

## 6. Dynamic Programming

Those who cannot remember the  
past are condemned to repeat it  
-Santayana

## Algorithmic Paradigms

**Greed.** Build up a solution incrementally, myopically optimizing some local criterion.

**Divide-and-conquer.** Break up a problem into two sub-problems, solve each sub-problem independently, and combine solution to sub-problems to form solution to original problem.

**Dynamic programming.** Break up a problem into a series of overlapping sub-problems, and build up solutions to larger and larger sub-problems.

## 6.6 Sequence Alignment

# String Similarity

How similar are two strings?

- **ocurrance**
- **occurrence**

o c u r r a n c e -

o c c u r r e n c e

5 mismatches, 1 gap

o c - u r r a n c e

o c c u r r e n c e

1 mismatch, 1 gap

o c - u r r - a n c e

o c c u r r e - n c e

0 mismatches, 3 gaps

# Edit Distance

## Applications.

- Basis for Unix diff.
- Speech recognition.
- Computational biology.

**Edit distance.** [Levenshtein 1966, Needleman-Wunsch 1970, Smith-Waterman 1981]

- Gap penalty  $\delta$ ; mismatch penalty  $\alpha_{pq}$ .
- Cost = sum of gap and mismatch penalties.

C	T	G	A	C	C	T	A	C	C	T
---	---	---	---	---	---	---	---	---	---	---

C	C	T	G	A	C	T	A	C	A	T
---	---	---	---	---	---	---	---	---	---	---

$$\alpha_{TC} + \alpha_{GT} + \alpha_{AG} + 2\alpha_{CA}$$

-	C	T	G	A	C	C	T	A	C	C	T
---	---	---	---	---	---	---	---	---	---	---	---

C	C	T	G	A	C	-	T	A	C	A	T
---	---	---	---	---	---	---	---	---	---	---	---

$$2\delta + \alpha_{CA}$$

## Sequence Alignment

**Goal:** Given two strings  $X = x_1 x_2 \dots x_m$  and  $Y = y_1 y_2 \dots y_n$  find alignment of minimum cost.

**Def.** An **alignment**  $M$  is a set of ordered pairs  $x_i - y_j$  such that each item occurs in at most one pair and no crossings.

**Def.** The pair  $x_i - y_j$  and  $x_{i'} - y_{j'}$  **cross** if  $i < i'$ , but  $j > j'$ .

$$\text{cost}(M) = \underbrace{\sum_{(x_i, y_j) \in M} \alpha_{x_i y_j}}_{\text{mismatch}} + \underbrace{\sum_{i: x_i \text{ unmatched}} \delta + \sum_{j: y_j \text{ unmatched}} \delta}_{\text{gap}}$$

**Ex:** CTACCG vs. TACATG.

**Sol:**  $M = x_2 - y_1, x_3 - y_2, x_4 - y_3, x_5 - y_4, x_6 - y_6$ .

$x_1$	$x_2$	$x_3$	$x_4$	$x_5$		$x_6$
C	T	A	C	C	-	G
	-	T	A	A	T	G
	$y_1$	$y_2$	$y_3$	$y_4$	$y_5$	$y_6$

## Sequence Alignment: Problem Structure

**Def.**  $OPT(i, j)$  = min cost of aligning strings  $x_1 x_2 \dots x_i$  and  $y_1 y_2 \dots y_j$ .

- Case 1:  $OPT$  matches  $x_i$ - $y_j$ .
  - pay mismatch for  $x_i$ - $y_j$  + min cost of aligning two strings  $x_1 x_2 \dots x_{i-1}$  and  $y_1 y_2 \dots y_{j-1}$
- Case 2a:  $OPT$  leaves  $x_i$  unmatched.
  - pay gap for  $x_i$  and min cost of aligning  $x_1 x_2 \dots x_{i-1}$  and  $y_1 y_2 \dots y_j$
- Case 2b:  $OPT$  leaves  $y_j$  unmatched.
  - pay gap for  $y_j$  and min cost of aligning  $x_1 x_2 \dots x_i$  and  $y_1 y_2 \dots y_{j-1}$

$$OPT(i, j) = \begin{cases} j\delta & \text{if } i = 0 \\ \min \begin{cases} \alpha_{x_i y_j} + OPT(i-1, j-1) \\ \delta + OPT(i-1, j) \\ \delta + OPT(i, j-1) \end{cases} & \text{otherwise} \\ i\delta & \text{if } j = 0 \end{cases}$$

## Sequence Alignment: Algorithm

```
Sequence-Alignment(m, n,  $x_1x_2\dots x_m$ ,  $y_1y_2\dots y_n$ ,  $\delta$ ,  $\alpha$ ) {  
  for i = 0 to m  
    M[0, i] =  $i\delta$   
  for j = 0 to n  
    M[j, 0] =  $j\delta$   
  
  for i = 1 to m  
    for j = 1 to n  
      M[i, j] = min( $\alpha[x_i, y_j] + M[i-1, j-1]$ ,  
                    $\delta + M[i-1, j]$ ,  
                    $\delta + M[i, j-1]$ )  
  
  return M[m, n]  
}
```

Analysis.  $\Theta(mn)$  time and space.

English words or sentences:  $m, n \leq 10$ .

Computational biology:  $m = n = 100,000$ . 10 billions ops OK, but 10GB array?



## 6.7 Sequence Alignment in Linear Space

## Sequence Alignment: Linear Space

Q. Can we avoid using quadratic **space**?

Easy. Optimal **value** in  $O(m + n)$  space and  $O(mn)$  time.

- Compute  $\text{OPT}(i, \cdot)$  from  $\text{OPT}(i-1, \cdot)$ .
- No longer a simple way to recover alignment itself.

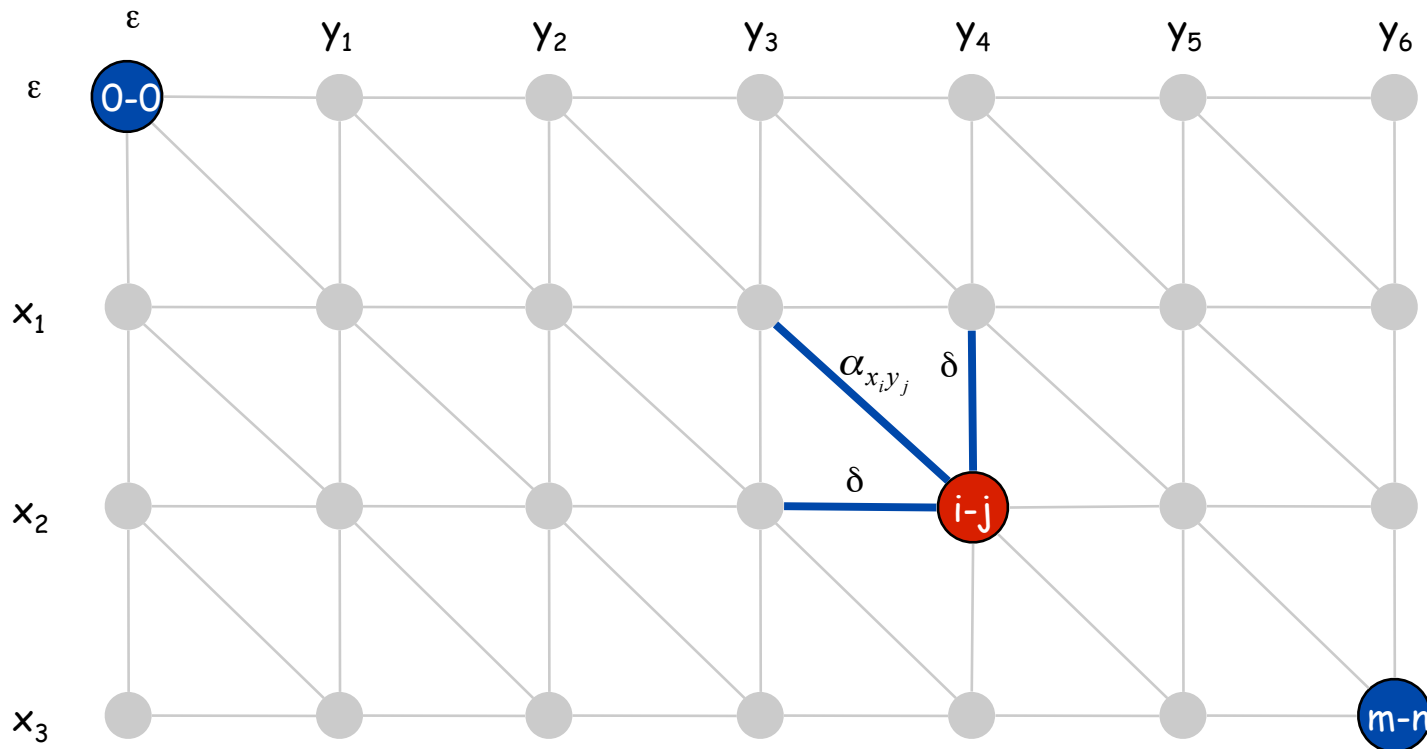
Theorem. [Hirschberg, 1975] Optimal **alignment** in  $O(m + n)$  space and  $O(mn)$  time.

- Clever combination of divide-and-conquer and dynamic programming.
- Inspired by idea of Savitch from complexity theory.

## Sequence Alignment: Linear Space

### Edit distance graph.

- Let  $f(i, j)$  be shortest path from  $(0,0)$  to  $(i, j)$ .
- Observation:  $f(i, j) = \text{OPT}(i, j)$ .



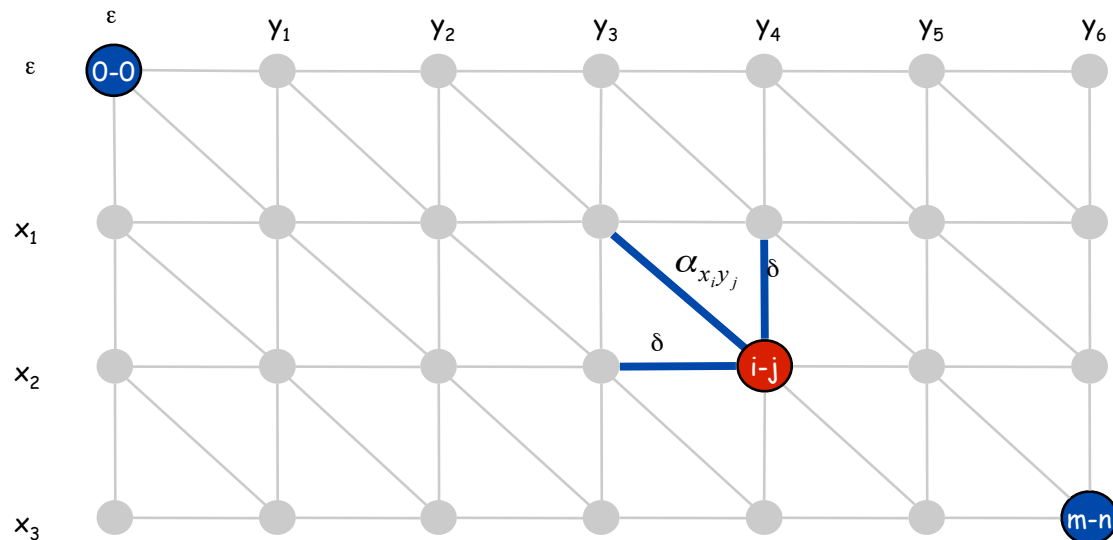
## Sequence Alignment: Linear Space

**Claim.**  $f(i, j) = \text{OPT}(i, j)$ .

**Pf.** (by induction on  $i + j$ )

- Base case:  $f(0, 0) = \text{OPT}(0, 0) = 0$ .
- Inductive step: assume  $f(i', j') = \text{OPT}(i', j')$  for all  $i' + j' < i + j$ .
- Last edge on path to  $(i, j)$  is either from  $(i-1, j-1)$ ,  $(i-1, j)$ , or  $(i, j-1)$ .

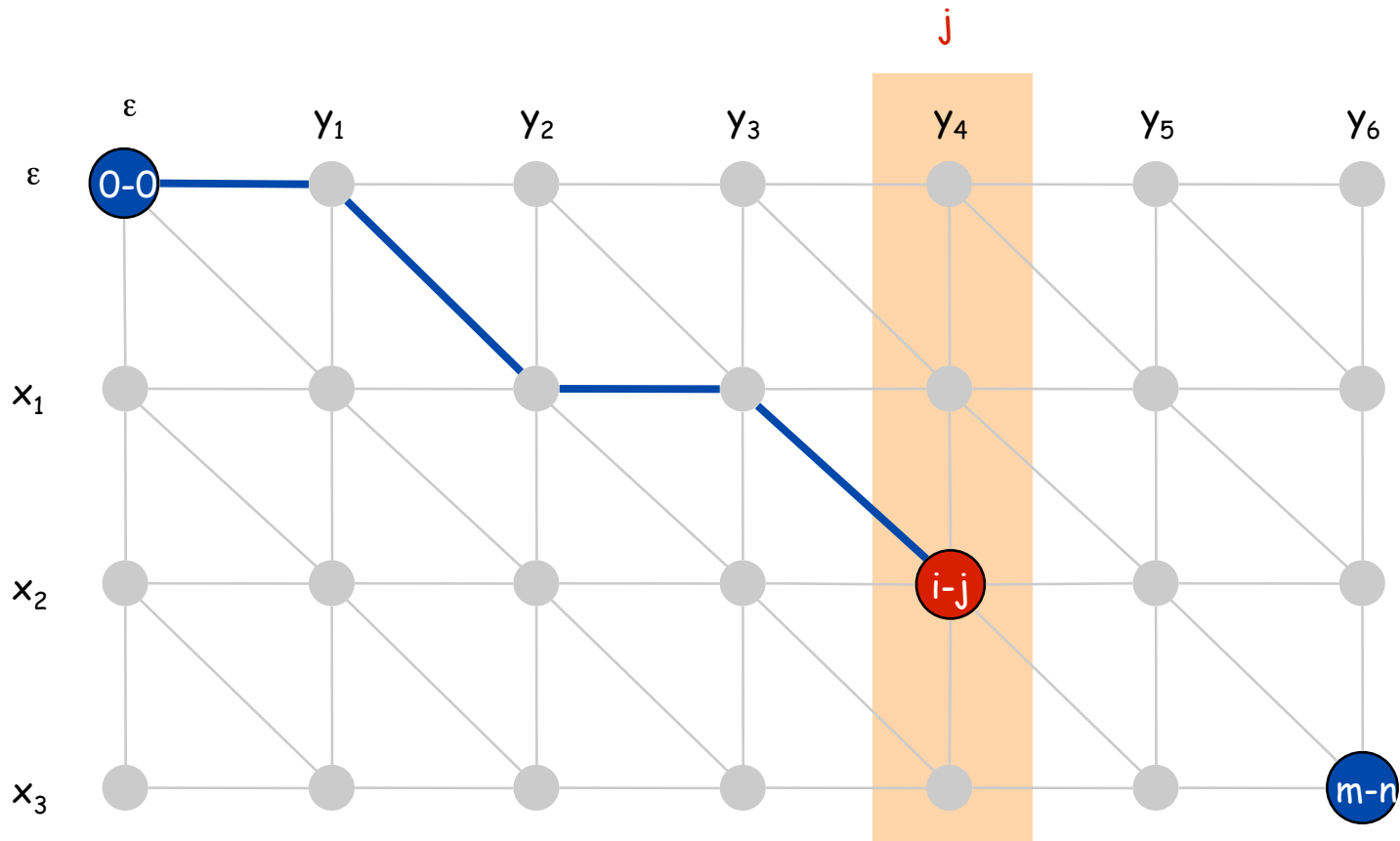
$$\begin{aligned}
 f(i, j) &= \min \{ \alpha_{x_i y_j} + f(i-1, j-1), \delta + f(i-1, j), \delta + f(i, j-1) \} \\
 &= \min \{ \alpha_{x_i y_j} + \text{OPT}(i-1, j-1), \delta + \text{OPT}(i-1, j), \delta + \text{OPT}(i, j-1) \} \\
 &= \text{OPT}(i, j)
 \end{aligned}$$



## Sequence Alignment: Linear Space

### Edit distance graph.

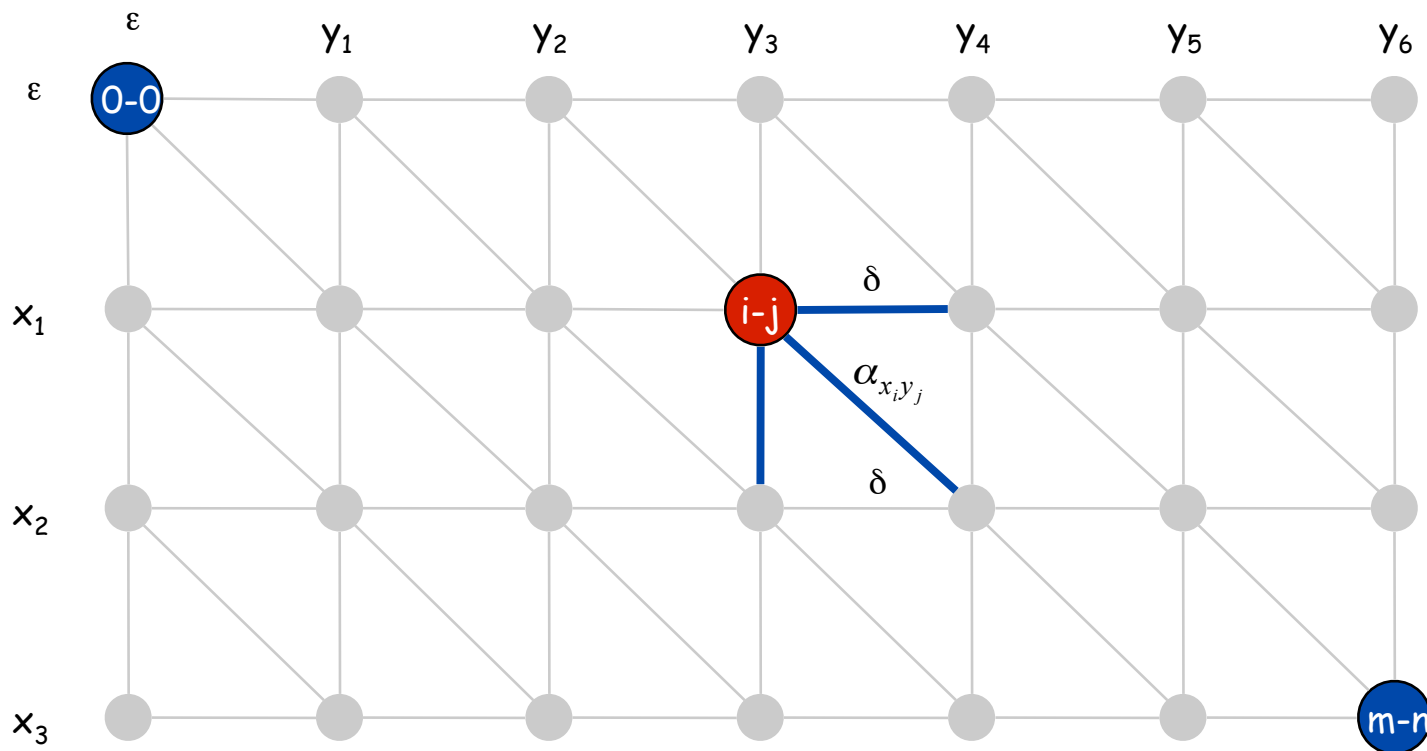
- Let  $f(i, j)$  be shortest path from  $(0,0)$  to  $(i, j)$ .
- Can compute  $f(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.



## Sequence Alignment: Linear Space

### Edit distance graph.

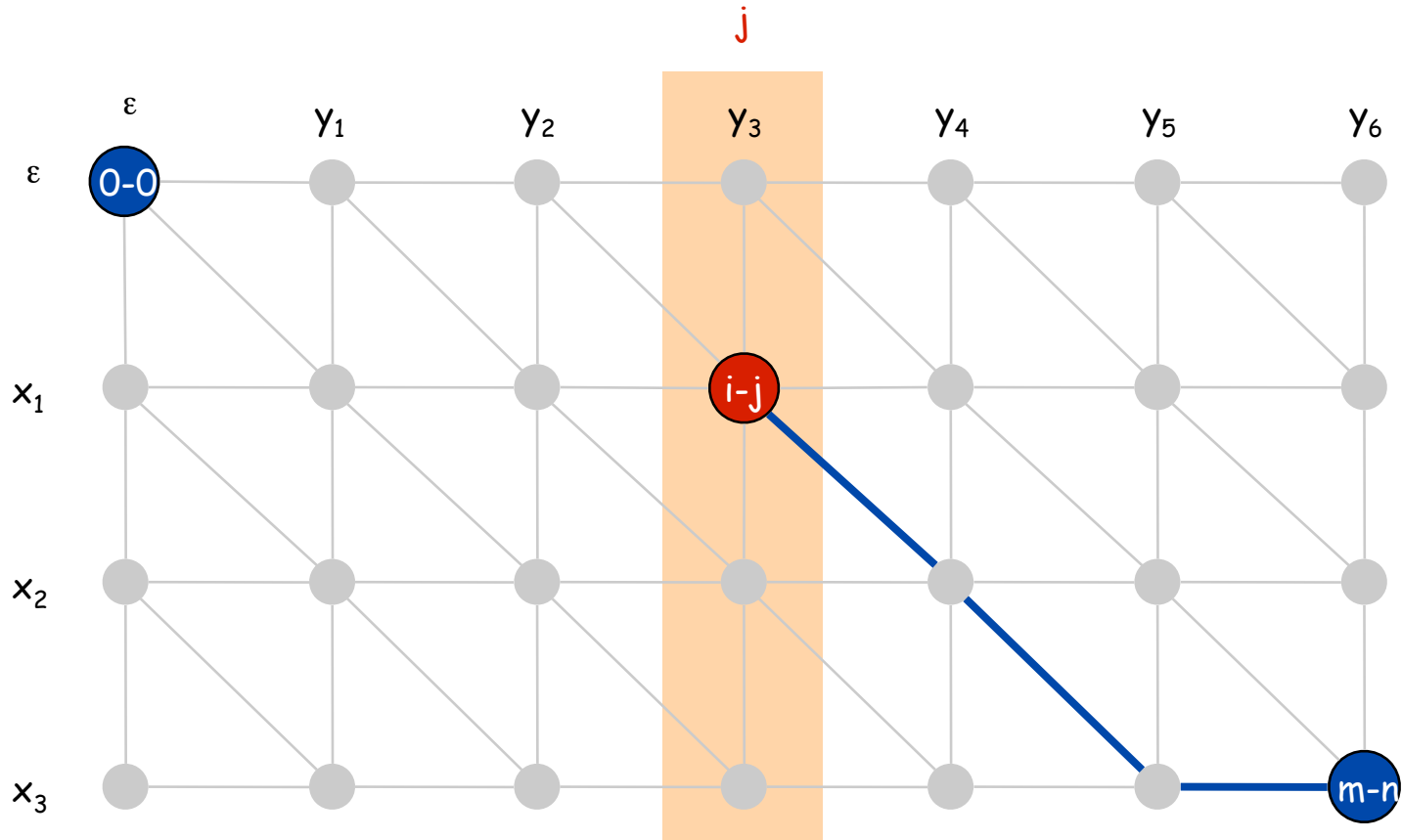
- Let  $g(i, j)$  be shortest path from  $(i, j)$  to  $(m, n)$ .
- Can compute by reversing the edge orientations and inverting the roles of  $(0, 0)$  and  $(m, n)$



## Sequence Alignment: Linear Space

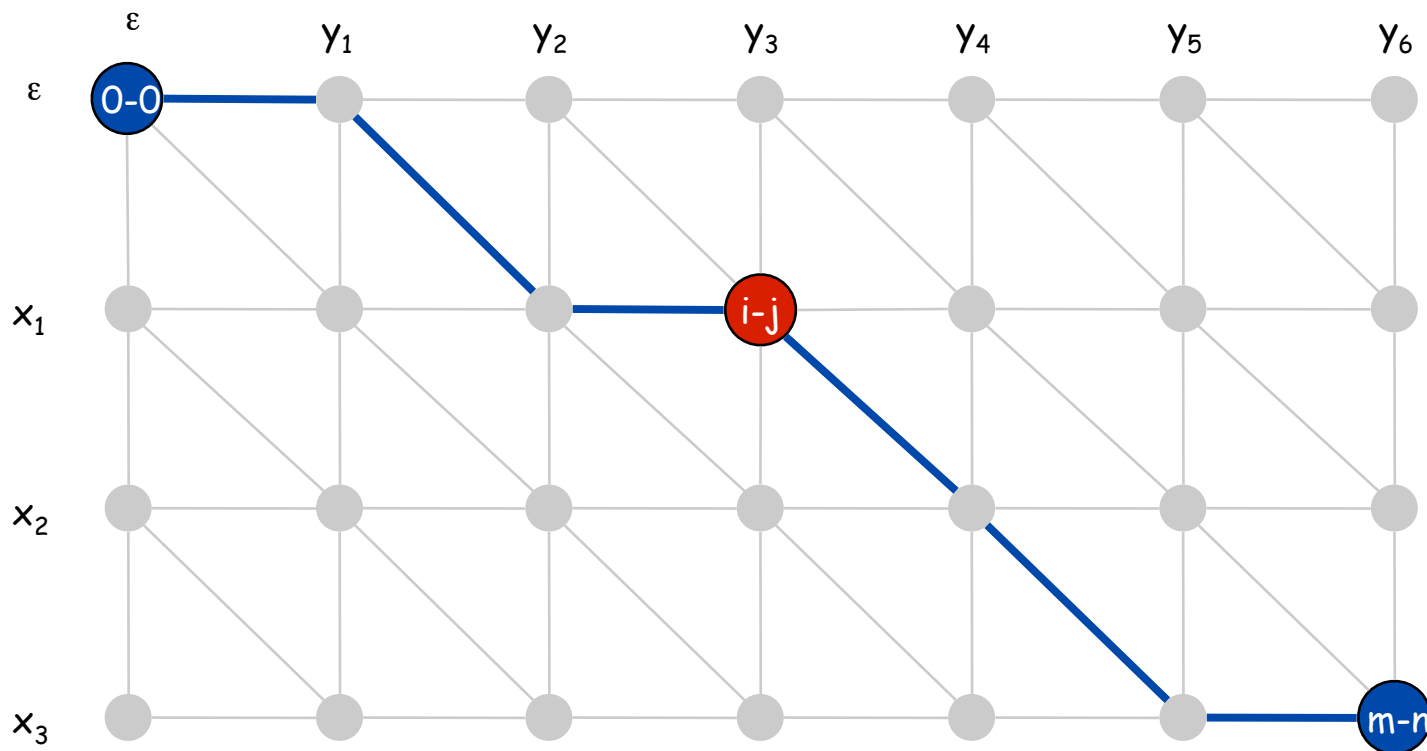
### Edit distance graph.

- Let  $g(i, j)$  be shortest path from  $(i, j)$  to  $(m, n)$ .
- Can compute  $g(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.



## Sequence Alignment: Linear Space

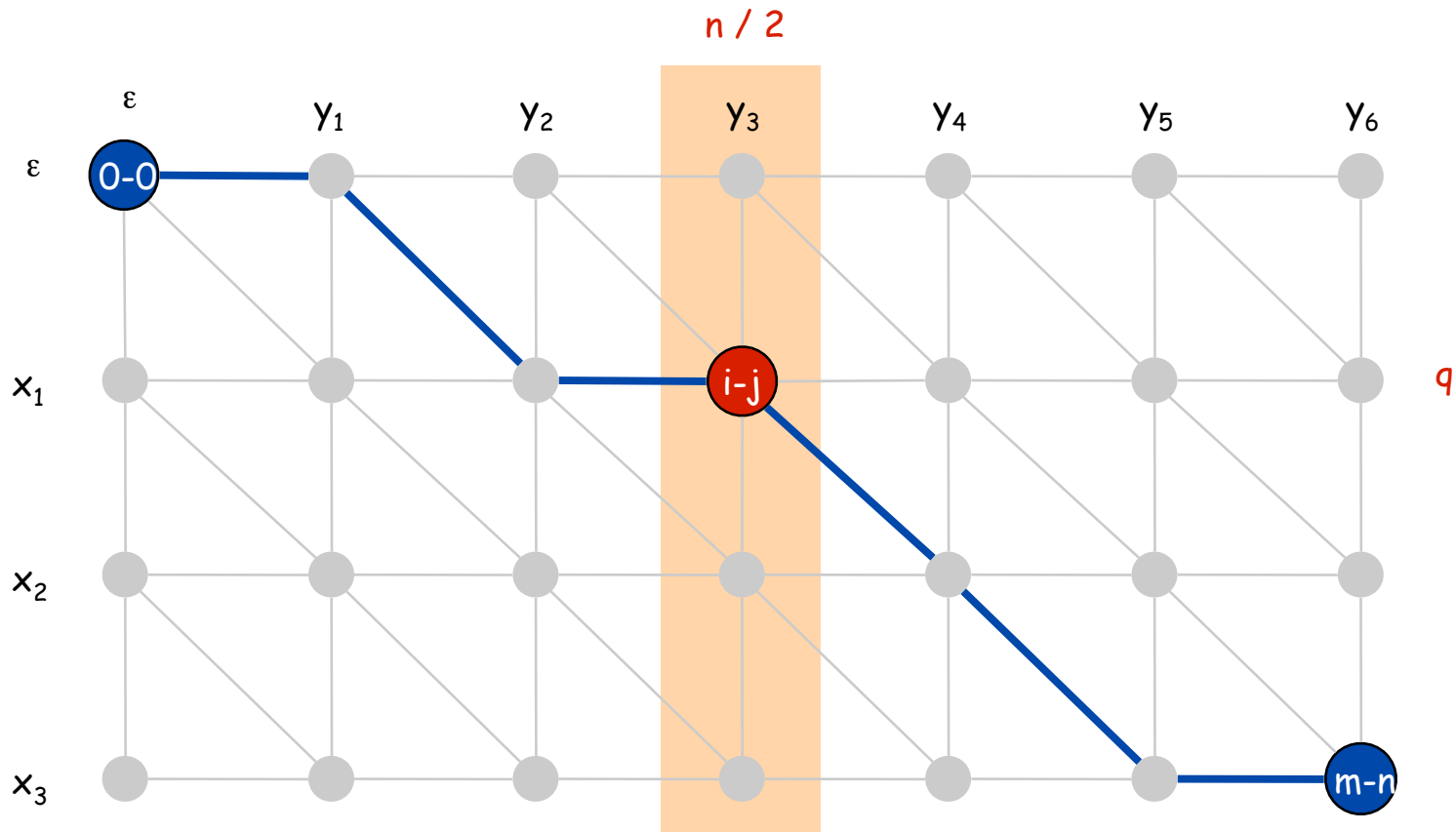
**Observation 1.** The cost of the shortest path that uses  $(i, j)$  is  $f(i, j) + g(i, j)$ .





## Sequence Alignment: Linear Space

**Observation 2.** let  $q$  be an index that minimizes  $f(q, n/2) + g(q, n/2)$ . Then, the shortest path from  $(0, 0)$  to  $(m, n)$  uses  $(q, n/2)$ .

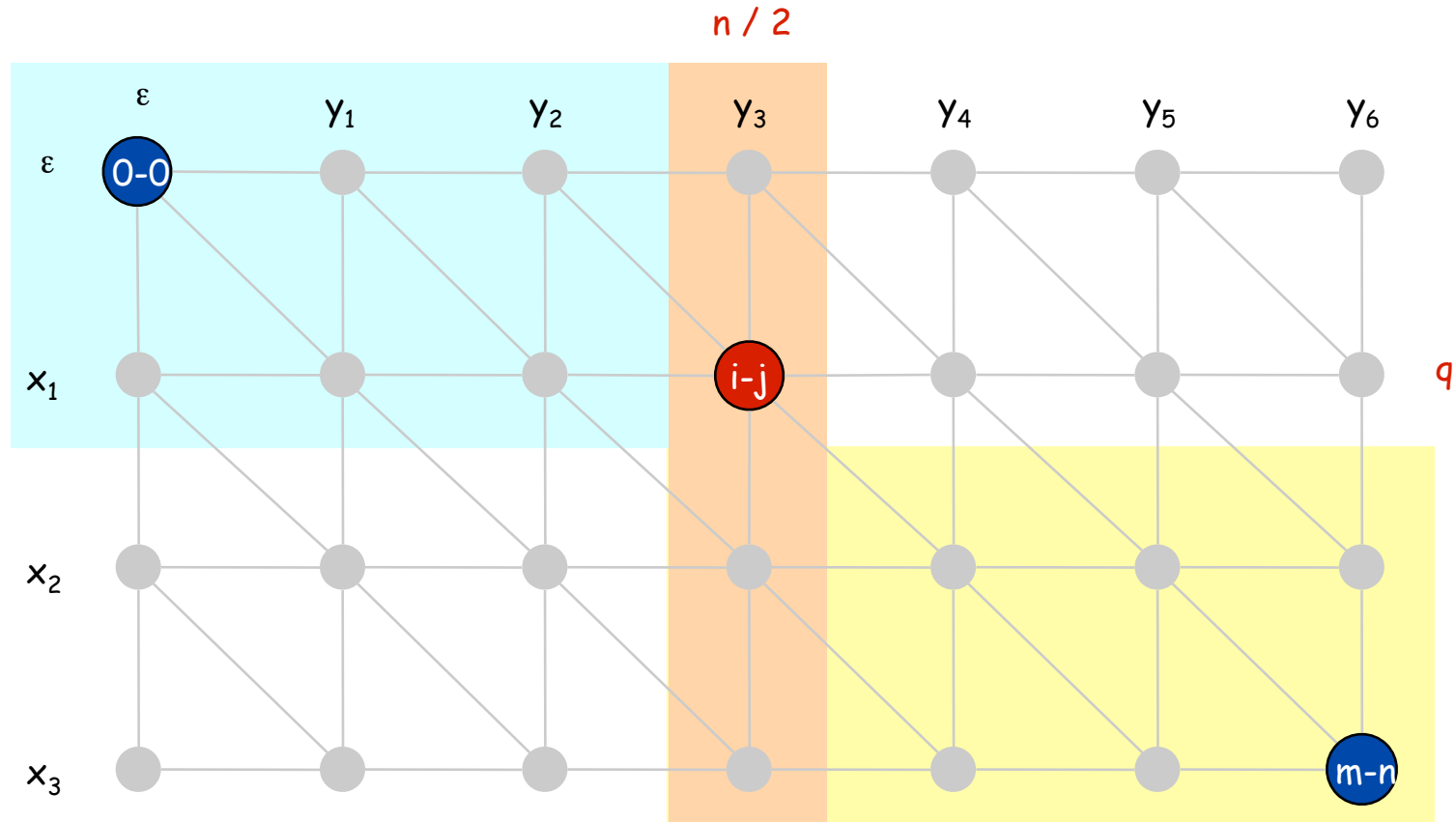


## Sequence Alignment: Linear Space

**Divide:** find index  $q$  that minimizes  $f(q, n/2) + g(q, n/2)$  using DP.

- Align  $x_q$  and  $y_{n/2}$ .

**Conquer:** recursively compute optimal alignment in each piece.



## Sequence Alignment: Running Time Analysis Warmup

**Theorem.** Let  $T(m, n)$  = max running time of algorithm on strings of length at most  $m$  and  $n$ .  $T(m, n) = O(mn \log n)$ .

$$T(m, n) \leq 2T(m, n/2) + O(mn) \Rightarrow T(m, n) = O(mn \log n)$$

**Remark.** Analysis is not tight because two sub-problems are of size  $(q, n/2)$  and  $(m - q, n/2)$ . In next slide, we save  $\log n$  factor.

## Sequence Alignment: Running Time Analysis

**Theorem.** Let  $T(m, n)$  = max running time of algorithm on strings of length  $m$  and  $n$ .  $T(m, n) = O(mn)$ .

**Pf.** (by induction on  $n$ )

- $O(mn)$  time to compute  $f(\cdot, n/2)$  and  $g(\cdot, n/2)$  and find index  $q$ .
- $T(q, n/2) + T(m - q, n/2)$  time for two recursive calls.
- Choose constant  $c$  so that:

$$T(m, 2) \leq cm$$

$$T(2, n) \leq cn$$

$$T(m, n) \leq cmn + T(q, n/2) + T(m - q, n/2)$$

- Base cases:  $m = 2$  or  $n = 2$ .
- Inductive hypothesis: for  $m' < m$  or  $n' < n$ ,  $T(m', n') \leq 2cm'n'$ .

$$\begin{aligned} T(m, n) &\leq T(q, n/2) + T(m - q, n/2) + cmn \\ &\leq 2cq n/2 + 2c(m - q)n/2 + cmn \\ &= cq n + cmn - cq n + cmn \\ &= 2cmn \end{aligned}$$





