

# Chapter 6

## Dynamic Programming



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## Algorithmic Paradigms

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

**Divide-and-conquer.** Break up a problem into 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.

# Dynamic Programming History

**Bellman.** [1950s] Pioneered the systematic study of dynamic programming.

## Etymology.

- Dynamic programming = planning over time.
- Secretary of Defense was hostile to mathematical research.
- Bellman sought an impressive name to avoid confrontation.

"it's impossible to use dynamic in a pejorative sense"  
"something not even a Congressman could object to"

Reference: Bellman, R. E. *Eye of the Hurricane, An Autobiography*.

# Dynamic Programming Applications

## Areas.

- Bioinformatics.
- Control theory.
- Information theory.
- Operations research.
- Computer science: theory, graphics, AI, compilers, systems, ....

## Some famous dynamic programming algorithms.

- Unix diff for comparing two files.
- Viterbi for hidden Markov models.
- Smith-Waterman for genetic sequence alignment.
- Bellman-Ford for shortest path routing in networks.
- Cocke-Kasami-Younger for parsing context free grammars.

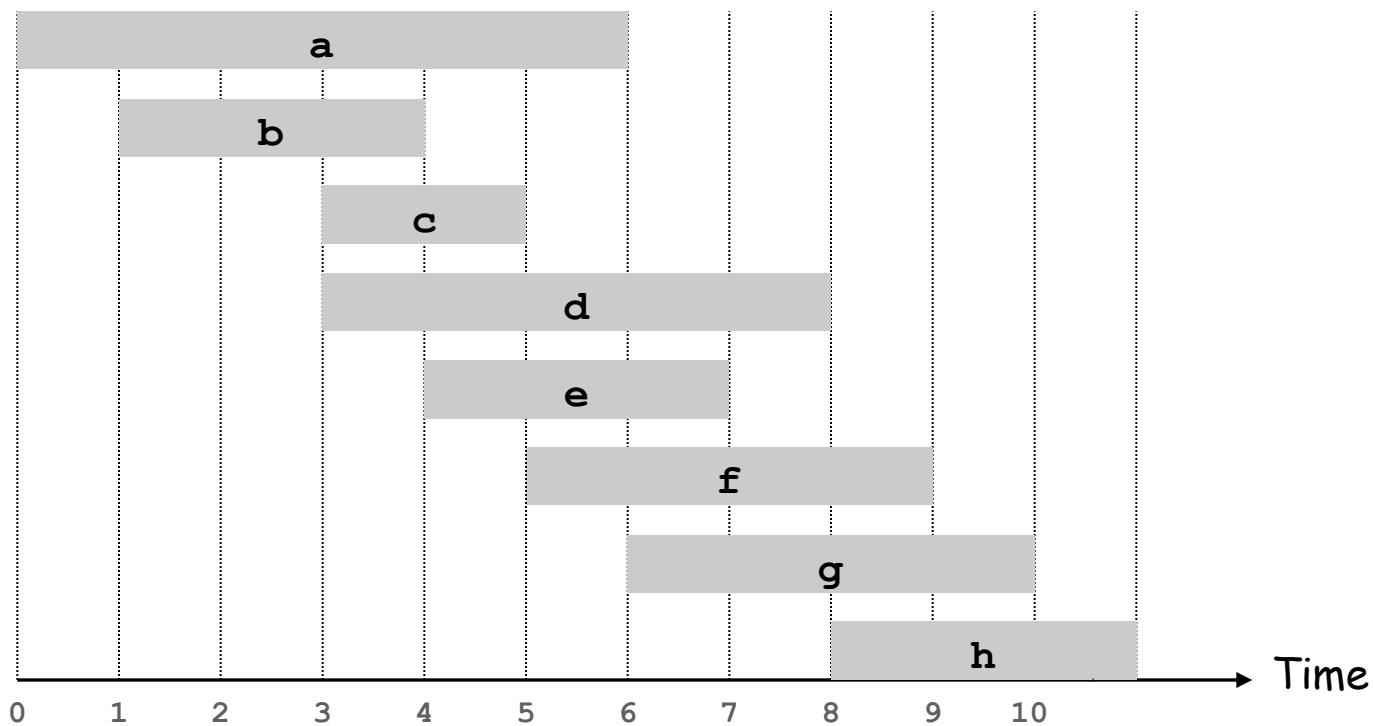
## 6.1 Weighted Interval Scheduling

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# Weighted Interval Scheduling

## Weighted interval scheduling problem.

- Job  $j$  starts at  $s_j$ , finishes at  $f_j$ , and has weight or value  $v_j$ .
- Two jobs **compatible** if they don't overlap.
- Goal: find maximum **weight** subset of mutually compatible jobs.

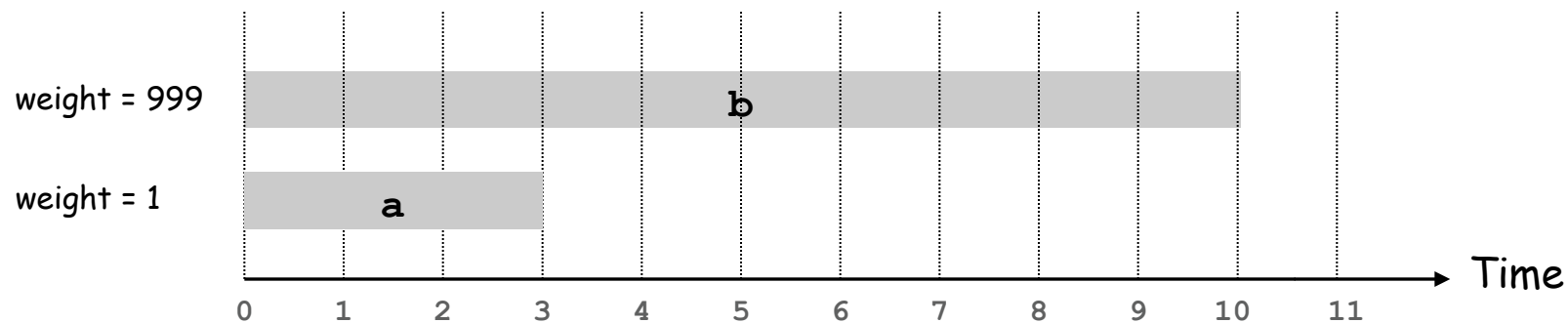


## Unweighted Interval Scheduling Review

**Recall.** Greedy algorithm works if all weights are 1.

- Consider jobs in ascending order of finish time.
- Add job to subset if it is compatible with previously chosen jobs.

**Observation.** Greedy algorithm can fail spectacularly if arbitrary weights are allowed.

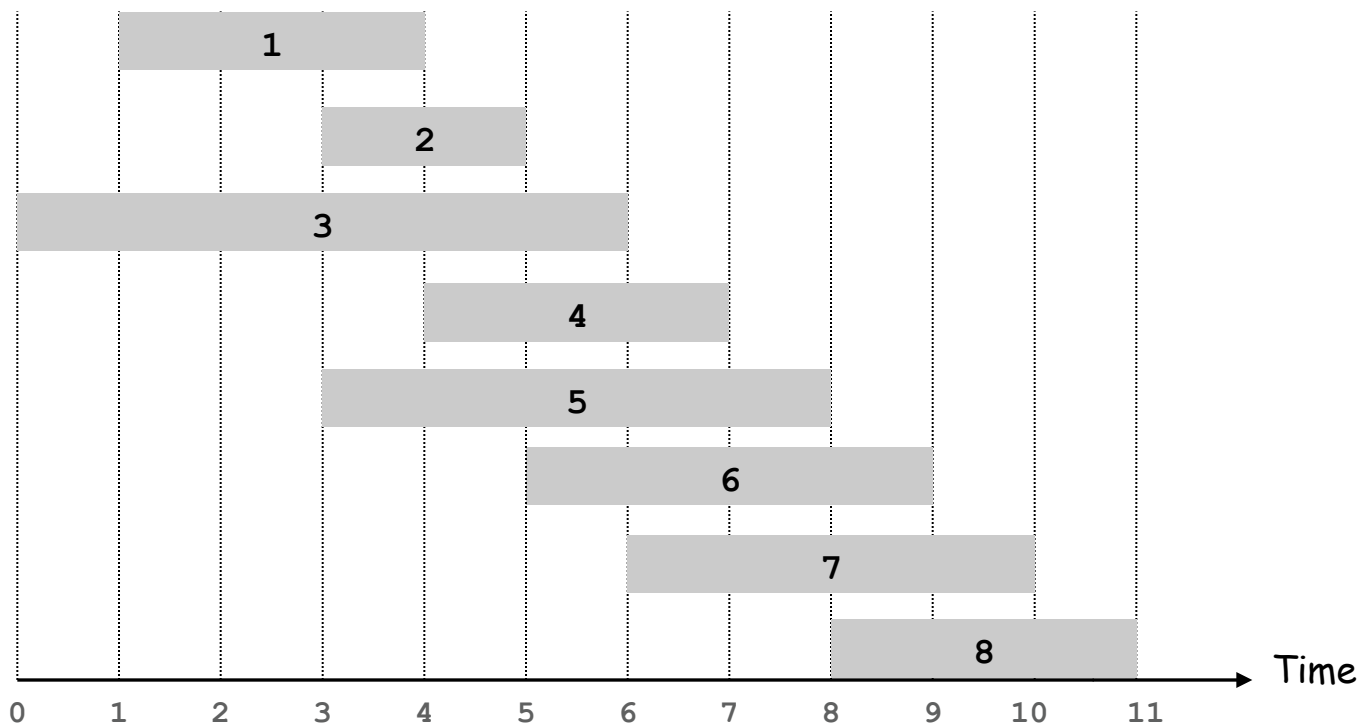


# Weighted Interval Scheduling

**Notation.** Label jobs by finishing time:  $f_1 \leq f_2 \leq \dots \leq f_n$ .

**Def.**  $p(j)$  = largest index  $i < j$  such that job  $i$  is compatible with  $j$ .

**Ex:**  $p(8) = 5$ ,  $p(7) = 3$ ,  $p(2) = 0$ .





## Dynamic Programming: Binary Choice

**Notation.**  $OPT(j)$  = value of optimal solution to the problem consisting of job requests  $1, 2, \dots, j$ .

- Case 1:  $OPT$  selects job  $j$ .
  - collect profit  $v_j$
  - can't use incompatible jobs  $\{ p(j) + 1, p(j) + 2, \dots, j - 1 \}$
  - must include optimal solution to problem consisting of remaining compatible jobs  $1, 2, \dots, p(j)$
- Case 2:  $OPT$  does not select job  $j$ .
  - must include optimal solution to problem consisting of remaining compatible jobs  $1, 2, \dots, j-1$

optimal substructure

$$OPT(j) = \begin{cases} 0 & \text{if } j = 0 \\ \max \{ v_j + OPT(p(j)), OPT(j-1) \} & \text{otherwise} \end{cases}$$

## Weighted Interval Scheduling: Brute Force

Brute force algorithm.

```
Input:  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$ 
```

```
Sort jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .
```

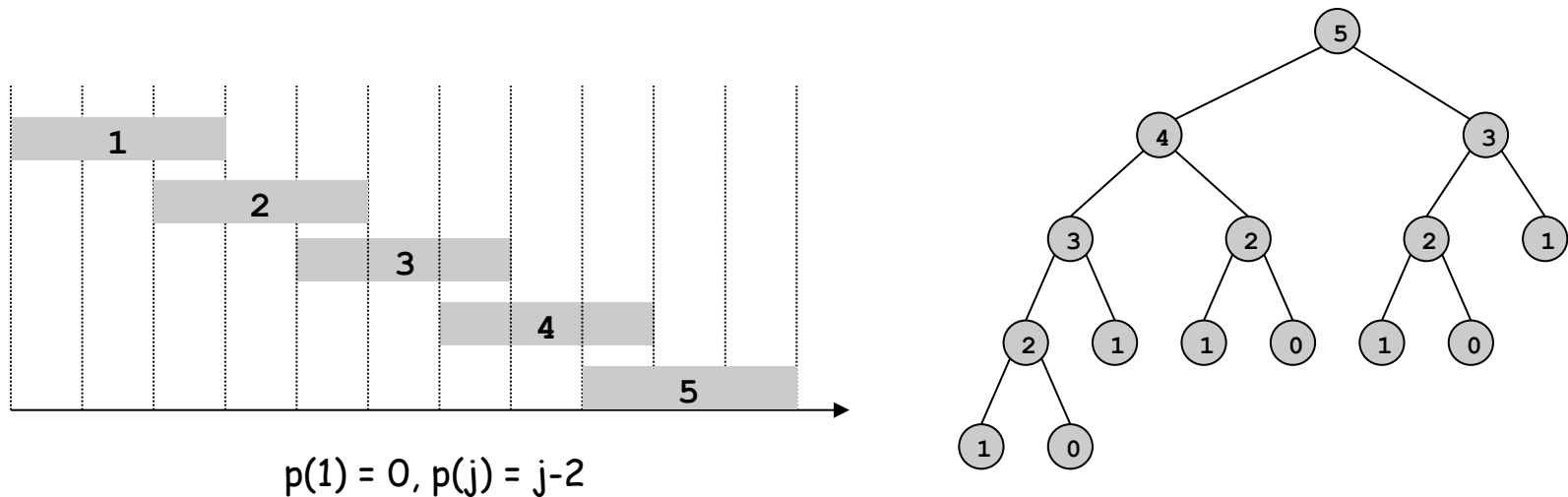
```
Compute  $p(1), p(2), \dots, p(n)$ 
```

```
Compute-Opt( $j$ ) {  
    if ( $j = 0$ )  
        return 0  
    else  
        return  $\max(v_j + \text{Compute-Opt}(p(j)), \text{Compute-Opt}(j-1))$   
}
```

# Weighted Interval Scheduling: Brute Force

**Observation.** Recursive algorithm fails spectacularly because of redundant sub-problems  $\Rightarrow$  exponential algorithms.

**Ex.** Number of recursive calls for family of "layered" instances grows like Fibonacci sequence.



## Weighted Interval Scheduling: Memoization

**Memoization.** Store results of each sub-problem in a cache; lookup as needed.

```
Input:  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$ 
```

```
Sort jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .
```

```
Compute  $p(1), p(2), \dots, p(n)$ 
```

```
for  $j = 1$  to  $n$ 
```

```
     $M[j] = \text{empty}$ 
```

```
 $M[0] = 0$ 
```

← global array

```
M-Compute-Opt( $j$ ) {
```

```
    if ( $M[j]$  is empty)
```

```
         $M[j] = \max(v_j + \text{M-Compute-Opt}(p(j)), \text{M-Compute-Opt}(j-1))$ 
```

```
    return  $M[j]$ 
```

```
}
```

## Weighted Interval Scheduling: Running Time

**Claim.** Memoized version of algorithm takes  $O(n \log n)$  time.

- Sort by finish time:  $O(n \log n)$ .
- Computing  $p(\cdot)$ :  $O(n \log n)$  via sorting by start time.
- $M\text{-Compute-Opt}(j)$ : each invocation takes  $O(1)$  time and either
  - (i) returns an existing value  $M[j]$
  - (ii) fills in one new entry  $M[j]$  and makes two recursive calls
- Progress measure  $\Phi = \#$  nonempty entries of  $M[\ ]$ .
  - initially  $\Phi = 0$ , throughout  $\Phi \leq n$ .
  - (ii) increases  $\Phi$  by 1  $\Rightarrow$  at most  $2n$  recursive calls.
- Overall running time of  $M\text{-Compute-Opt}(n)$  is  $O(n)$ . ▪

**Remark.**  $O(n)$  if jobs are pre-sorted by start and finish times.

## Weighted Interval Scheduling: Finding a Solution

Q. Dynamic programming algorithms computes optimal value.

What if we want the solution itself?

A. Do some post-processing.

```
Run M-Compute-Opt(n)
Run Find-Solution(n)

Find-Solution(j) {
    if (j = 0)
        output nothing
    else if ( $v_j + M[p(j)] > M[j-1]$ )
        print j
        Find-Solution(p(j))
    else
        Find-Solution(j-1)
}
```

- # of recursive calls  $\leq n \Rightarrow O(n)$ .

## Weighted Interval Scheduling: Bottom-Up

Bottom-up dynamic programming. Unwind recursion.

```
Input:  $n, s_1, \dots, s_n, f_1, \dots, f_n, v_1, \dots, v_n$ 
```

```
Sort jobs by finish times so that  $f_1 \leq f_2 \leq \dots \leq f_n$ .
```

```
Compute  $p(1), p(2), \dots, p(n)$ 
```

```
Iterative-Compute-Opt {  
     $M[0] = 0$   
    for  $j = 1$  to  $n$   
         $M[j] = \max(v_j + M[p(j)], M[j-1])$   
}
```

## 6.3 Segmented Least Squares

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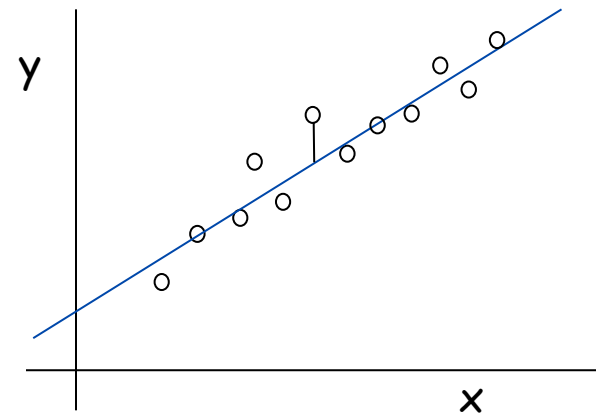


# Segmented Least Squares

## Least squares.

- Foundational problem in statistic and numerical analysis.
- Given  $n$  points in the plane:  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$ .
- Find a line  $y = ax + b$  that minimizes the sum of the squared error:

$$SSE = \sum_{i=1}^n (y_i - ax_i - b)^2$$



**Solution.** Calculus  $\Rightarrow$  min error is achieved when

$$a = \frac{n \sum_i x_i y_i - (\sum_i x_i) (\sum_i y_i)}{n \sum_i x_i^2 - (\sum_i x_i)^2}, \quad b = \frac{\sum_i y_i - a \sum_i x_i}{n}$$

# Segmented Least Squares

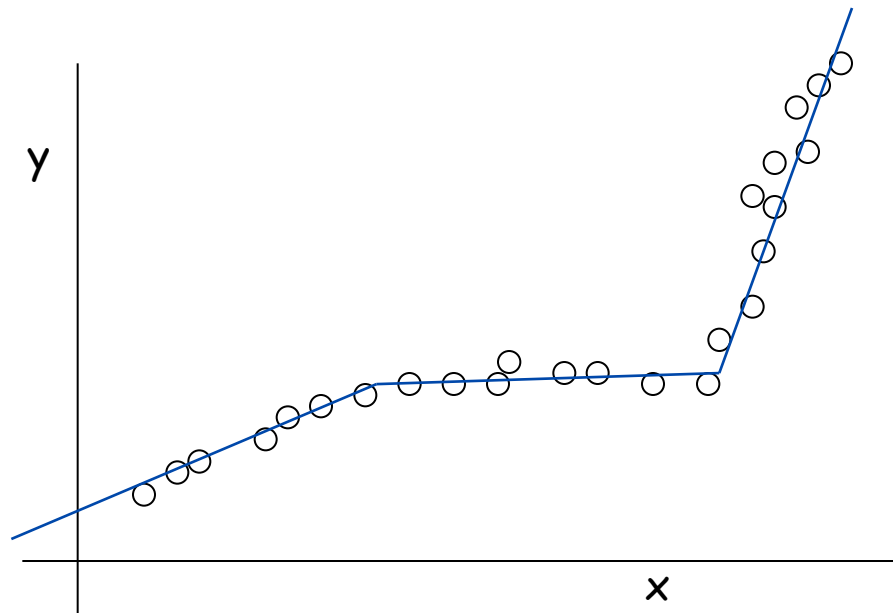
## Segmented least squares.

- Points lie roughly on a sequence of several line segments.
- Given  $n$  points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$  with
- $x_1 < x_2 < \dots < x_n$ , find a sequence of lines that minimizes  $f(x)$ .

Q. What's a reasonable choice for  $f(x)$  to balance accuracy and parsimony?

↑  
number of lines

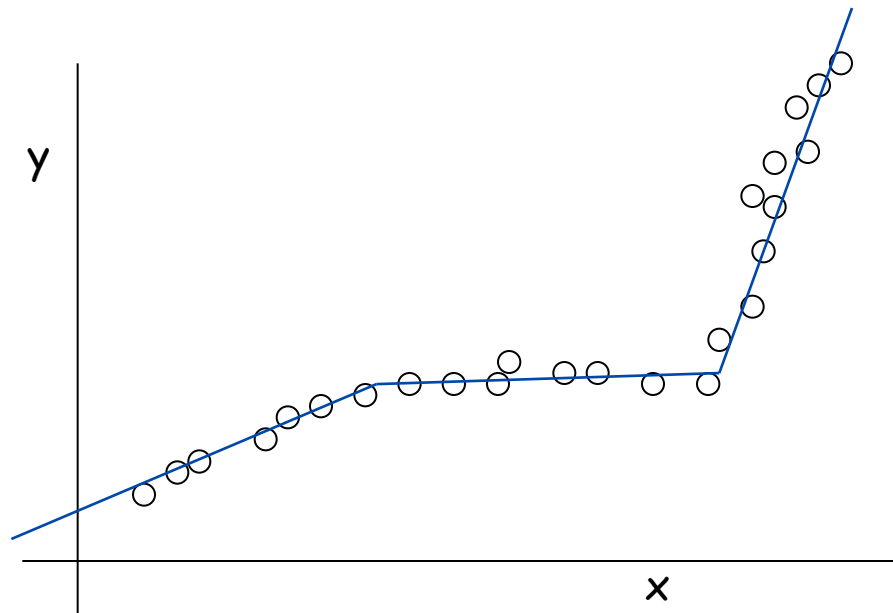
↑  
goodness of fit



# Segmented Least Squares

## Segmented least squares.

- Points lie roughly on a sequence of several line segments.
- Given  $n$  points in the plane  $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$  with
- $x_1 < x_2 < \dots < x_n$ , find a sequence of lines that minimizes:
  - the sum of the sums of the squared errors  $E$  in each segment
  - the number of lines  $L$
- Tradeoff function:  $E + c L$ , for some constant  $c > 0$ .



## Dynamic Programming: Multiway Choice

### Notation.

- $OPT(j)$  = minimum cost for points  $p_1, p_{i+1}, \dots, p_j$ .
- $e(i, j)$  = minimum sum of squares for points  $p_i, p_{i+1}, \dots, p_j$ .

### To compute $OPT(j)$ :

- Last segment uses points  $p_i, p_{i+1}, \dots, p_j$  for some  $i$ .
- Cost =  $e(i, j) + c + OPT(i-1)$ .

$$OPT(j) = \begin{cases} 0 & \text{if } j = 0 \\ \min_{1 \leq i \leq j} \{ e(i, j) + c + OPT(i-1) \} & \text{otherwise} \end{cases}$$


## Segmented Least Squares: Algorithm

```
INPUT:  $n, p_1, \dots, p_N, c$ 

Segmented-Least-Squares () {
  M[0] = 0
  for j = 1 to n
    for i = 1 to j
      compute the least square error  $e_{ij}$  for
      the segment  $p_i, \dots, p_j$ 

  for j = 1 to n
    M[j] =  $\min_{1 \leq i \leq j} (e_{ij} + c + M[i-1])$ 

  return M[n]
}
```

Running time.  $O(n^3)$ .  can be improved to  $O(n^2)$  by pre-computing various statistics

- Bottleneck = computing  $e(i, j)$  for  $O(n^2)$  pairs,  $O(n)$  per pair using previous formula.

## 6.4 Knapsack Problem

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# Knapsack Problem

## Knapsack problem.

- Given  $n$  objects and a "knapsack."
- Item  $i$  weighs  $w_i > 0$  kilograms and has value  $v_i > 0$ .
- Knapsack has capacity of  $W$  kilograms.
- Goal: fill knapsack so as to maximize total value.

Ex: { 3, 4 } has value 40.

$$W = 11$$

#	value	weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

*Greedy:* repeatedly add item with maximum ratio  $v_i / w_i$ .

Ex: { 5, 2, 1 } achieves only value = 35  $\Rightarrow$  greedy not optimal.

## Dynamic Programming: False Start

**Def.**  $OPT(i)$  = max profit subset of items  $1, \dots, i$ .

- Case 1:  $OPT$  does not select item  $i$ .
  - $OPT$  selects best of  $\{ 1, 2, \dots, i-1 \}$
- Case 2:  $OPT$  selects item  $i$ .
  - accepting item  $i$  does not immediately imply that we will have to reject other items
  - without knowing what other items were selected before  $i$ , we don't even know if we have enough room for  $i$

**Conclusion.** Need more sub-problems!



## Dynamic Programming: Adding a New Variable

Def.  $OPT(i, w)$  = max profit subset of items 1, ..., i with weight limit  $w$ .

- Case 1:  $OPT$  does not select item  $i$ .
  - $OPT$  selects best of  $\{ 1, 2, \dots, i-1 \}$  using weight limit  $w$
- Case 2:  $OPT$  selects item  $i$ .
  - new weight limit =  $w - w_i$
  - $OPT$  selects best of  $\{ 1, 2, \dots, i-1 \}$  using this new weight limit

$$OPT(i, w) = \begin{cases} 0 & \text{if } i = 0 \\ OPT(i-1, w) & \text{if } w_i > w \\ \max\{ OPT(i-1, w), v_i + OPT(i-1, w - w_i) \} & \text{otherwise} \end{cases}$$

## Knapsack Problem: Bottom-Up

Knapsack. Fill up an  $n$ -by- $W$  array.

```
Input:  $n, W, w_1, \dots, w_N, v_1, \dots, v_N$ 

for  $w = 0$  to  $W$ 
     $M[0, w] = 0$ 

for  $i = 1$  to  $n$ 
    for  $w = 1$  to  $W$ 
        if ( $w_i > w$ )
             $M[i, w] = M[i-1, w]$ 
        else
             $M[i, w] = \max \{M[i-1, w], v_i + M[i-1, w-w_i]\}$ 

return  $M[n, W]$ 
```

# Knapsack Algorithm

←—————  $W + 1$  —————→

		0	1	2	3	4	5	6	7	8	9	10	11
$n + 1$	$\phi$	0	0	0	0	0	0	0	0	0	0	0	0
	{1}	0	1	1	1	1	1	1	1	1	1	1	1
	{1, 2}	0	1	6	7	7	7	7	7	7	7	7	7
	{1, 2, 3}	0	1	6	7	7	18	19	24	25	25	25	25
	{1, 2, 3, 4}	0	1	6	7	7	18	22	24	28	29	29	40
	{1, 2, 3, 4, 5}	0	1	6	7	7	18	22	28	29	34	34	40

OPT: { 4, 3 }  
 value = 22 + 18 = 40

$W = 11$

Item	Value	Weight
1	1	1
2	6	2
3	18	5
4	22	6
5	28	7

## Knapsack Problem: Running Time

Running time.  $\Theta(nW)$ .

- Not polynomial in input size!
- "Pseudo-polynomial."
- Decision version of Knapsack is NP-complete. [Chapter 8]

Knapsack approximation algorithm. There exists a poly-time algorithm that produces a feasible solution that has value within 0.01% of optimum. [Section 11.8]

## 6.5 RNA Secondary Structure

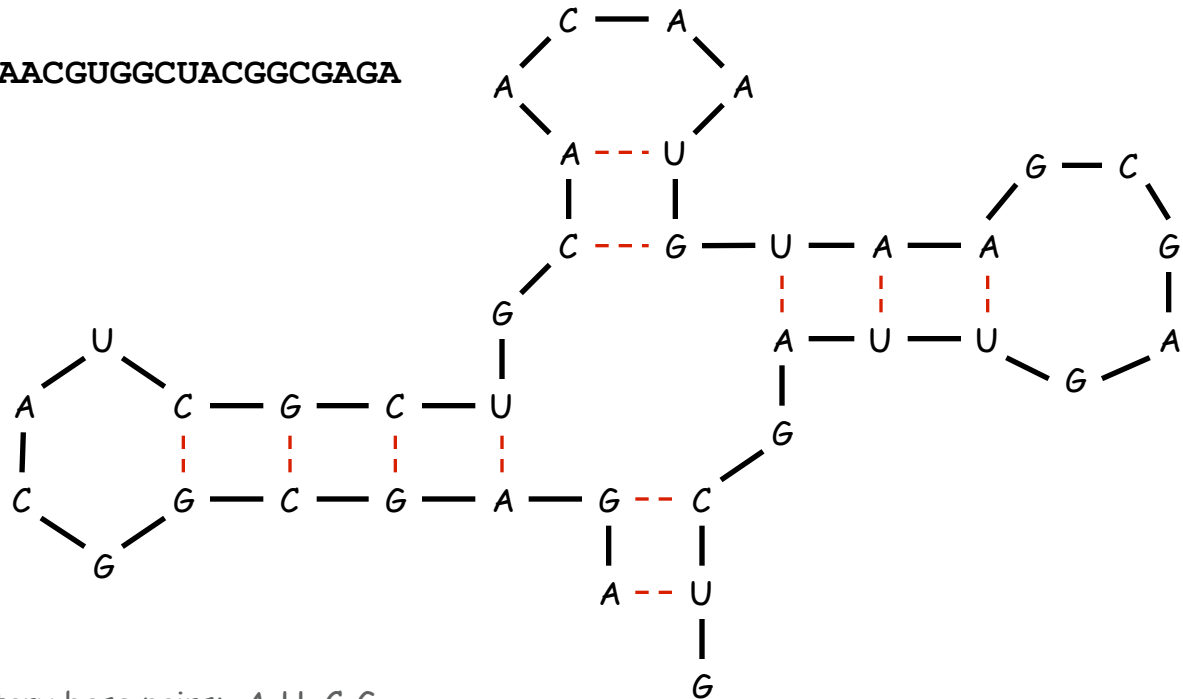
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# RNA Secondary Structure

**RNA.** String  $B = b_1b_2\dots b_n$  over alphabet  $\{A, C, G, U\}$ .

**Secondary structure.** RNA is single-stranded so it tends to loop back and form base pairs with itself. This structure is essential for understanding behavior of molecule.

**Ex:** GUCGAUUGAGCGAAUGUAACAACGUGGCUACGGCGAGA



# RNA Secondary Structure

**Secondary structure.** A set of pairs  $S = \{ (b_i, b_j) \}$  that satisfy:

- [Watson-Crick.]  $S$  is a matching and each pair in  $S$  is a Watson-Crick complement:  $A-U$ ,  $U-A$ ,  $C-G$ , or  $G-C$ .
- [No sharp turns.] The ends of each pair are separated by at least 4 intervening bases. If  $(b_i, b_j) \in S$ , then  $i < j - 4$ .
- [Non-crossing.] If  $(b_i, b_j)$  and  $(b_k, b_l)$  are two pairs in  $S$ , then we cannot have  $i < k < j < l$ .

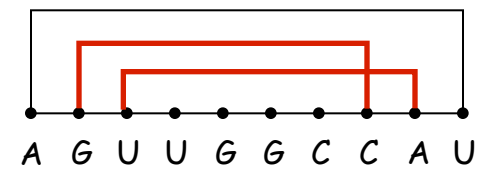
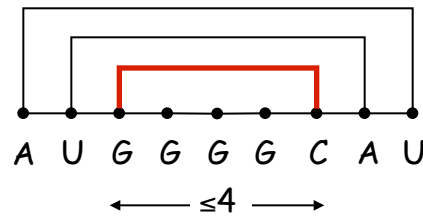
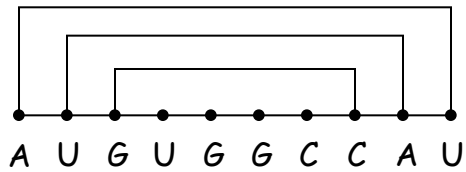
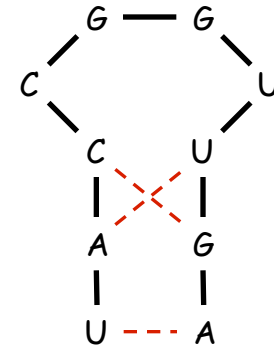
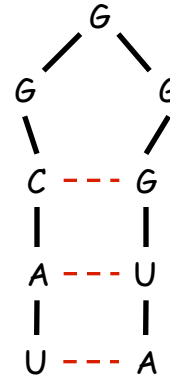
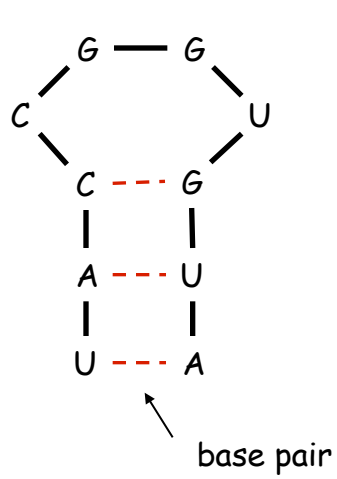
**Free energy.** Usual hypothesis is that an RNA molecule will form the secondary structure with the optimum total free energy.

↑  
approximate by number of base pairs

**Goal.** Given an RNA molecule  $B = b_1b_2\dots b_n$ , find a secondary structure  $S$  that maximizes the number of base pairs.

# RNA Secondary Structure: Examples

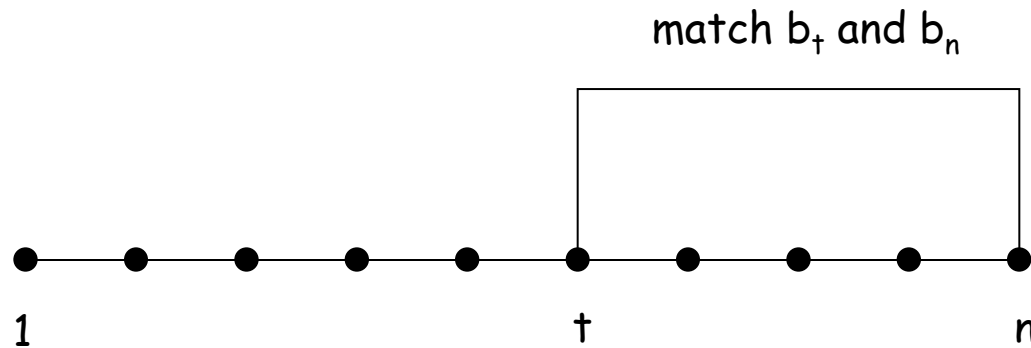
Examples.





## RNA Secondary Structure: Subproblems

**First attempt.**  $OPT(j)$  = maximum number of base pairs in a secondary structure of the substring  $b_1b_2\dots b_j$ .



**Difficulty.** Results in two sub-problems.

- Finding secondary structure in:  $b_1b_2\dots b_{t-1}$ . ←  $OPT(t-1)$
- Finding secondary structure in:  $b_{t+1}b_{t+2}\dots b_{n-1}$ . ← need more sub-problems

## Dynamic Programming Over Intervals

**Notation.**  $OPT(i, j)$  = maximum number of base pairs in a secondary structure of the substring  $b_i b_{i+1} \dots b_j$ .

- Case 1. If  $i \geq j - 4$ .
  - $OPT(i, j) = 0$  by no-sharp turns condition.
- Case 2. Base  $b_j$  is not involved in a pair.
  - $OPT(i, j) = OPT(i, j-1)$
- Case 3. Base  $b_j$  pairs with  $b_t$  for some  $i \leq t < j - 4$ .
  - non-crossing constraint decouples resulting sub-problems
  - $OPT(i, j) = 1 + \max_t \{ OPT(i, t-1) + OPT(t+1, j-1) \}$

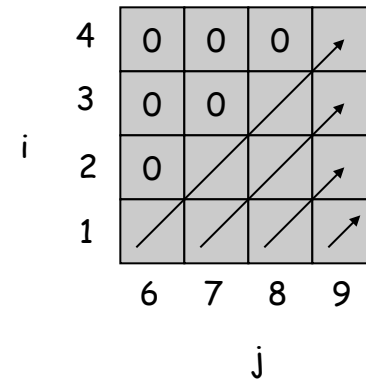
take max over  $t$  such that  $i \leq t < j-4$  and  $b_t$  and  $b_j$  are Watson-Crick complements

**Remark.** Same core idea in CKY algorithm to parse context-free grammars.

## Bottom Up Dynamic Programming Over Intervals

- Q. What order to solve the sub-problems?  
A. Do shortest intervals first.

```
RNA( $b_1, \dots, b_n$ ) {  
  for  $k = 5, 6, \dots, n-1$   
    for  $i = 1, 2, \dots, n-k$   
       $j = i + k$   
      Compute  $M[i, j]$   
  
  return  $M[1, n]$  using recurrence  
}
```



Running time.  $O(n^3)$ .

# Dynamic Programming Summary

## Recipe.

- Characterize structure of problem.
- Recursively define value of optimal solution.
- Compute value of optimal solution.
- Construct optimal solution from computed information.

## Dynamic programming techniques.

- Binary choice: weighted interval scheduling.
- Multi-way choice: segmented least squares.
- Adding a new variable: knapsack.
- Dynamic programming over intervals: RNA secondary structure.

Viterbi algorithm for HMM also uses DP to optimize a maximum likelihood tradeoff between parsimony and accuracy

CKY parsing algorithm for context-free grammar has similar structure

Top-down vs. bottom-up: different people have different intuitions.

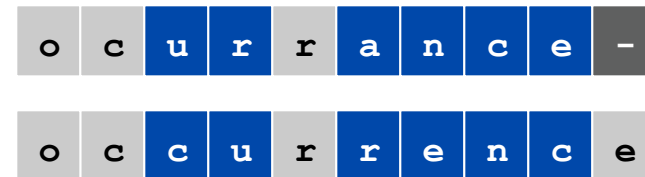
## 6.6 Sequence Alignment

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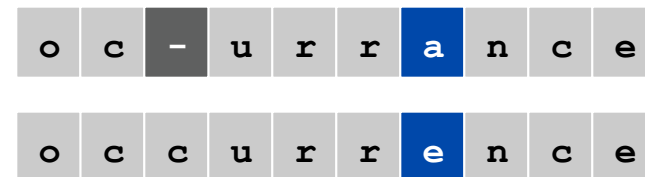
# String Similarity

How similar are two strings?

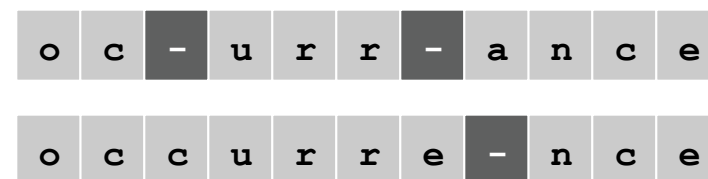
- **ocurrance**
- **occurrence**



6 mismatches, 1 gap



1 mismatch, 1 gap



0 mismatches, 3 gaps

# Edit Distance

## Applications.

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

Edit distance. [Levenshtein 1966, Needleman-Wunsch 1970]

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

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

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

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

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

## 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	C	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[i, 0] =  $i\delta$   
  for j = 0 to n  
    M[0, j] =  $j\delta$   
  
  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.

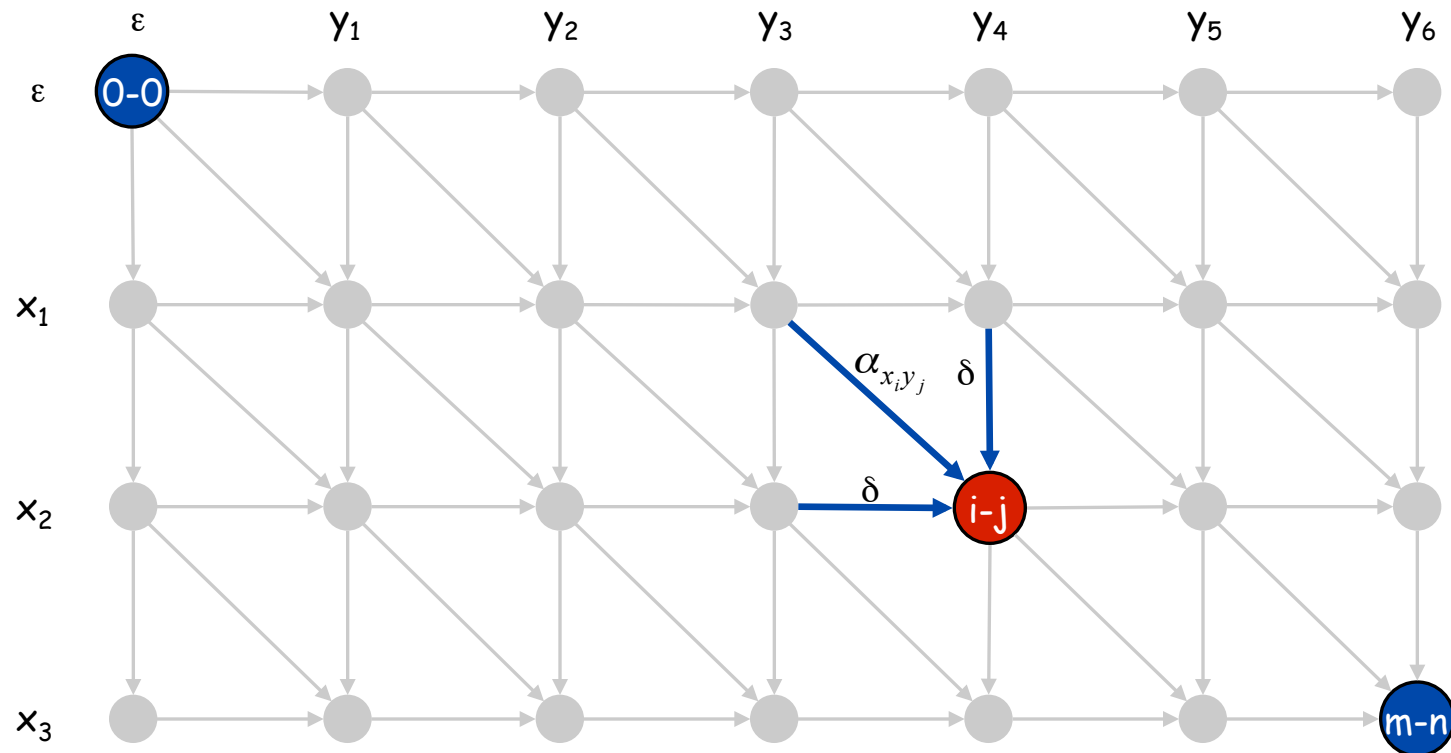
Theorem. [Hirschberg 1975] Optimal **alignment** in  $O(m + n)$  space and  $O(mn)$  time.

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

# Sequence Alignment: Linear Space

## Edit distance graph.

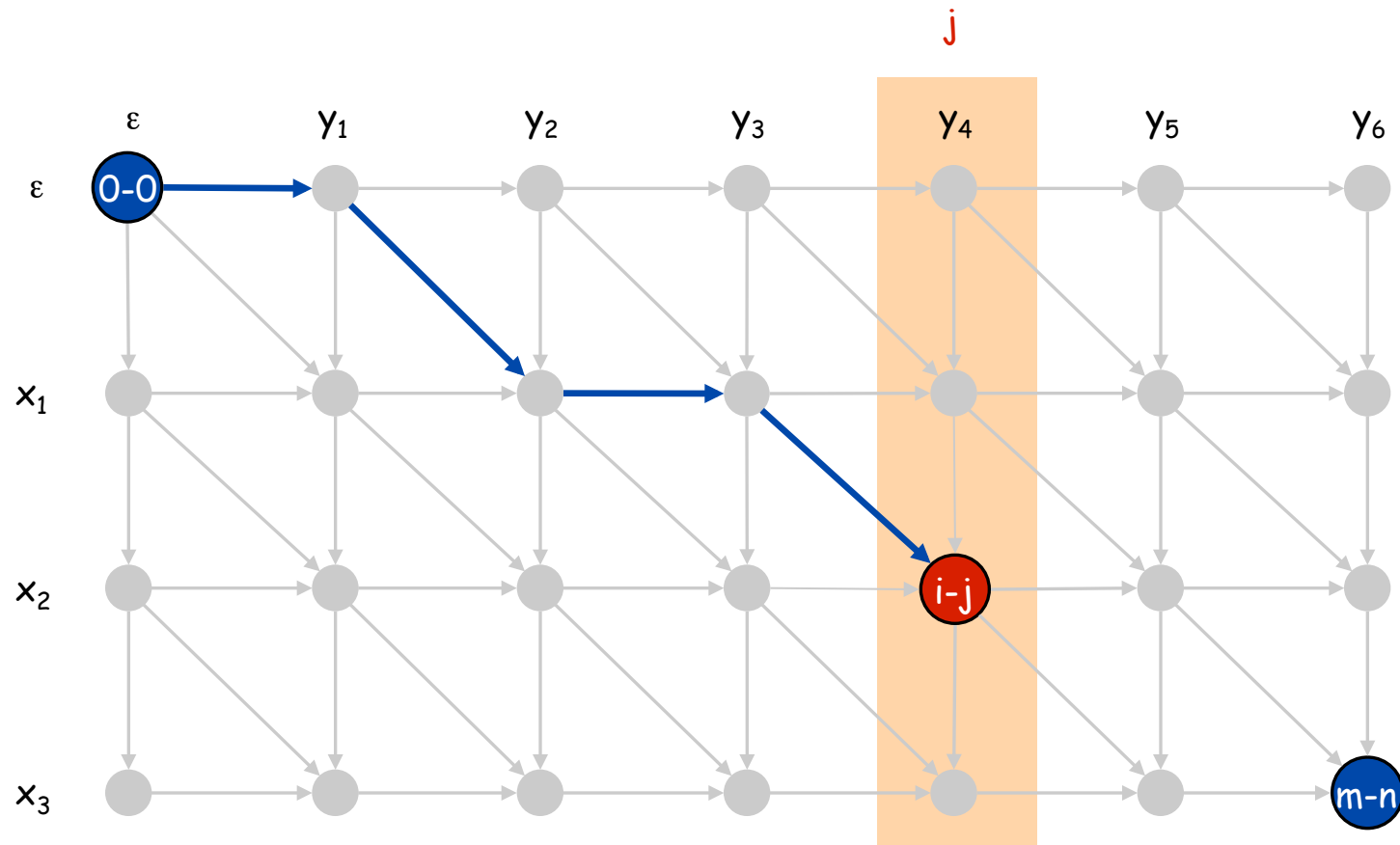
- Let  $f(i, j)$  be shortest path from  $(0,0)$  to  $(i, j)$ .
- Observation:  $f(i, j) = \text{OPT}(i, j)$ .



# Sequence Alignment: Linear Space

## Edit distance graph.

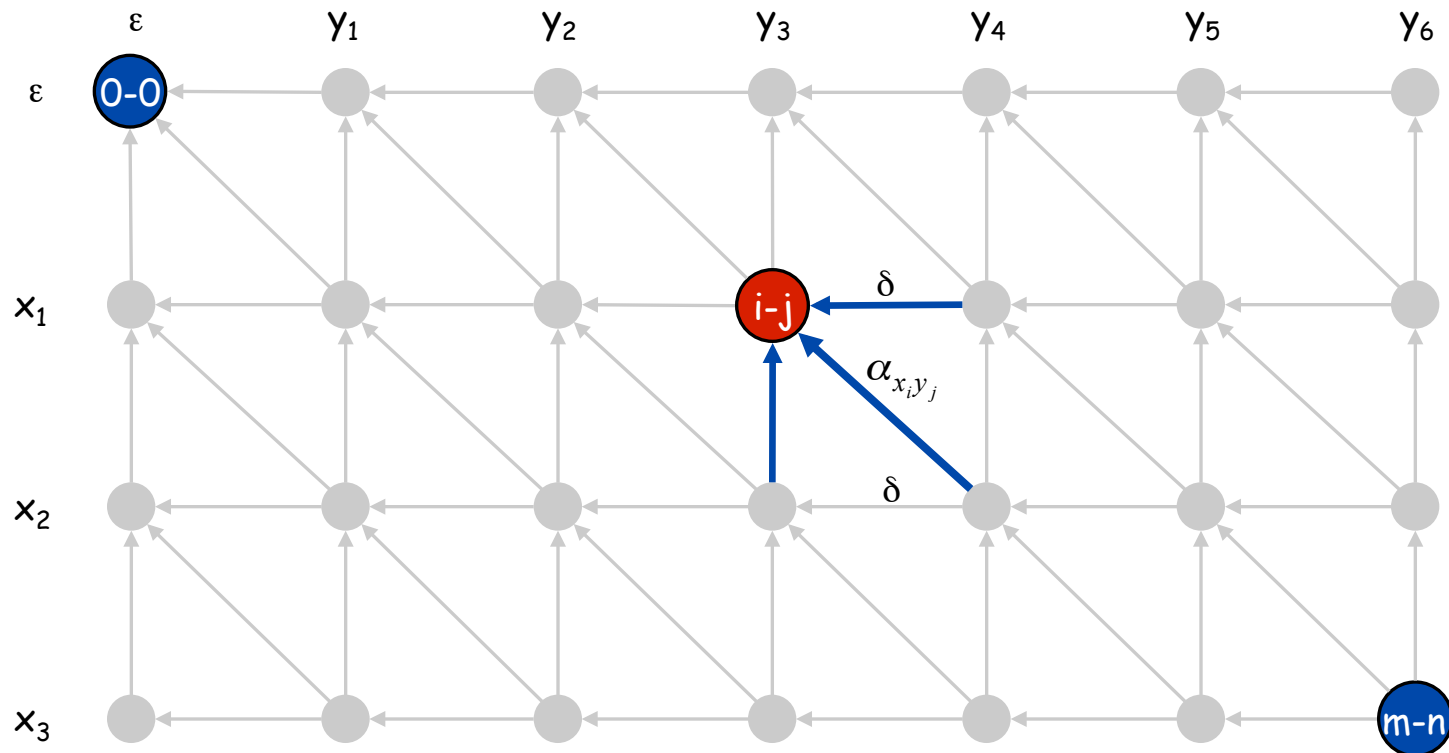
- Let  $f(i, j)$  be shortest path from  $(0,0)$  to  $(i, j)$ .
- Can compute  $f(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.



# Sequence Alignment: Linear Space

## Edit distance graph.

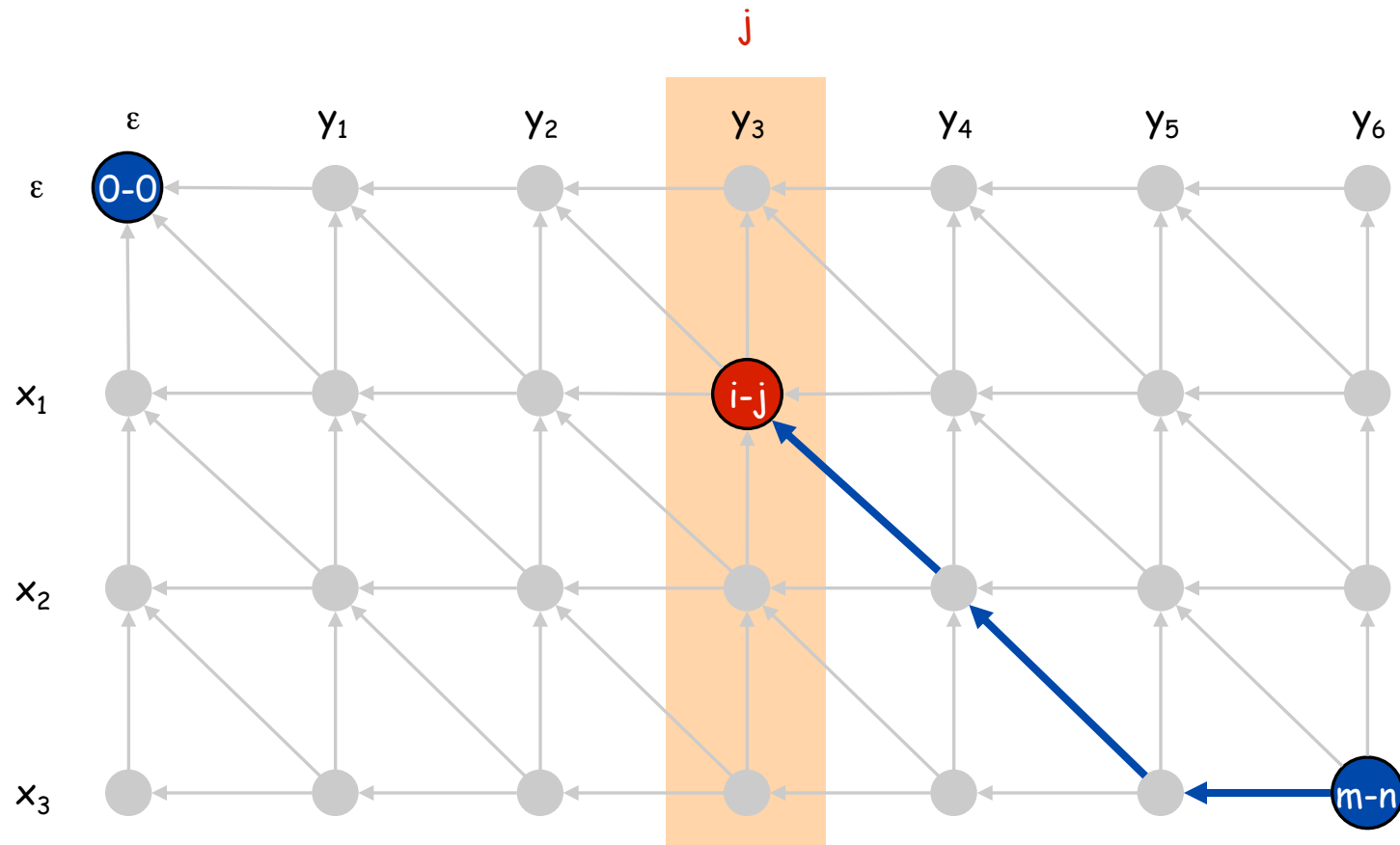
- Let  $g(i, j)$  be shortest path from  $(i, j)$  to  $(m, n)$ .
- Can compute by reversing the edge orientations and inverting the roles of  $(0, 0)$  and  $(m, n)$



# Sequence Alignment: Linear Space

## Edit distance graph.

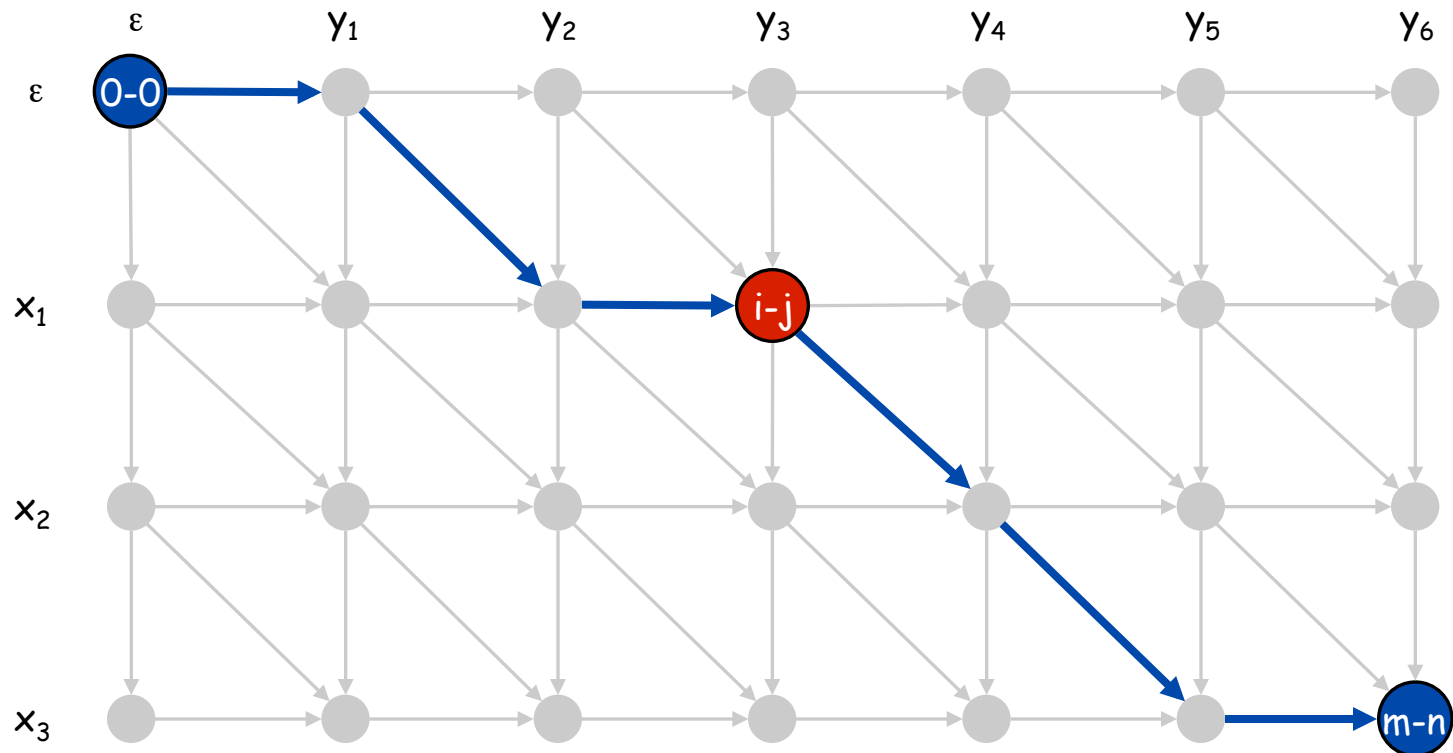
- Let  $g(i, j)$  be shortest path from  $(i, j)$  to  $(m, n)$ .
- Can compute  $g(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.





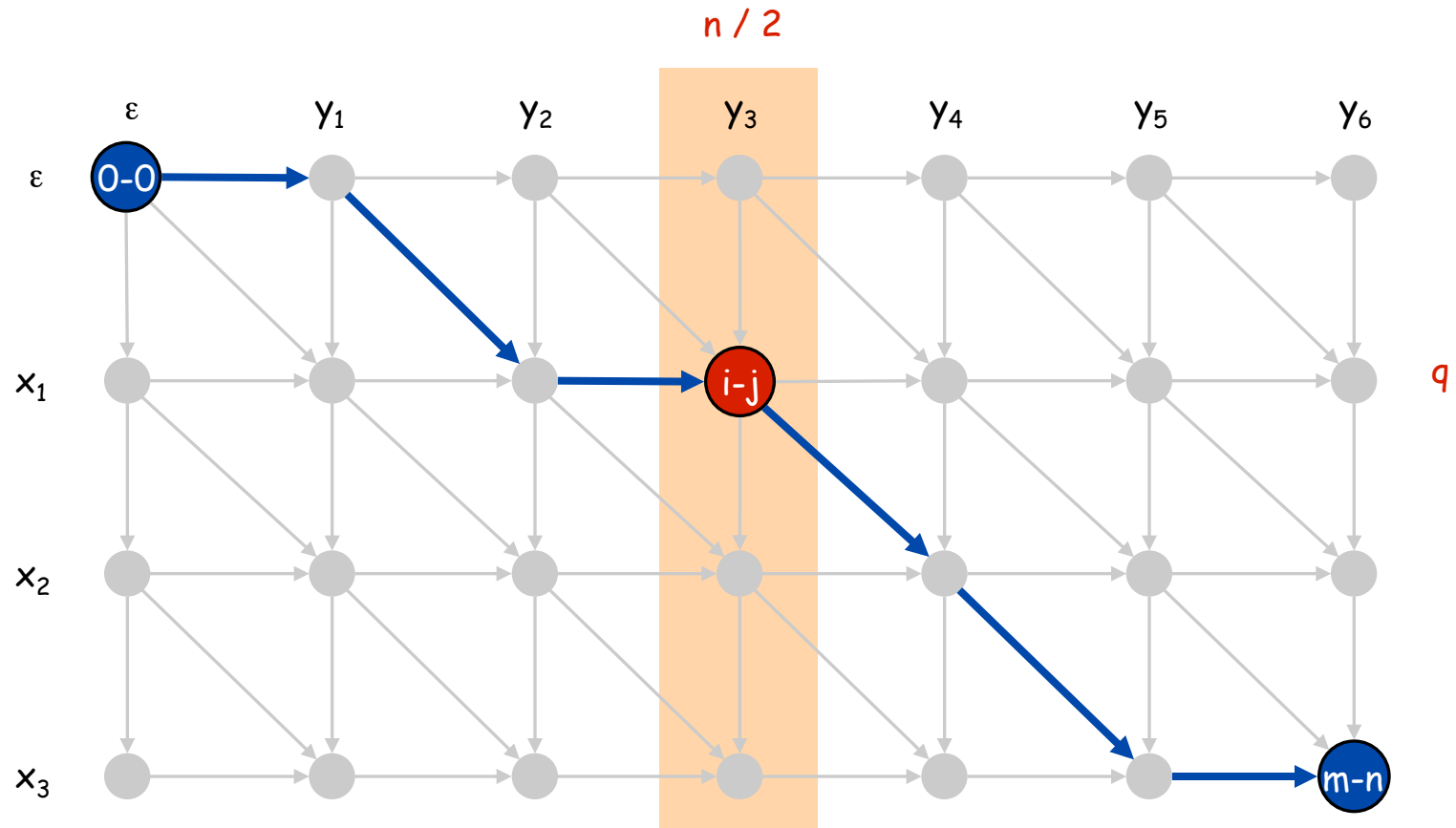
## Sequence Alignment: Linear Space

**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  $(q, n/2)$ .

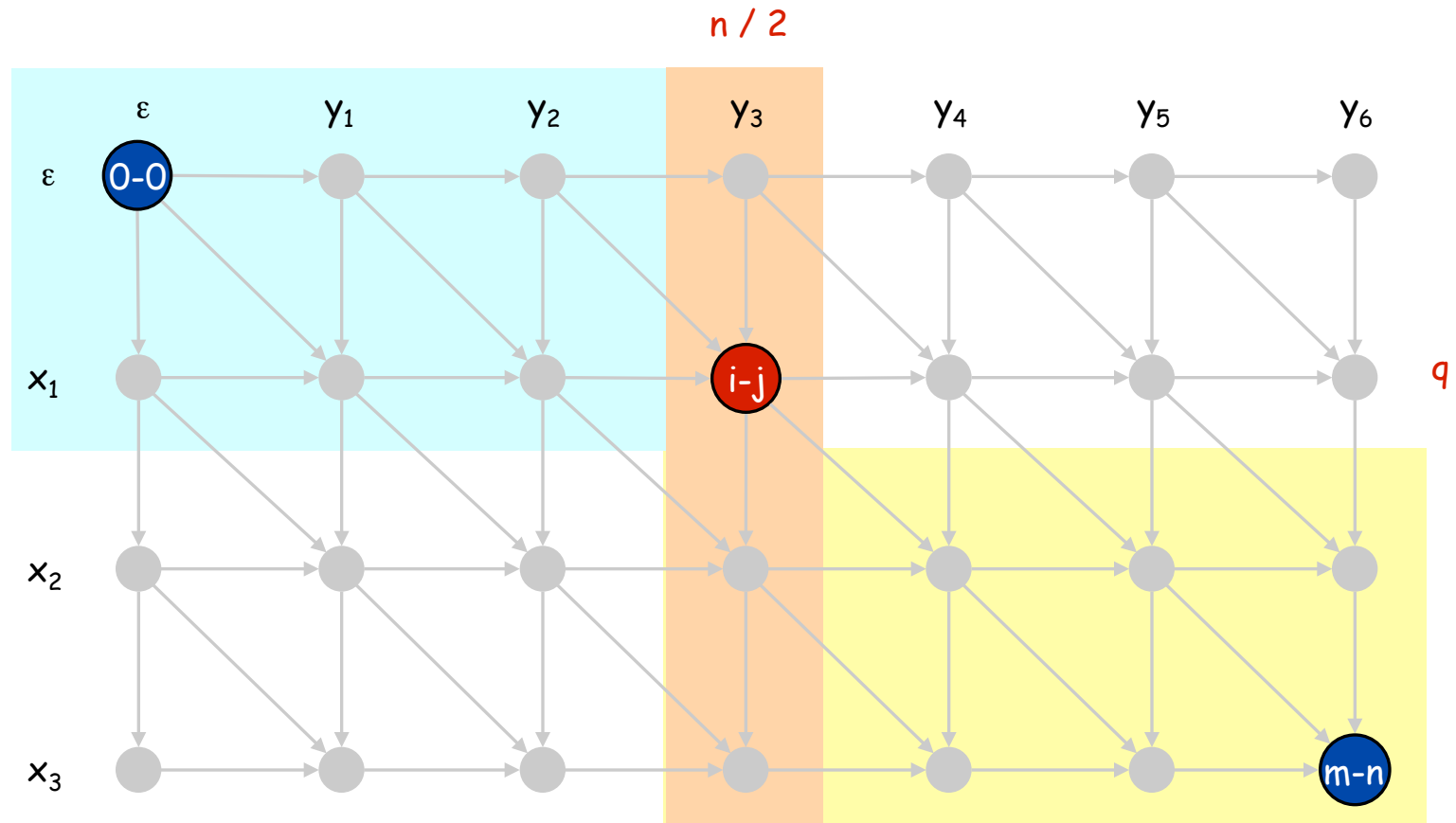


# Sequence Alignment: Linear Space

**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:  $T(m, n) \leq 2cmn$ .

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