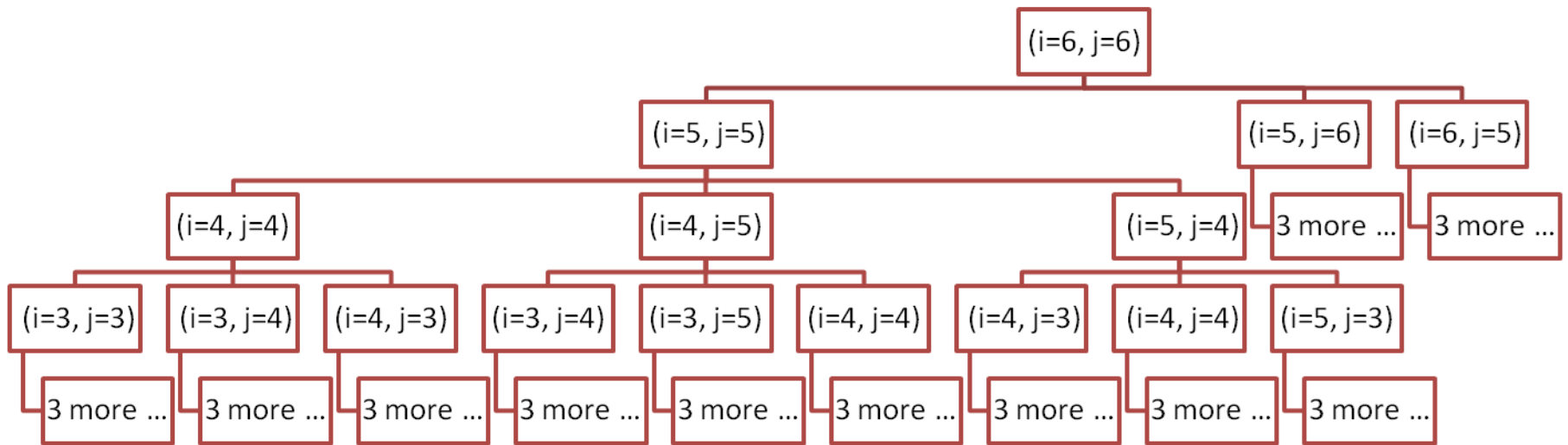
**return** *j*

**if** *j*=0 **then**

**return** *i*

**return** **min** ( *cost* ( *i*-1, *j* ) +1, *cost* ( *i*, *j*-1 ) +1, *cost* ( *i*-1, *j*-1 ) + *diagonalCost* [ *i* ] [ *j* ] )

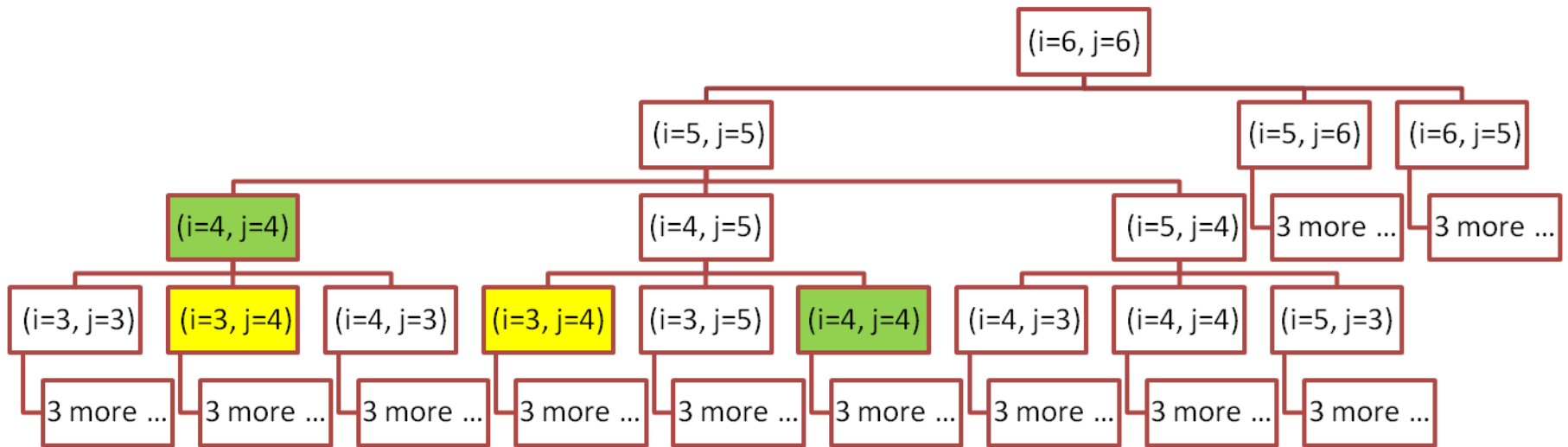
# The recursion tree: $O(3^N)$



$O(3^N)$  ?

But there are only  $N \cdot M$  different combinations  $(i, j)$ !

# Recursive algorithm: $O(3^N)$



The algorithm is exponential in  $N$  because **we call the recursive function multiple times with the same parameters!**

# Idea 1: store intermediate results

- Store the results of the  $cost(i,j)$  in a 2D table – so they do not need to be recomputed when needed again
- There are at most  $N^2$  different combinations of  $(i,j)$
- For each combination of  $(i,j)$  we compute the  $cost(i,j)$  only once
- When we need  $cost(i,j)$  again, we first check if it is already computed
- This gives a total running time  $O(N^2)$
- The method of storing the results of recursive calls in a lookup table is called ***recursion with memoization***



# Idea 2: The bottom-up computation

- In this problem we would need to compute the cost for all combinations of  $(i, j)$
- Instead of starting from  $\text{cost}(N, M)$  - fill in the best values for each cell of  $N * M$  table **starting from the lowest values**

# The bottom-up computation

- Create a table of size  $(N \times M)$  to store results of  $\text{cost}(i, j)$  for each  $0 \leq i \leq N$  and  $0 \leq j \leq M$
- First, fill-in the basic values of recursion – for  $i=0$  and for  $j=0$
- Apply recursive formula for computing the value of each cell from the lowest numbers of  $i$  and  $j$  to the highest (by rows or by columns)
- At the end, we will have the cost of the best path in the cell  $(N, M)$

# The recurrence relation: stays the same

The base condition:

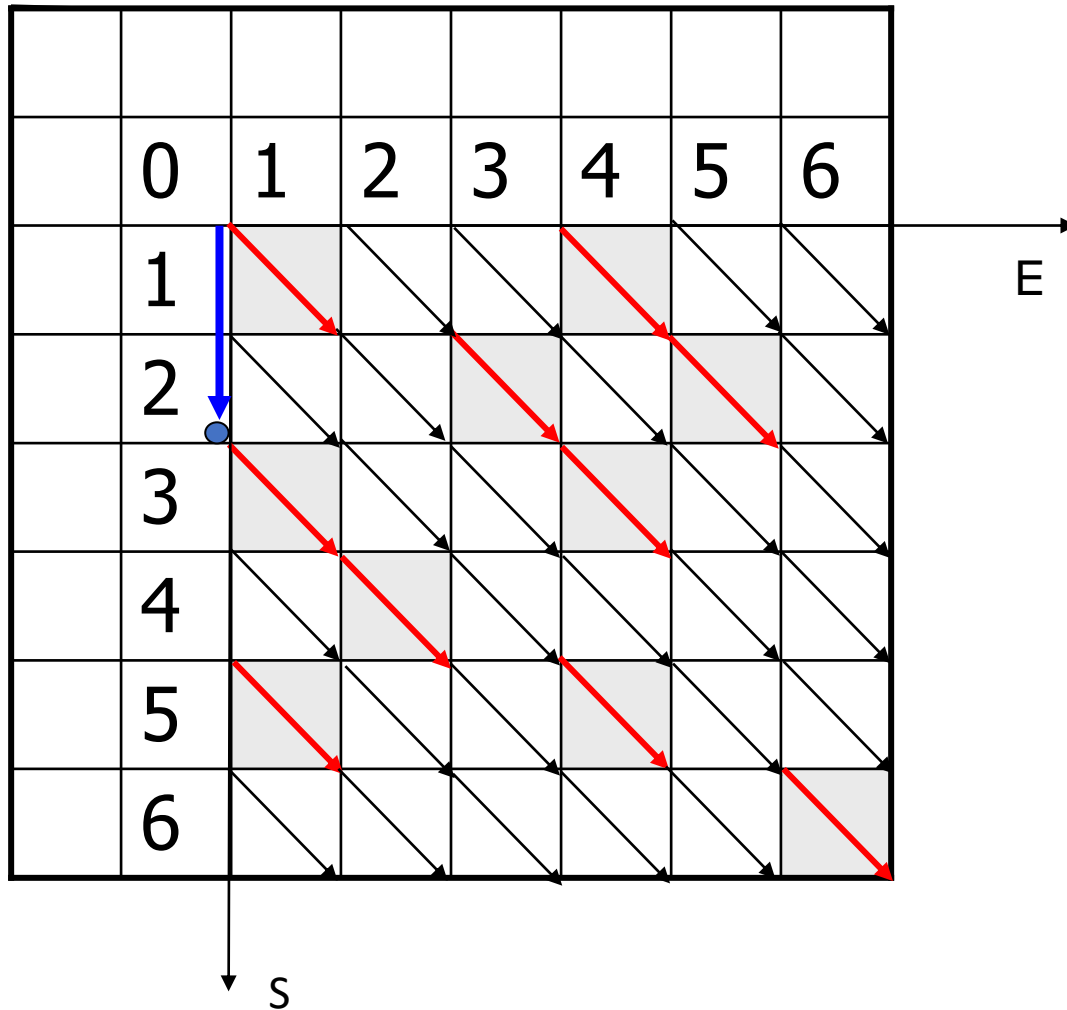
if  $i=0$  then  $COST(i,j)=j$   
if  $j=0$  then  $COST(i,j)=i$

The main relation ( for  $i>0$  and  $j>0$ )

$$COST(i,j)=\min \begin{cases} COST(i-1,j)+1 \\ COST(i,j-1)+1 \\ COST(i-1,j-1)+DIAGONAL(i,j) \end{cases}$$

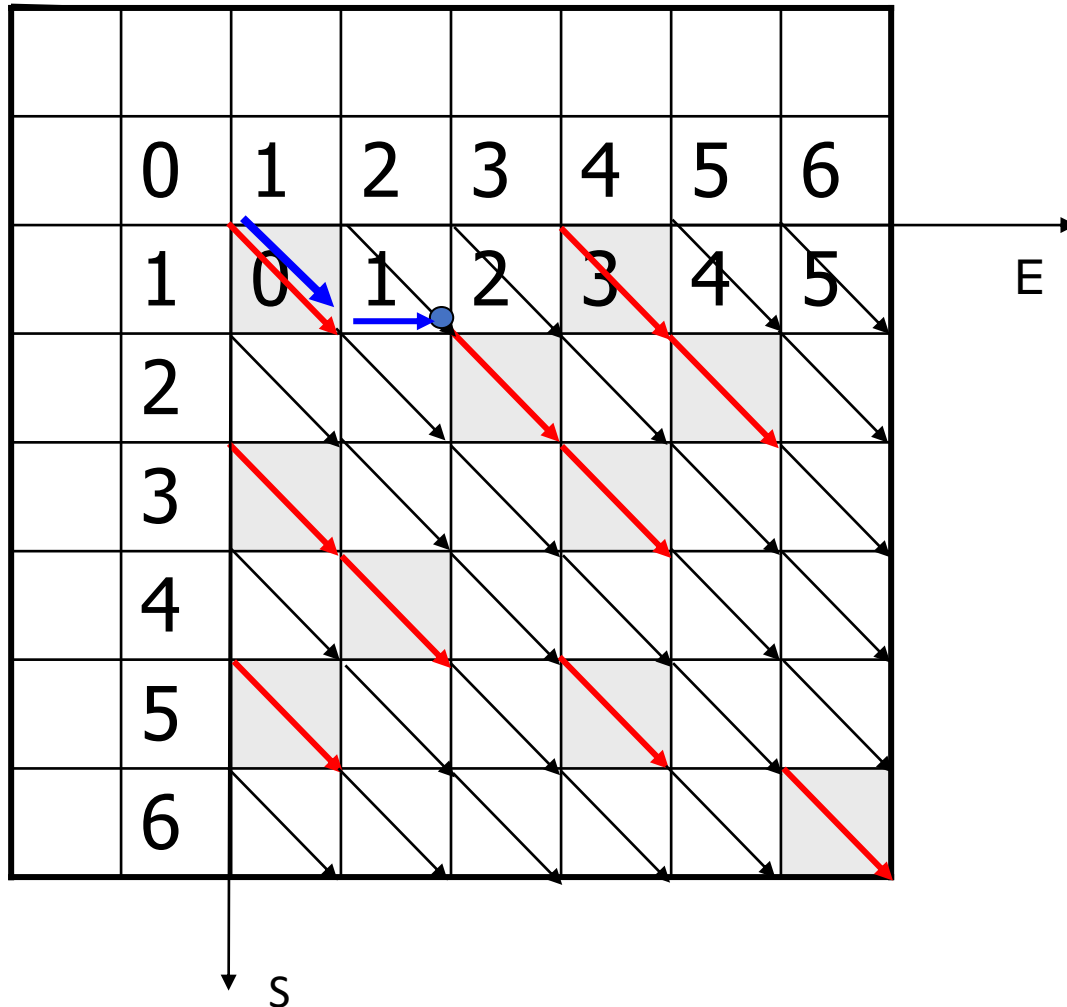
**We change:**  
**the order of computation**

Fill values for  $i=0$  and for  $j=0$   
(the base recursion condition)



There is no cheaper way  
of going to the point  
(2,0) than paying 2 \$

# Fill values for $i=1$ (from left to right)



Cell(1,2)=1

since the cheapest possible way is to continue the free path through the cell (1,1)

Fill the entire table  
(left-to-right top-down)

	0	1	2	3	4	5	6
1	0	1	2	3	4	5	
2	1	1	1	2	3	4	
3	2	2	2	1	2	3	
4	3	2	3	2	2	3	
5	4	3	3	3	3	3	
6	5	4	4	4	4	4	3

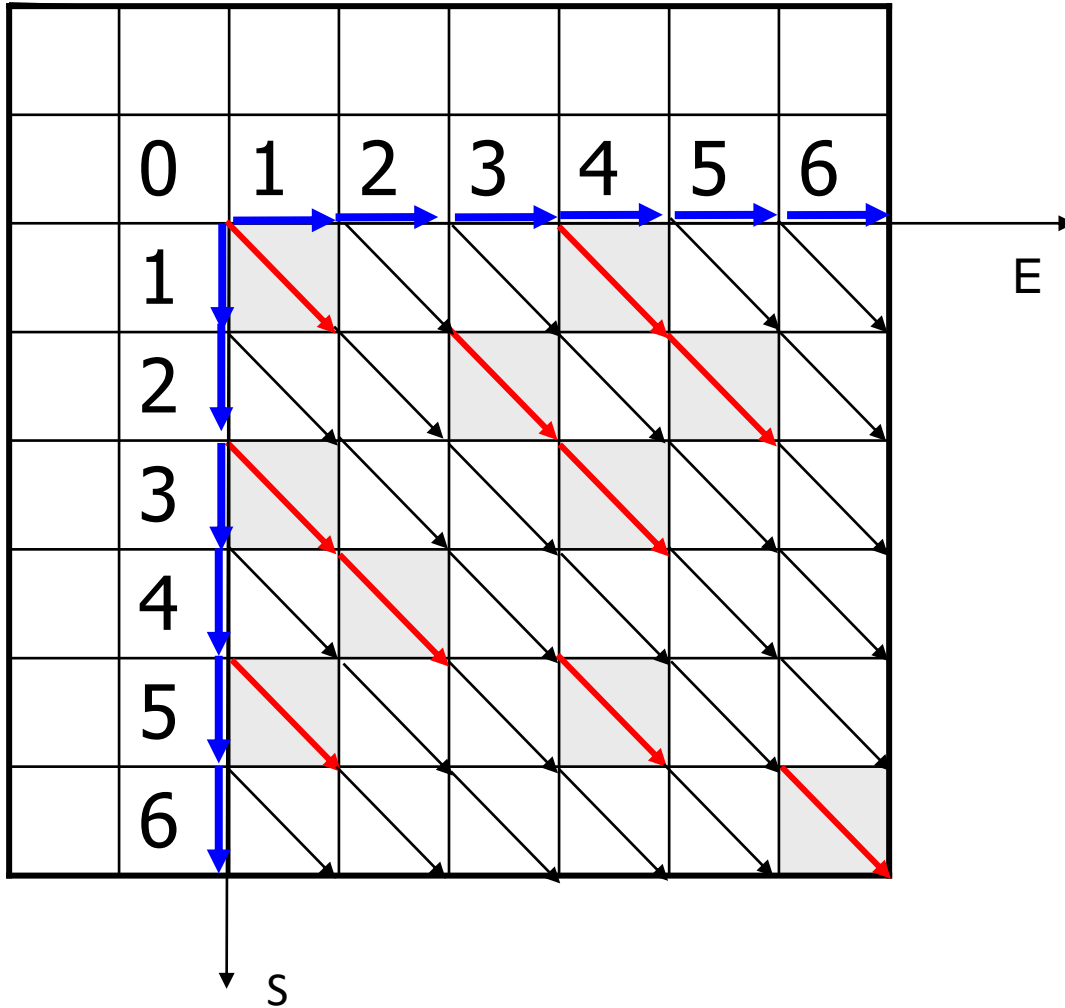
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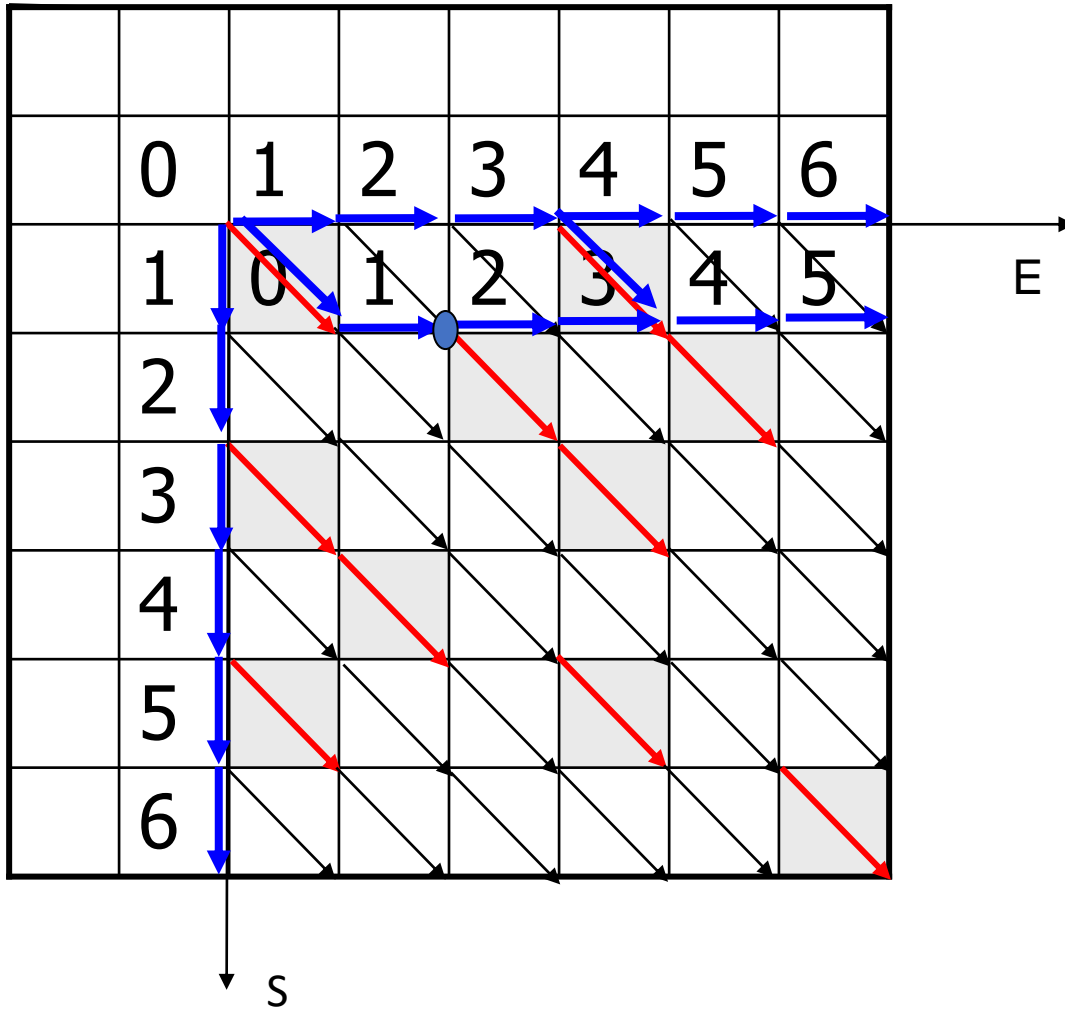
The overall cheapest possible path costs 3\$

**But what is this path?**

# Keeping track of the source

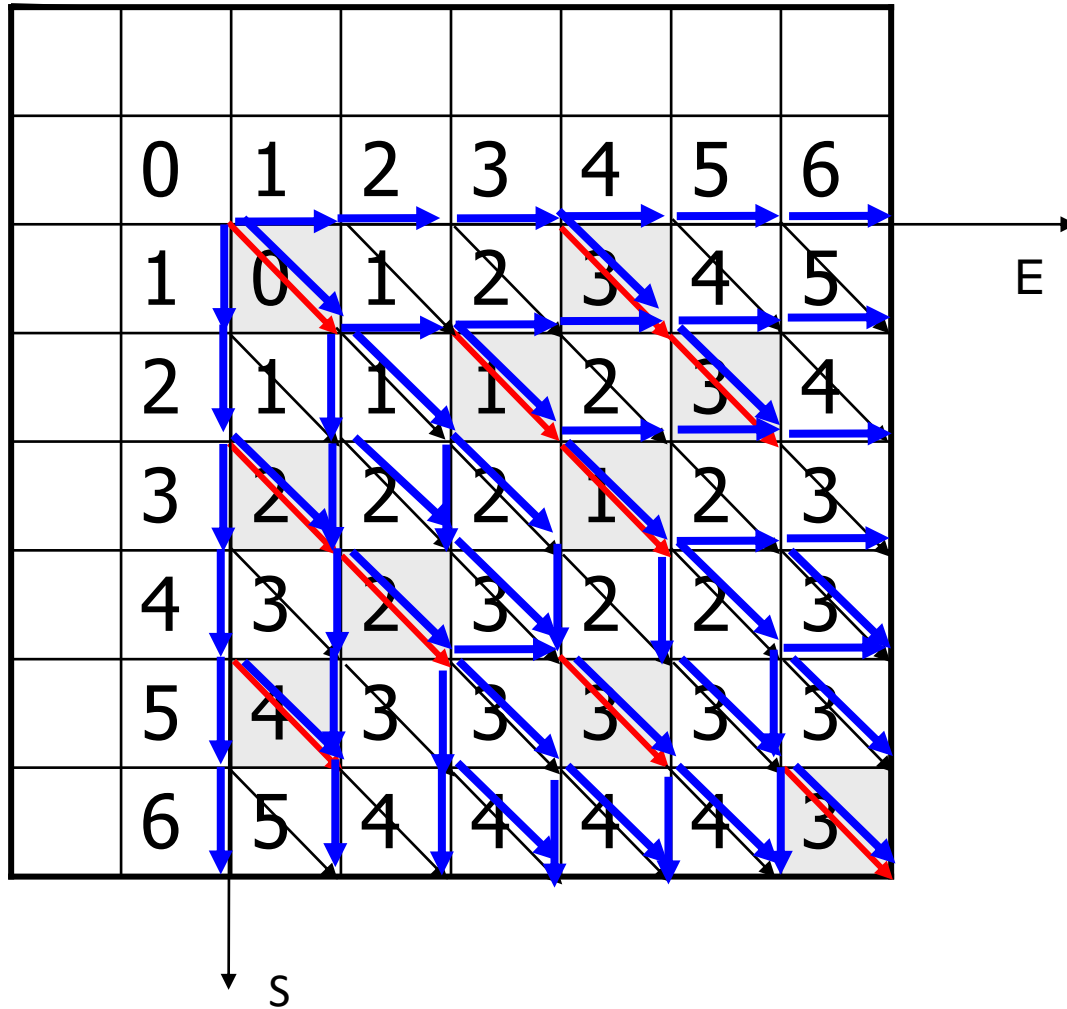


# Keeping track of the source

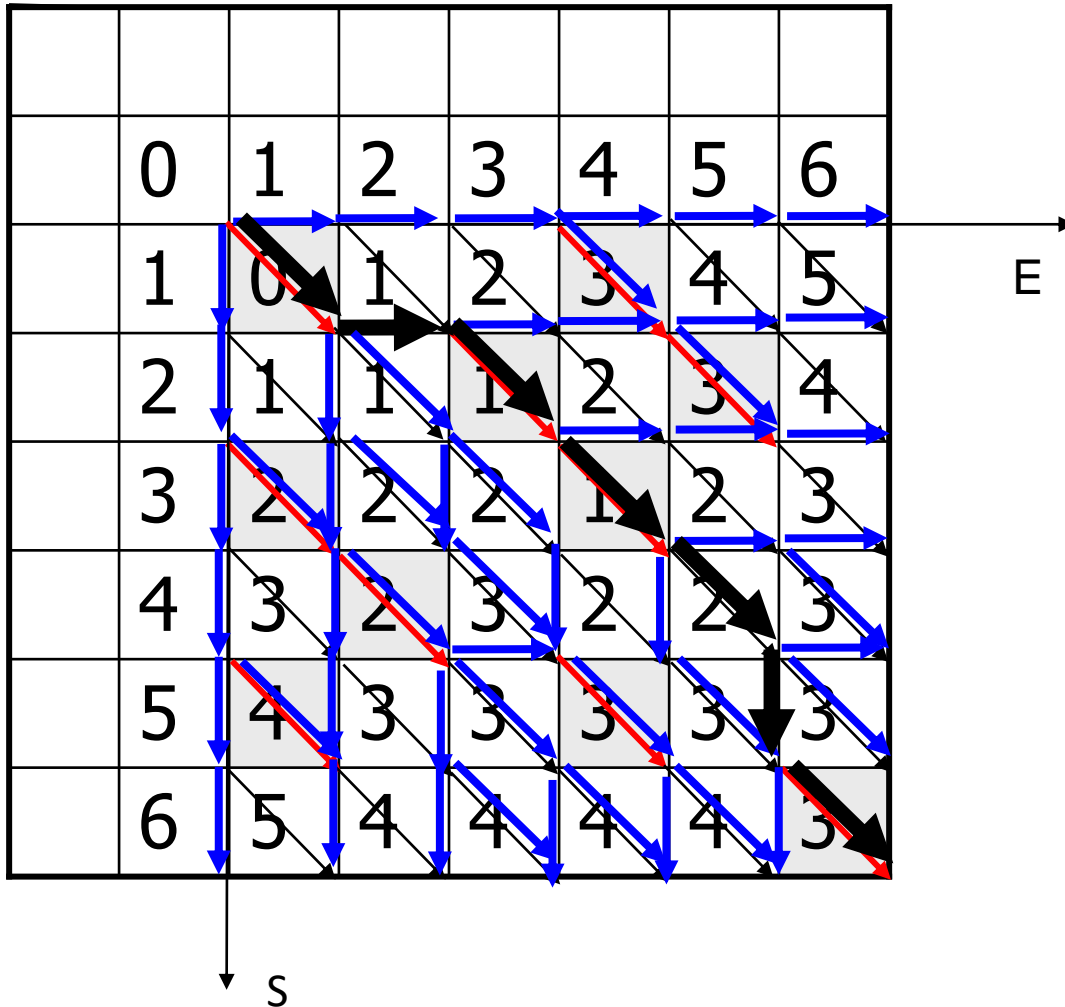




# Keeping track of the source



Trace back –  
how did we get the path with the cost 3?



# Our *Dynamic Programming* algorithm

Algorithm: cheapestPath (diagonalCost NxM)

allocate array *DPTable* ( $N \times M$ )

*DPTable* [0][0]:=0

for *i* from 1 to *N*:

*DPTable* [*i*][0]:=*i*

for *j* from 1 to *M*:

*DPTable* [0][*j*]:=*j*

for *i* from 1 to *N*:

    for *j* from 1 to *M*:

*DPtable* [*i*][*j*]:=min (*DPtable* [*i*-1][*j*-1]+ *diagonalCost* [*i*][*j*],  
                                  *DPtable* [*i*-1][*j*]+1, *DPtable* [*i*][*j*-1]+ 1)

return *DPTable* [*N*][*M*]

2 nested loops:  $O(N^2)$

# Dynamic programming: when

- ❑ We want to **optimize** something: min, max
- ❑ The solution to the problem depends on the solutions to **subproblems**
- ❑ We would need the solutions to **all subproblems**
- ❑ Subproblems **overlap**

# Dynamic programming: how

- ❑ The recurrence relation
- ❑ The bottom-up computation
- ❑ The traceback

# “Programming” in “Dynamic programming” has nothing to do with programming!

- Richard Bellman developed this idea in 1950s working on an Air Force project
- At that time, his approach seemed completely impractical
- He wanted to hide that he is really doing pure math from the Secretary of Defense



**Richard  
Bellman**

... What name could I choose? I was interested in planning but *planning* is not a good word for various reasons. I decided therefore to use the word “programming” and I wanted to get across the idea that this was dynamic. **It was something not even a Congressman could object to.** So I used it as an umbrella for my activities.

**Edit distance**

# Transforming one sequence into another: *edit operations*

- ❑ We can transform the first string S1 into the second S2 by applying a sequence of **edit operations** on S1 :
  - ❑ Deleting 1 symbol
  - ❑ Inserting 1 symbol
  - ❑ Replacing 1 symbol

S1	a	c	t			a	t	g
S2	a	Delete c	t	Insert a	Insert c	a	Delete t	g

In total, 4 edit operations



# String alignment

- An *alignment* of 2 strings is obtained by first inserting spaces (gaps), either into or at the end of both strings, and then placing 2 resulting strings one above the other, so that every character or space in either string is opposite a single character or space in the other string

Alignment

S1	a	c	t	-	-	a	t	g
S2	a	-	t	a	c	a	-	g

4 gaps,  
no mismatches

# Edit distance: definition

- The **edit distance** between two strings is defined as the **minimum number of edit operations** needed to transform one string into another

S1	a	c	t	a	t		g
S2	a	Delete c	t	a	Replace t	Insert a	g

In total, 3 edit operations

# Optimal alignment

- An optimal alignment is obtained from the calculation of the edit distance

## Optimal Alignment

S1	a	c	t	a	t		g
S2	a	Delete c	t	a	Replace t	Insert a	g

Edit distance=3

Is this really the smallest number of edit operations?

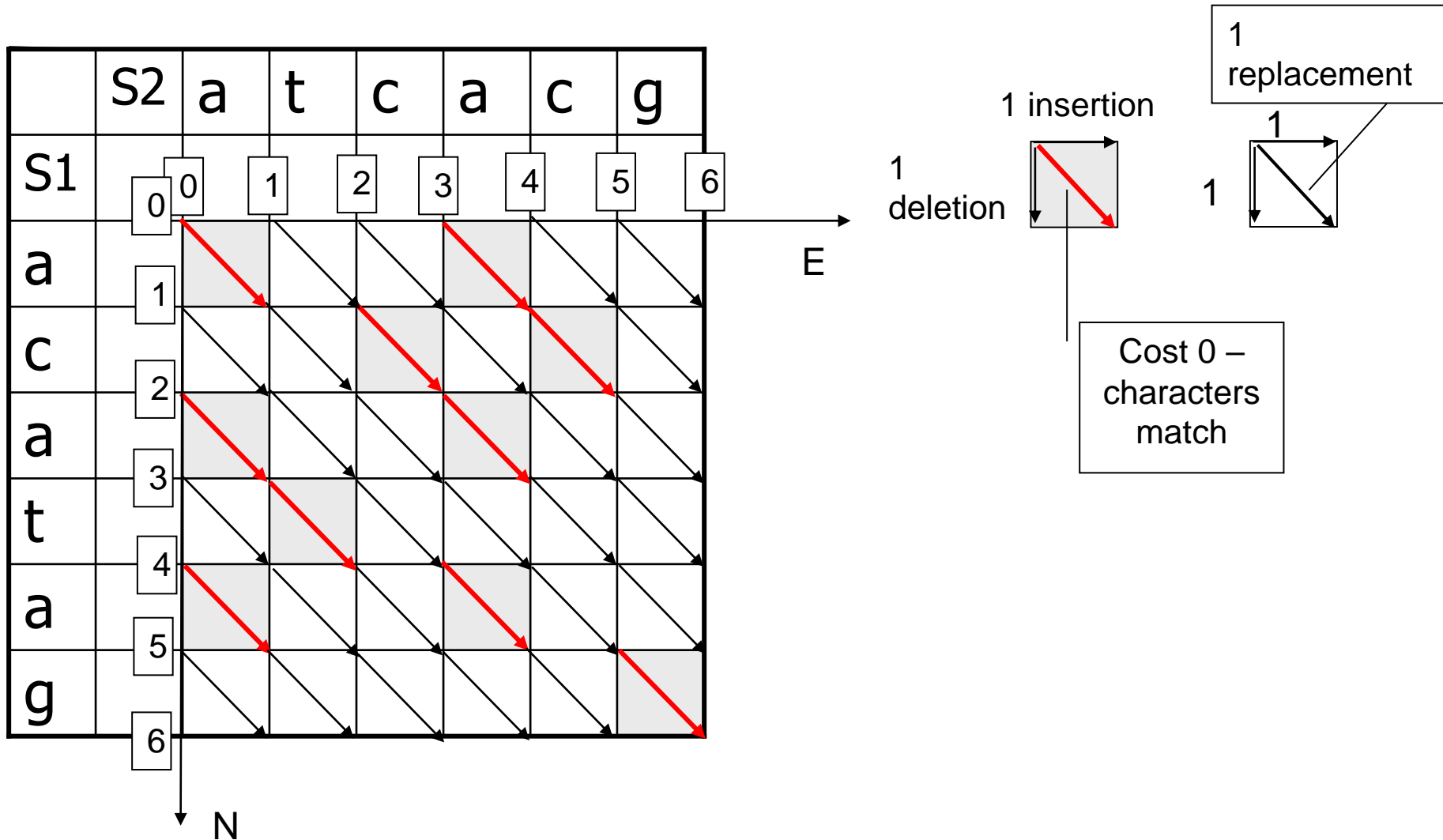
How do we compute edit distance in general?

## The edit distance problem

**Input:** 2 strings  $S_1$  and  $S_2$

**Output:** the *edit distance* between two strings along with a sequence of the operations which describe the transformation

# Full analogy with the cheapest path

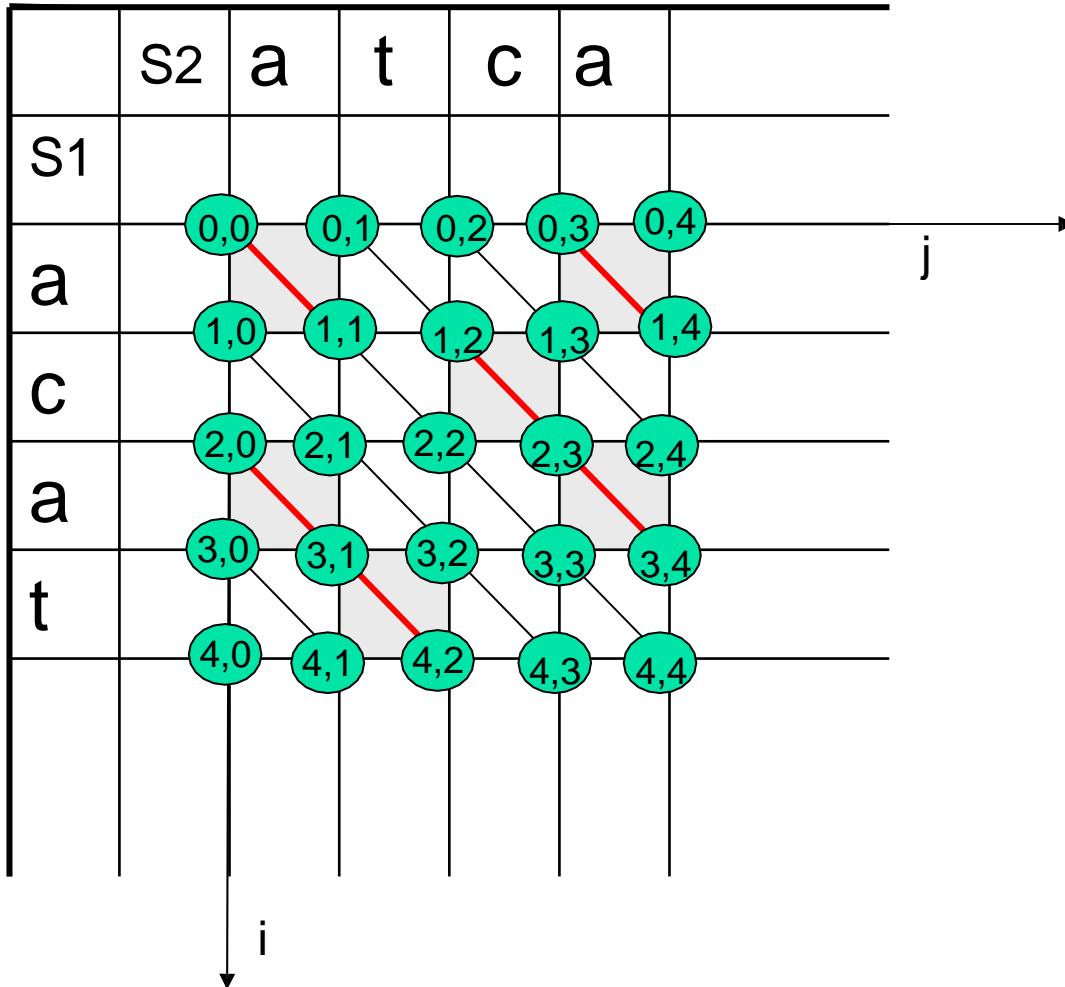


# The dynamic programming solution to the edit distance problem

Trivially follows from the solution for the cheapest path:

- ◆ If we moved strictly down in the grid, we inserted 1 symbol from S1
- ◆ If we moved strictly to the right, we deleted (ignored) 1 symbol of S1
- ◆ If we moved by diagonal of cost 0, we matched the corresponding characters
- ◆ If we moved by diagonal of cost 1, we replaced one symbol in S1 with the corresponding symbol in S2

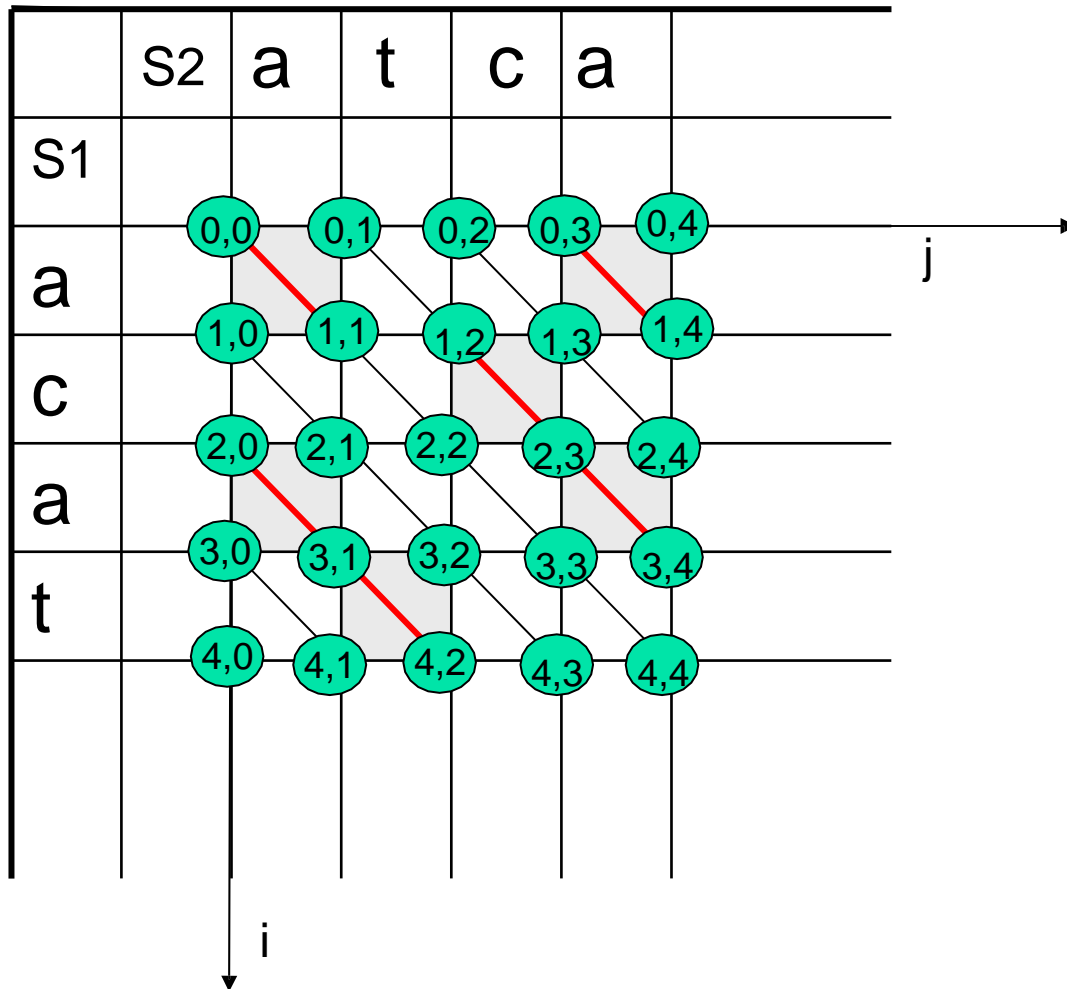
# Useful abstraction: *edit graph*



An **edit graph** for a pair of strings  $S_1$  and  $S_2$  has  $(N+1)*(M+1)$  vertices, each labeled with a corresponding pair  $(i,j)$ ,  $0 \leq i \leq N$ ,  $0 \leq j \leq M$

The edges are **directed** and their weight depends on the specific string problem: for the edit distance problem – red edges have cost 0, black edges have cost 1

# The cheapest path in the edit graph

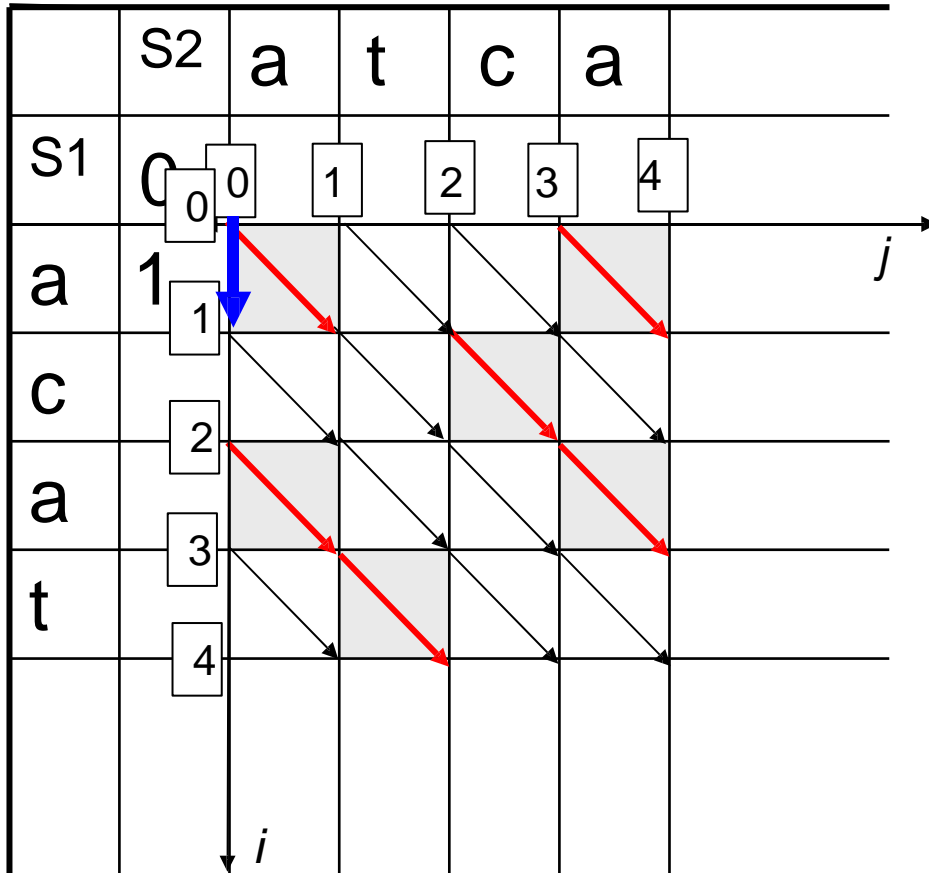


The cost of a **cheapest path** from vertex  $(0,0)$  to vertex  $(N,M)$  in this edit graph corresponds to the **edit distance** between S1 and S2, and the path itself represents a series of edit operations and an optimal alignment of S1 with S2



# Calculating edit distance.

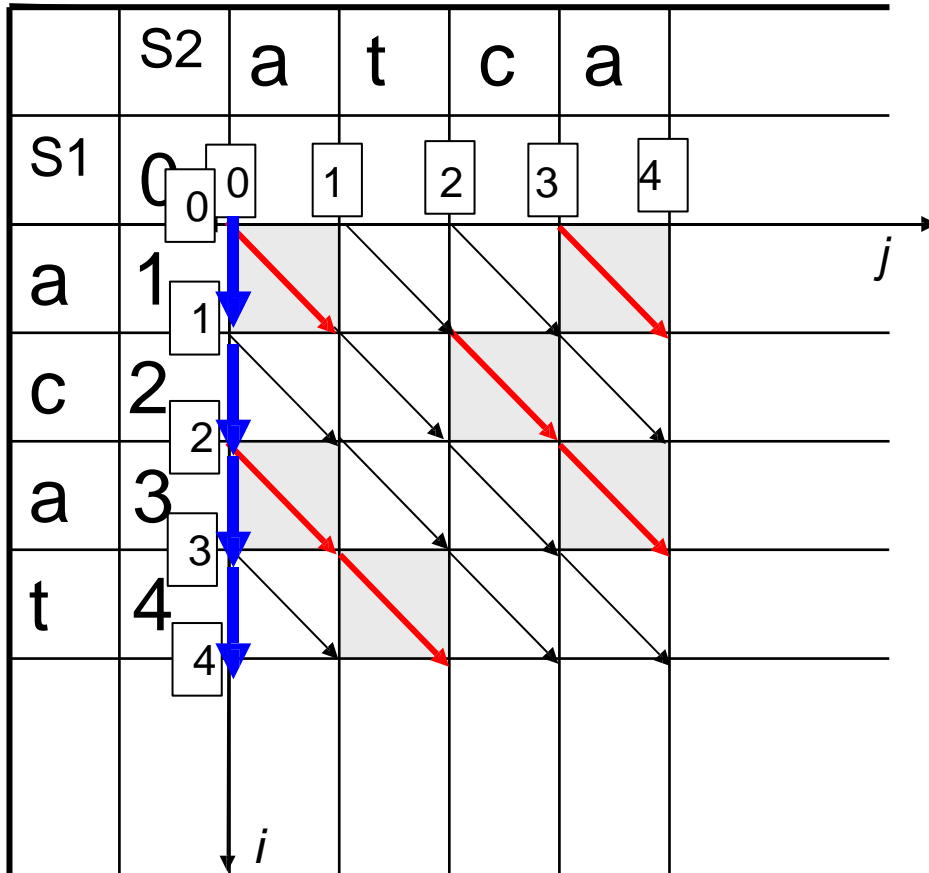
## Base condition



The minimum number of edit operations we need in order to transform string  $a$  into an empty string (of length 0) is 1 (deletion)

Therefore the minimum edit distance between  $\epsilon$  and  $a$  is 1

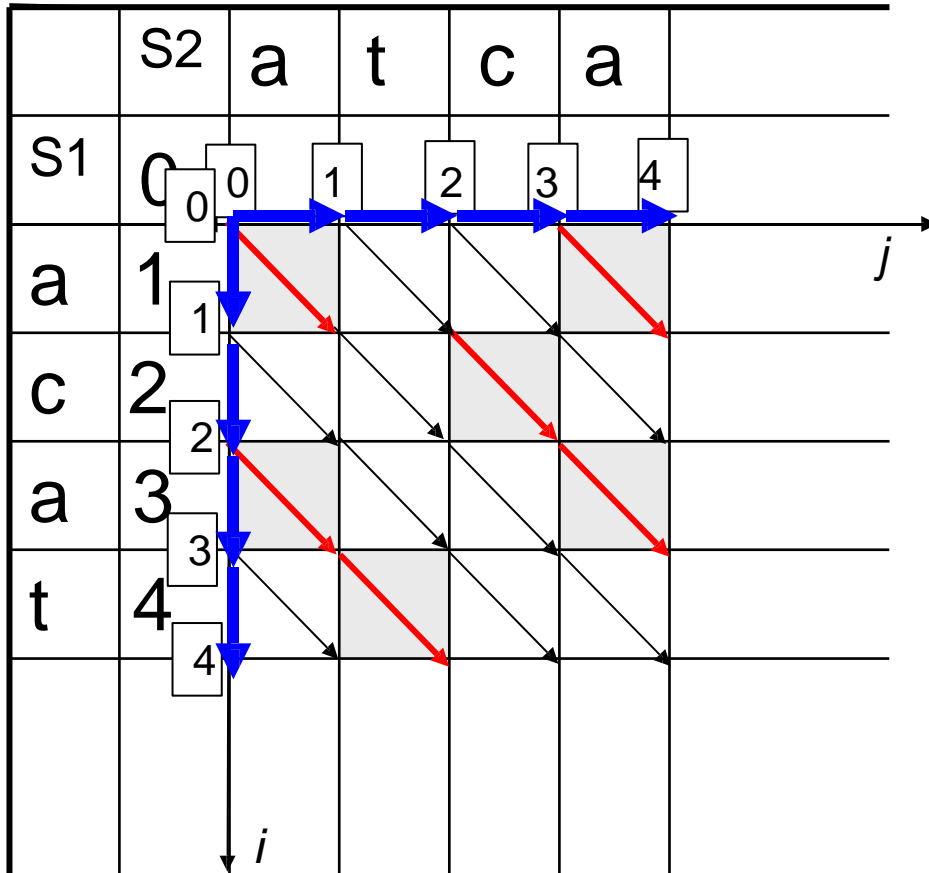
# Calculating edit distance. Base condition



The same is true for  $\epsilon$   
and  $ac, aca, acat$

# Calculating edit distance.

## Base condition



In order to transform  $\epsilon$  into *a*, we need to insert 1 character. This is the best way to do it, there is no cheaper way.

The same for transforming  $\epsilon$  into *at*, *atc*, *atca* with 2, 3, 4 insertions respectively

# Calculating edit distance.

## Filling cells for $i > 0$ and $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	<i>j</i>
<i>a</i>	1					
<i>c</i>	2					
<i>a</i>	3					
<i>t</i>	4					

There are only 3 different ways to move through the next cell in the graph:

1. Increase both  $i$  and  $j$  (diagonal)  
if  $S_1[i] \neq S_2[j]$  : 1 edit  
if  $S_1[i] = S_2[j]$  : 0 edits
1. Increase only  $i$  (insert  $S_1[i]$ ) with the cost 1
2. Increase only  $j$  (delete - ignore  $S_1[i]$ ) with the cost 1

# Calculating edit distance.

## Filling cells for $i > 0$ and $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	
<i>a</i>	1					$j$
<i>c</i>	2					
<i>a</i>	3					
<i>t</i>	4					

Thus, if we know the edit distance  $D[i-1, j-1]$ ,  $D[i-1, j]$  and  $D[i, j-1]$ , we can correctly calculate  $D[i, j]$

This is true since there are no other ways of moving through cell  $[i][j]$ .

Reaching the top, left and top-left corners by different paths cannot produce a better value than is already in these 3 cells, since they contain the minimum cost by definition

# Calculating edit distance.

Filling cells for  $i > 0$  and  $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	
<i>a</i>	1	0	1	2	3	$j$
<i>c</i>	2					
<i>a</i>	3					
<i>t</i>	4					

$i$

$$D(i,j) = \min \begin{cases} D(i-1,j)+1 \\ D(i,j-1)+1 \\ D(i-1,j-1)+c(i,j) \end{cases}$$

$$\text{where } c(i,j) = \begin{cases} 0 & \text{if } S1[j]=S2[j] \\ 1 & \text{if } S1[j] \neq S2[j] \end{cases}$$

# Calculating edit distance.

## Filling cells for $i > 0$ and $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	
<i>a</i>	1	0	1	2	3	$j$
<i>c</i>	2	1	1	1	2	
<i>a</i>	3					
<i>t</i>	4					

$i$

$$D(i,j) = \min \begin{cases} D(i-1,j)+1 \\ D(i,j-1)+1 \\ D(i-1,j-1)+c(i,j) \end{cases}$$

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# Calculating edit distance.

## Filling cells for $i > 0$ and $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	
<i>a</i>	1	0	1	2	3	$j$
<i>c</i>	2	1	1	1	2	
<i>a</i>	3	2	2	2	1	
<i>t</i>	4					

$i$

$$D(i,j) = \min \begin{cases} D(i-1,j)+1 \\ D(i,j-1)+1 \\ D(i-1,j-1)+c(i,j) \end{cases}$$

$$\text{where } c(i,j) = \begin{cases} 0 & \text{if } S1[j]=S2[j] \\ 1 & \text{if } S1[j] \neq S2[j] \end{cases}$$



# Calculating edit distance.

## Filling cells for $i > 0$ and $j > 0$

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	
<i>a</i>	1	0	1	2	3	$j$
<i>c</i>	2	1	1	1	2	
<i>a</i>	3	2	2	2	1	
<i>t</i>	4	3	2	3	2	

$i$

$$D(i,j) = \min \begin{cases} D(i-1,j)+1 \\ D(i,j-1)+1 \\ D(i-1,j-1)+c(i,j) \end{cases}$$

$$\text{where } c(i,j) = \begin{cases} 0 & \text{if } S1[j]=S2[j] \\ 1 & \text{if } S1[j] \neq S2[j] \end{cases}$$

# The sequence of edit operations

	$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	
$S_1$	0	1	2	3	4	$j$
<i>a</i>	1	0	1	2	3	
<i>c</i>	2	1	1	1	2	
<i>a</i>	3	2	2	2	1	
<i>t</i>	4	3	2	3	2	
		$i$				



Place a character in  $S_1$  opposite to a character in  $S_2$



Place a character in  $S_1$  opposite to a gap in  $S_2$



Place a character in  $S_2$  opposite to a gap in  $S_1$

$S_1$	<i>a</i>	-	<i>c</i>	<i>a</i>	<i>t</i>
$S_2$	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	-

# Optimal alignment

<i>S1</i>	<i>a</i>	-	<i>c</i>	<i>a</i>	<i>t</i>
<i>S2</i>	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	-

Explanation:

$S_2$  can be obtained from  $S_1$  by a series of the following edit operations:

Insertion of *t* at position 2

Deletion of *t* at position 5

# An optimal alignment is not unique

<i>S1</i>	-	<i>a</i>	<i>t</i>	<i>t</i>	<i>a</i>	<i>a</i>	<i>g</i>
<i>S2</i>	<i>t</i>	<i>a</i>	-	<i>t</i>	<i>c</i>	<i>a</i>	<i>g</i>

<i>S1</i>	-	<i>a</i>	<i>t</i>	<i>t</i>	<i>a</i>	<i>a</i>	<i>g</i>
<i>S2</i>	<i>t</i>	<i>a</i>	<i>t</i>	<i>c</i>	<i>a</i>	-	<i>g</i>

2 different alignments with the optimal edit distance 3