

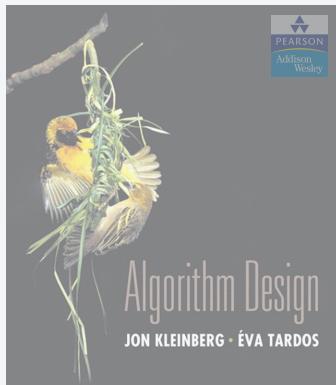
## 6. DYNAMIC PROGRAMMING II

- ▶ *sequence alignment*
- ▶ *Hirschberg's algorithm*
- ▶ *Bellman–Ford–Moore algorithm*
- ▶ *distance-vector protocols*
- ▶ *negative cycles*

Lecture slides by Kevin Wayne  
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<http://www.cs.princeton.edu/~wayne/kleinberg-tardos>

Last updated on 5/19/21 10:15 AM



## 6. DYNAMIC PROGRAMMING II

- ▶ *sequence alignment*
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- ▶ *distance-vector protocols*
- ▶ *negative cycles*

### SECTION 6.6

## String similarity

Q. How similar are two strings?

Ex. occurrence and occurrence.

o   c   u   r   r   a   n   c   e   -	o   c   -   u   r   r   a   n   c   e
o   c   c   u   r   r   e   n   c   e	o   c   c   u   r   r   e   n   c   e

6 mismatches, 1 gap

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

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

$$\text{cost} = \delta + \alpha_{CG} + \alpha_{TA}$$

assuming  $\alpha_{AA} = \alpha_{CC} = \alpha_{GG} = \alpha_{TT} = 0$

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

**Edit distance.** [Levenshtein 1966, Needleman–Wunsch 1970]

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

**Applications.** Bioinformatics, spell correction, machine translation, speech recognition, information extraction, ...

Spokesperson confirms senior government adviser was found  
Spokesperson said the senior adviser was found



## Sequence alignment: bottom-up algorithm

**SEQUENCE-ALIGNMENT**( $m, n, x_1, \dots, x_m, y_1, \dots, y_n, \delta, \alpha$ )

```

FOR i = 0 TO m
    M[i, 0] ← i δ.
FOR j = 0 TO n
    M[0, j] ← j δ.

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].

```

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## Sequence alignment: traceback

	P	A	L	A	T	E	
P	0	2	4	6	8	10	12
A	2	0	2	4	6	8	10
L	4	2	0	2	4	6	8
E	6	4	2	0	2	4	6
T	8	6	4	2	1	3	4
T	10	8	6	4	3	1	3
E	12	10	8	6	5	3	2
E	14	12	10	8	7	5	3

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## Sequence alignment: analysis

**Theorem.** The DP algorithm computes the edit distance (and an optimal alignment) of two strings of lengths  $m$  and  $n$  in  $\Theta(mn)$  time and space.

**Pf.**

- Algorithm computes edit distance.
- Can trace back to extract optimal alignment itself. ▀

**Theorem.** [Backurs-Indyk 2015] If can compute edit distance of two strings of length  $n$  in  $O(n^{2-\varepsilon})$  time for some constant  $\varepsilon > 0$ , then can solve SAT with  $n$  variables and  $m$  clauses in  $\text{poly}(m) 2^{(1-\delta)n}$  time for some constant  $\delta > 0$ .

Edit Distance Cannot Be Computed in Strongly Subquadratic Time (unless SETH is false)\*

which would disprove SETH  
(strong exponential time hypothesis)

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Piotr Indyk<sup>‡</sup>  
MIT

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## Dynamic programming: quiz 3

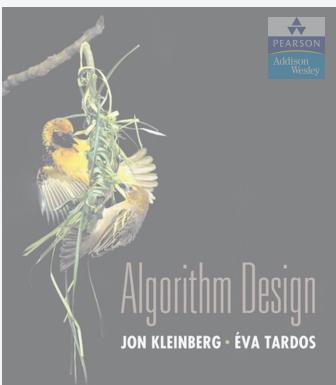


It is easy to modify the DP algorithm for edit distance to...

- Compute edit distance in  $O(mn)$  time and  $O(m + n)$  space.
- Compute an optimal alignment in  $O(mn)$  time and  $O(m + n)$  space.
- Both A and B.
- Neither A nor B.

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

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SECTION 6.7

## 6. DYNAMIC PROGRAMMING II

- ▶ sequence alignment
- ▶ Hirschberg's algorithm
- ▶ Bellman–Ford–Moore algorithm
- ▶ distance-vector protocols
- ▶ negative cycles

### Sequence alignment in linear space

**Theorem.** [Hirschberg] There exists an algorithm to find an optimal alignment in  $O(mn)$  time and  $O(m + n)$  space.

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

Programming Techniques G. Manacher Editor  
A Linear Space Algorithm for Computing Maximal Common Subsequences  
D.S. Hirschberg Princeton University

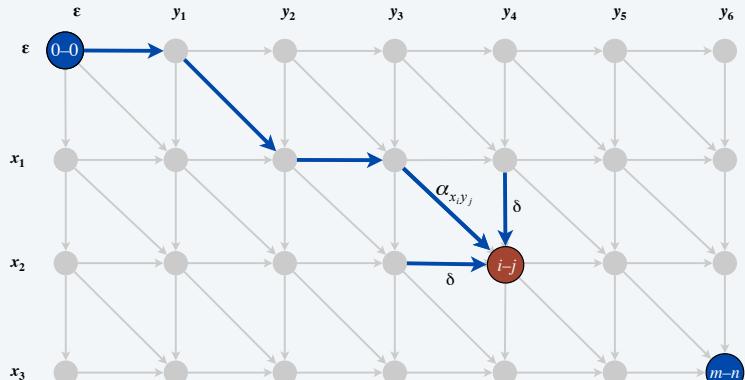
The problem of finding a longest common subsequence of two strings has been solved in quadratic time and space. An algorithm is presented which will solve this problem in quadratic time and in linear space.  
Key Words and Phrases: subsequence, longest common subsequence, string correction, editing  
CR Categories: 3.63, 3.73, 3.79, 4.22, 5.25

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### Hirschberg's algorithm

#### Edit distance graph.

- Let  $f(i, j)$  denote length of shortest path from  $(0, 0)$  to  $(i, j)$ .
- Lemma:  $f(i, j) = OPT(i, j)$  for all  $i$  and  $j$ .



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### Hirschberg's algorithm

#### Edit distance graph.

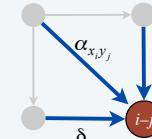
- Let  $f(i, j)$  denote length of shortest path from  $(0, 0)$  to  $(i, j)$ .
- Lemma:  $f(i, j) = OPT(i, j)$  for all  $i$  and  $j$ .

**Pf of Lemma.** [ by strong induction on  $i + j$  ]

- Base case:  $f(0, 0) = OPT(0, 0) = 0$ .
- Inductive hypothesis: assume true for all  $(i', j')$  with  $i' + j' < i + j$ .
- Last edge on shortest path to  $(i, j)$  is from  $(i - 1, j - 1)$ ,  $(i - 1, j)$ , or  $(i, j - 1)$ .
- Thus,

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

inductive hypothesis  
Bellman equation

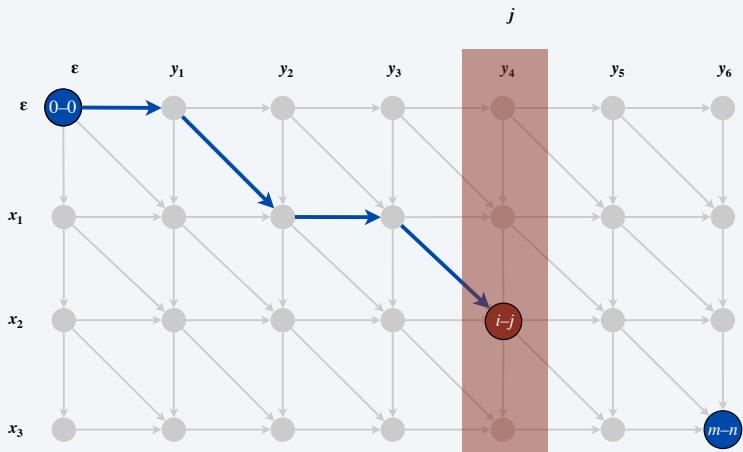


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## Hirschberg's algorithm

### Edit distance graph.

- Let  $f(i, j)$  denote length of shortest path from  $(0, 0)$  to  $(i, j)$ .
- Lemma:  $f(i, j) = OPT(i, j)$  for all  $i$  and  $j$ .
- Can compute  $f(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.

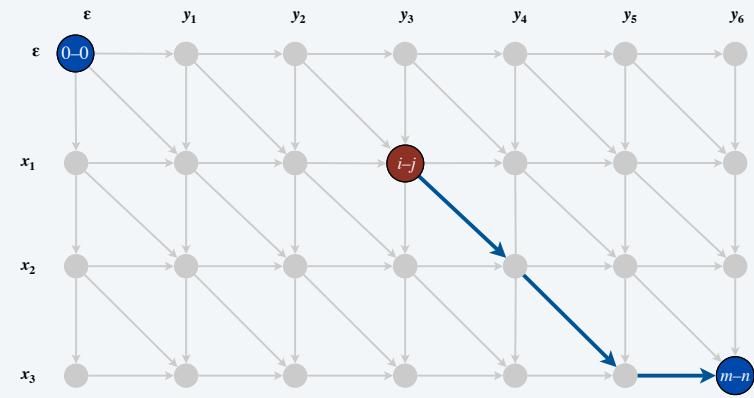


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## Hirschberg's algorithm

### Edit distance graph.

- Let  $g(i, j)$  denote length of shortest path from  $(i, j)$  to  $(m, n)$ .

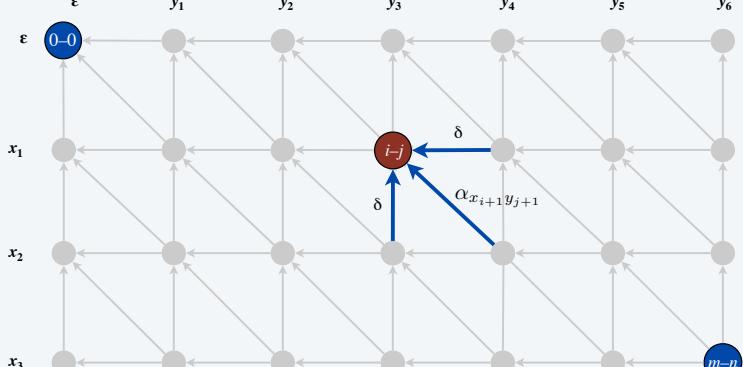


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## Hirschberg's algorithm

### Edit distance graph.

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

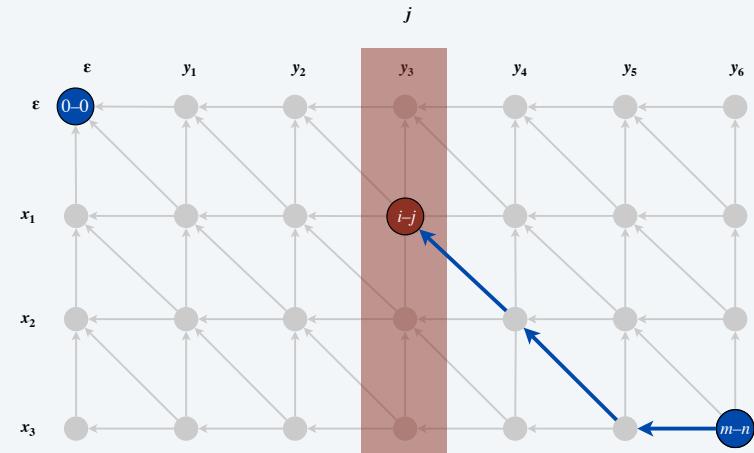


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## Hirschberg's algorithm

### Edit distance graph.

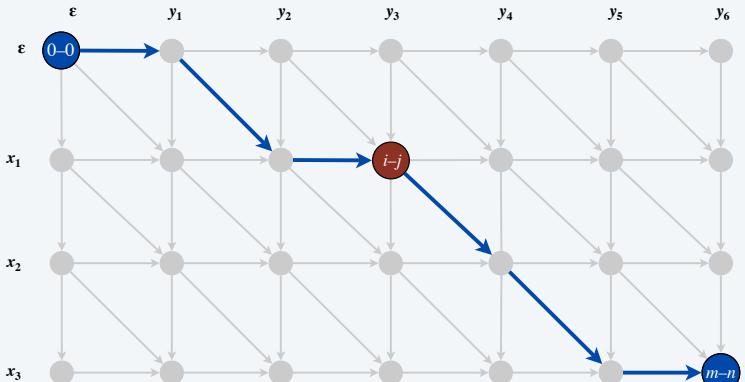
- Let  $g(i, j)$  denote length of shortest path from  $(i, j)$  to  $(m, n)$ .
- Can compute  $g(\cdot, j)$  for any  $j$  in  $O(mn)$  time and  $O(m + n)$  space.



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## Hirschberg's algorithm

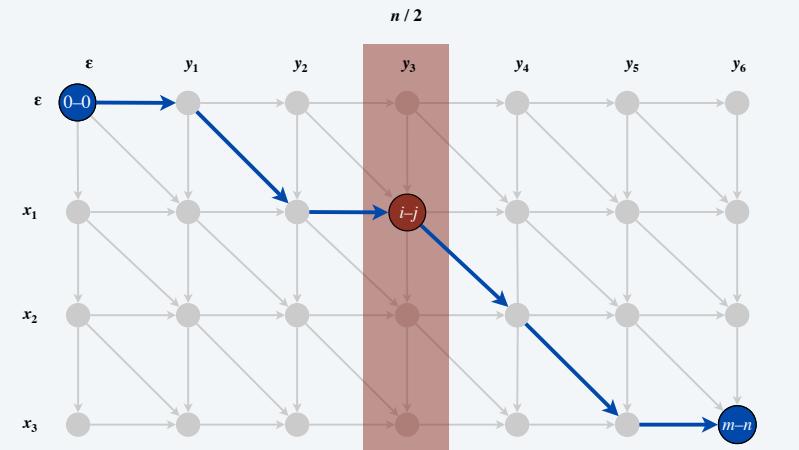
**Observation 1.** The length of a shortest path that uses  $(i, j)$  is  $f(i, j) + g(i, j)$ .



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## Hirschberg's algorithm

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

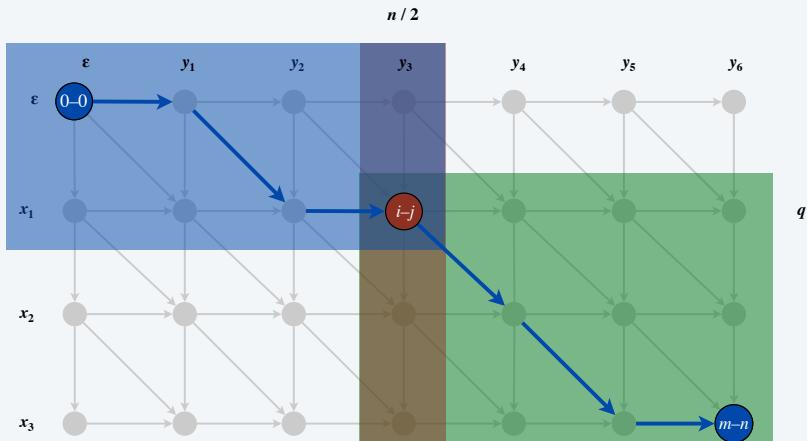


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## Hirschberg's algorithm

**Divide.** Find index  $q$  that minimizes  $f(q, n/2) + g(q, n/2)$ ; save node  $i-j$  as part of solution.

**Conquer.** Recursively compute optimal alignment in each piece.



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## Hirschberg's algorithm: space analysis

**Theorem.** Hirschberg's algorithm uses  $\Theta(m + n)$  space.

**Pf.**

- Each recursive call uses  $\Theta(m)$  space to compute  $f(\cdot, n/2)$  and  $g(\cdot, n/2)$ .
- Only  $\Theta(1)$  space needs to be maintained per recursive call.
- Number of recursive calls  $\leq n$ . ■

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## What is the worst-case running time of Hirschberg's algorithm?

- A.  $O(mn)$
- B.  $O(mn \log m)$
- C.  $O(mn \log n)$
- D.  $O(mn \log m \log n)$

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## Hirschberg's algorithm: running time analysis

**Theorem.** Let  $T(m, n) = \max$  running time of Hirschberg's algorithm on strings of lengths at most  $m$  and  $n$ . Then,  $T(m, n) = O(mn)$ .

**Pf.** [ by strong induction on  $m + n$  ]

- $O(mn)$  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:  $T(m, 2) \leq cm$
- $$T(2, n) \leq cn$$
- $$T(m, n) \leq cmn + T(q, n/2) + T(m - q, n/2)$$
- Claim.  $T(m, n) \leq 2cmn$ .
  - Base cases:  $m = 2$  and  $n = 2$ .
  - Inductive hypothesis:  $T(m, n) \leq 2cmn$  for all  $(m', n')$  with  $m' + n' < m + n$ .

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

inductive hypothesis

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## Hirschberg's algorithm: running time analysis warmup

**Theorem.** Let  $T(m, n) = \max$  running time of Hirschberg's algorithm on strings of lengths at most  $m$  and  $n$ . Then,  $T(m, n) = O(mn \log n)$ .

**Pf.**

- $T(m, n)$  is monotone nondecreasing in both  $m$  and  $n$ .
  - $T(m, n) \leq 2T(m, n/2) + O(mn)$
- $$\Rightarrow T(m, n) = O(mn \log n).$$

**Remark.** Analysis is not tight because two subproblems are of size  $(q, n/2)$  and  $(m - q, n/2)$ . Next, we prove  $T(m, n) = O(mn)$ .

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## LONGEST COMMON SUBSEQUENCE



**Problem.** Given two strings  $x_1 x_2 \dots x_m$  and  $y_1 y_2 \dots y_n$ , find a common subsequence that is as long as possible.

**Alternative viewpoint.** Delete some characters from  $x$ ; delete some character from  $y$ ; a common subsequence if it results in the same string.

**Ex.**  $\text{LCS}(\text{GGCACCAACG}, \text{ACGGCGGATACG}) = \text{GGCAACG}$ .

**Applications.** Unix diff, git, bioinformatics.

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**Solution 1.** Dynamic programming.

**Def.**  $OPT(i, j)$  = length of LCS of prefix strings  $x_1 x_2 \dots x_i$  and  $y_1 y_2 \dots y_j$ .

**Goal.**  $OPT(m, n)$ .

**Case 1.**  $x_i = y_j$ .

- $1 + \text{length of LCS of } x_1 x_2 \dots x_{i-1} \text{ and } y_1 y_2 \dots y_{j-1}$ . ← optimal substructure property (proof via exchange argument)

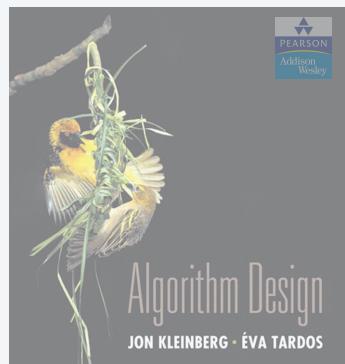
**Case 2.**  $x_i \neq y_j$ .

- Delete  $x_i$ : length of LCS of  $x_1 x_2 \dots x_{i-1}$  and  $y_1 y_2 \dots y_j$ .
- Delete  $y_j$ : length of LCS of  $x_1 x_2 \dots x_i$  and  $y_1 y_2 \dots y_{j-1}$ .

Bellman equation.

$$OPT(i, j) = \begin{cases} 0 & \text{if } i = 0 \text{ or } j = 0 \\ 1 + OPT(i - 1, j - 1) & \text{if } x_i = y_j \\ \max \{OPT(i - 1, j), OPT(i, j - 1)\} & \text{if } x_i \neq y_j \end{cases}$$

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SECTION 6.8

## 6. DYNAMIC PROGRAMMING II

- ▶ sequence alignment
- ▶ Hirschberg's algorithm
- ▶ Bellman–Ford–Moore algorithm
- ▶ distance-vector protocols
- ▶ negative cycles



**Solution 2.** Reduce to finding a min-cost alignment of  $x$  and  $y$  with

- Gap penalty  $\delta = 1$
- Mismatch penalty  $\alpha_{pq} = \begin{cases} 0 & \text{if } p = q \\ \infty & \text{if } p \neq q \end{cases}$
- Edit distance = # gaps = number of characters deleted from  $x$  and  $y$ .
- Length of LCS =  $(m + n - \text{edit distance}) / 2$ .

**Analysis.**  $O(mn)$  time and  $O(m + n)$  space.

**Lower bound.** No  $O(n^{2-\epsilon})$  algorithm unless SETH is false.

Tight Hardness Results for LCS and other Sequence Similarity Measures

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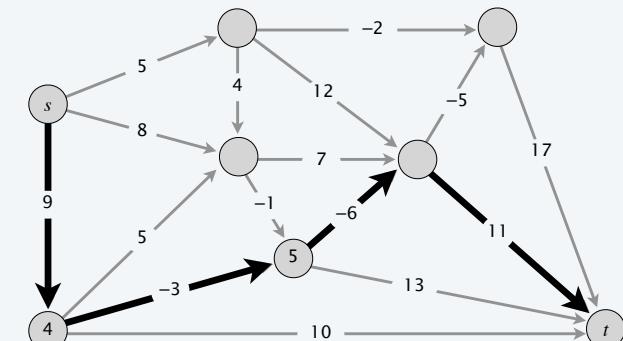
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## Shortest paths with negative weights

**Shortest-path problem.** Given a digraph  $G = (V, E)$ , with arbitrary edge lengths  $\ell_{vw}$ , find shortest path from source node  $s$  to destination node  $t$ .

assume there exists a path from every node to  $t$

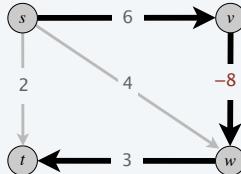


length of shortest  $s \xrightarrow{w} t$  path =  $9 - 3 - 6 + 11 = 11$

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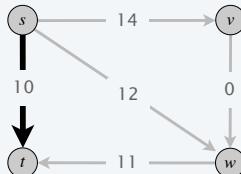
## Shortest paths with negative weights: failed attempts

Dijkstra. May not produce shortest paths when edge lengths are negative.



Dijkstra selects the vertices in the order  $s, t, w, v$ .  
But shortest path from  $s$  to  $t$  is  $s \rightarrow v \rightarrow w \rightarrow t$ .

**Reweighting.** Adding a constant to every edge length does not necessarily make Dijkstra's algorithm produce shortest paths.

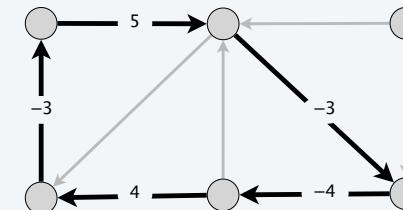


Adding 8 to each edge weight changes the shortest path from  $s \rightarrow v \rightarrow w \rightarrow t$  to  $s \rightarrow t$ .

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## Negative cycles

**Def.** A **negative cycle** is a directed cycle for which the sum of its edge lengths is negative.



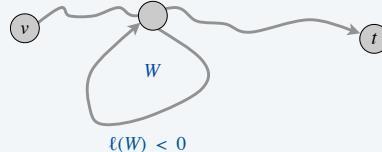
a negative cycle  $W$ :  $\ell(W) = \sum_{e \in W} \ell_e < 0$

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## Shortest paths and negative cycles

**Lemma 1.** If some  $v \rightsquigarrow t$  path contains a negative cycle, then there does not exist a shortest  $v \rightsquigarrow t$  path.

**Pf.** If there exists such a cycle  $W$ , then can build a  $v \rightsquigarrow t$  path of arbitrarily negative length by detouring around  $W$  as many times as desired. ▀



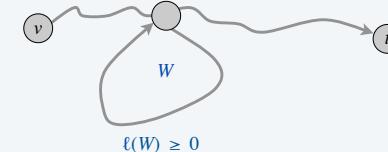
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## Shortest paths and negative cycles

**Lemma 2.** If  $G$  has no negative cycles, then there exists a shortest  $v \rightsquigarrow t$  path that is simple (and has  $\leq n - 1$  edges).

**Pf.**

- Among all shortest  $v \rightsquigarrow t$  paths, consider one that uses the fewest edges.
- If that path  $P$  contains a directed cycle  $W$ , can remove the portion of  $P$  corresponding to  $W$  without increasing its length. ▀

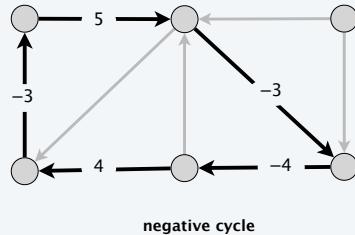
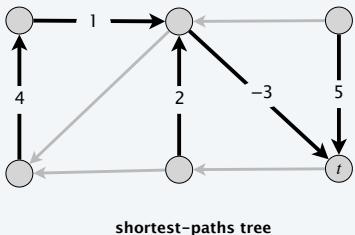


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## Shortest-paths and negative-cycle problems

**Single-destination shortest-paths problem.** Given a digraph  $G = (V, E)$  with edge lengths  $\ell_{vw}$  (but no negative cycles) and a distinguished node  $t$ , find a shortest  $v \rightsquigarrow t$  path for every node  $v$ .

**Negative-cycle problem.** Given a digraph  $G = (V, E)$  with edge lengths  $\ell_{vw}$ , find a negative cycle (if one exists).



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## Dynamic programming: quiz 5



Which subproblems to find shortest  $v \rightsquigarrow t$  paths for every node  $v$ ?

- A.  $OPT(i, v) =$  length of shortest  $v \rightsquigarrow t$  path that uses exactly  $i$  edges.
- B.  $OPT(i, v) =$  length of shortest  $v \rightsquigarrow t$  path that uses at most  $i$  edges.
- C. Neither A nor B.

## Shortest paths with negative weights: dynamic programming

**Def.**  $OPT(i, v) =$  length of shortest  $v \rightsquigarrow t$  path that uses  $\leq i$  edges.

**Goal.**  $OPT(n - 1, v)$  for each  $v$ .

by Lemma 2, if no negative cycles,  
there exists a shortest  $v \rightsquigarrow t$  path that is simple

**Case 1.** Shortest  $v \rightsquigarrow t$  path uses  $\leq i - 1$  edges.

- $OPT(i, v) = OPT(i - 1, v)$ .

optimal substructure property  
(proof via exchange argument)

**Case 2.** Shortest  $v \rightsquigarrow t$  path uses exactly  $i$  edges.

- if  $(v, w)$  is first edge in shortest such  $v \rightsquigarrow t$  path, incur a cost of  $\ell_{vw}$ .
- Then, select best  $w \rightsquigarrow t$  path using  $\leq i - 1$  edges.

### Bellman equation.

$$OPT(i, v) = \begin{cases} 0 & \text{if } i = 0 \text{ and } v = t \\ \infty & \text{if } i = 0 \text{ and } v \neq t \\ \min \left\{ OPT(i - 1, v), \min_{(v, w) \in E} \{ OPT(i - 1, w) + \ell_{vw} \} \right\} & \text{if } i > 0 \end{cases}$$

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## Shortest paths with negative weights: implementation

**SHORTEST-PATHS( $V, E, \ell, t$ )**

**FOREACH** node  $v \in V$ :

$M[0, v] \leftarrow \infty$ .

$M[0, t] \leftarrow 0$ .

**FOR**  $i = 1$  TO  $n - 1$

**FOREACH** node  $v \in V$ :

$M[i, v] \leftarrow M[i - 1, v]$ .

**FOREACH** edge  $(v, w) \in E$ :

$M[i, v] \leftarrow \min \{ M[i, v], M[i - 1, w] + \ell_{vw} \}$ .

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## Shortest paths with negative weights: implementation

**Theorem 1.** Given a digraph  $G = (V, E)$  with no negative cycles, the DP algorithm computes the length of a shortest  $v \rightsquigarrow t$  path for every node  $v$  in  $\Theta(mn)$  time and  $\Theta(n^2)$  space.

Pf.

- Table requires  $\Theta(n^2)$  space.
- Each iteration  $i$  takes  $\Theta(m)$  time since we examine each edge once. ▀

### Finding the shortest paths.

- Approach 1: Maintain  $\text{successor}[i, v]$  that points to next node on a shortest  $v \rightsquigarrow t$  path using  $\leq i$  edges.
- Approach 2: Compute optimal lengths  $M[i, v]$  and consider only edges with  $M[i, v] = M[i - 1, w] + \ell_{vw}$ . Any directed path in this subgraph is a shortest path.

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## Dynamic programming: quiz 6

It is easy to modify the DP algorithm for shortest paths to...

- A. Compute lengths of shortest paths in  $O(mn)$  time and  $O(m + n)$  space.
- B. Compute shortest paths in  $O(mn)$  time and  $O(m + n)$  space.
- C. Both A and B.
- D. Neither A nor B.

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## Shortest paths with negative weights: practical improvements

**Space optimization.** Maintain two 1D arrays (instead of 2D array).

- $d[v]$  = length of a shortest  $v \rightsquigarrow t$  path that we have found so far.
- $\text{successor}[v]$  = next node on a  $v \rightsquigarrow t$  path.

**Performance optimization.** If  $d[w]$  was not updated in iteration  $i - 1$ , then no reason to consider edges entering  $w$  in iteration  $i$ .

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## Bellman–Ford–Moore: efficient implementation

**BELLMAN–FORD–MOORE( $V, E, c, t$ )**

FOREACH node  $v \in V$ :

$d[v] \leftarrow \infty$ .  
 $\text{successor}[v] \leftarrow \text{null}$ .  
 $d[t] \leftarrow 0$ .

FOR  $i = 1$  TO  $n - 1$

FOREACH node  $w \in V$ :

IF ( $d[w]$  was updated in previous pass)  
FOREACH edge  $(v, w) \in E$  :  
IF ( $d[v] > d[w] + \ell_{vw}$ )  
 $d[v] \leftarrow d[w] + \ell_{vw}$ .  
 $\text{successor}[v] \leftarrow w$ .  
IF (no  $d[\cdot]$  value changed in pass  $i$ ) STOP.

pass  $i$   
 $O(m)$  time

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## Bellman–Ford–Moore: finding the shortest paths

**Lemma 6.** Any directed cycle  $W$  in the successor graph is a negative cycle.

Pf.

- If  $\text{successor}[v] = w$ , we must have  $d[v] \geq d[w] + \ell_{vw}$ .

(LHS and RHS are equal when  $\text{successor}[v]$  is set;  $d[w]$  can only decrease;  $d[v]$  decreases only when  $\text{successor}[v]$  is reset)

- Let  $v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_k \rightarrow v_1$  be the sequence of nodes in a directed cycle  $W$ .
- Assume that  $(v_k, v_1)$  is the last edge in  $W$  added to the successor graph.
- Just prior to that:  $d[v_1] \geq d[v_2] + \ell(v_1, v_2)$

$$d[v_2] \geq d[v_3] + \ell(v_2, v_3)$$

$$\vdots \quad \vdots \quad \vdots$$

$$d[v_{k-1}] \geq d[v_k] + \ell(v_{k-1}, v_k)$$

$$d[v_k] > d[v_1] + \ell(v_k, v_1) \quad \text{holds with strict inequality since we are updating } d[v_k]$$

- Adding inequalities yields  $\ell(v_1, v_2) + \ell(v_2, v_3) + \dots + \ell(v_{k-1}, v_k) + \ell(v_k, v_1) < 0$ . ▀

$\overbrace{\hspace{236pt}}$

W is a negative cycle

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## Bellman–Ford–Moore: finding the shortest paths

**Theorem 3.** Assuming no negative cycles, Bellman–Ford–Moore finds shortest  $v \rightsquigarrow t$  paths for every node  $v$  in  $O(mn)$  time and  $\Theta(n)$  extra space.

Pf.

- The successor graph cannot have a directed cycle. [Lemma 6]
- Thus, following the successor pointers from  $v$  yields a directed path to  $t$ .
- Let  $v = v_1 \rightarrow v_2 \rightarrow \dots \rightarrow v_k = t$  be the nodes along this path  $P$ .
- Upon termination, if  $\text{successor}[v] = w$ , we must have  $d[v] = d[w] + \ell_{vw}$ .

(LHS and RHS are equal when  $\text{successor}[v]$  is set;  $d[\cdot]$  did not change)

- Thus,  $d[v_1] = d[v_2] + \ell(v_1, v_2)$
- $d[v_2] = d[v_3] + \ell(v_2, v_3)$
- $\vdots \quad \vdots \quad \vdots$
- $d[v_{k-1}] = d[v_k] + \ell(v_{k-1}, v_k)$

since algorithm terminated

- Adding equations yields  $d[v] = d[t] + \ell(v_1, v_2) + \ell(v_2, v_3) + \dots + \ell(v_{k-1}, v_k)$ . ▀



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## Single-source shortest paths with negative weights

year	worst case	discovered by
1955	$O(n^4)$	Shimbel
1956	$O(m n^2 W)$	Ford
1958	$O(m n)$	Bellman, Moore
1983	$O(n^{3/4} m \log W)$	Gabow
1989	$O(m n^{1/2} \log(nW))$	Gabow–Tarjan
1993	$O(m n^{1/2} \log W)$	Goldberg
2005	$O(n^{2.38} W)$	Sankowski, Yuster–Zwick
2016	$\tilde{O}(n^{10/7} \log W)$	Cohen–Mądry–Sankowski–Vlădu
20xx	???	

single-source shortest paths with weights between  $-W$  and  $W$



SECTION 6.9

## 6. DYNAMIC PROGRAMMING II

- ▶ sequence alignment
- ▶ Hirschberg's algorithm
- ▶ Bellman–Ford–Moore algorithm
- ▶ distance-vector protocols
- ▶ negative cycles

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## Distance-vector routing protocols

### Communication network.

- Node  $\approx$  router.
- Edge  $\approx$  direct communication link.
- Length of edge  $\approx$  latency of link.  $\leftarrow$  non-negative, but Bellman–Ford–Moore used anyway!

Dijkstra's algorithm. Requires global information of network.

Bellman–Ford–Moore. Uses only local knowledge of neighboring nodes.

**Synchronization.** We don't expect routers to run in lockstep. The order in which each edges are processed in Bellman–Ford–Moore is not important. Moreover, algorithm converges even if updates are asynchronous.

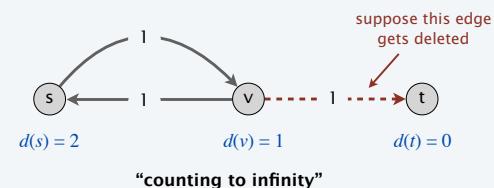
## Distance-vector routing protocols

### Distance-vector routing protocols. [ “routing by rumor” ]

- Each router maintains a vector of shortest-path lengths to every other node (distances) and the first hop on each path (directions).
- Algorithm: each router performs  $n$  separate computations, one for each potential destination node.

**Ex.** RIP, Xerox XNS RIP, Novell's IPX RIP, Cisco's IGRP, DEC's DNA Phase IV, AppleTalk's RTMP.

**Caveat.** Edge lengths may change during algorithm (or fail completely).



## Path-vector routing protocols

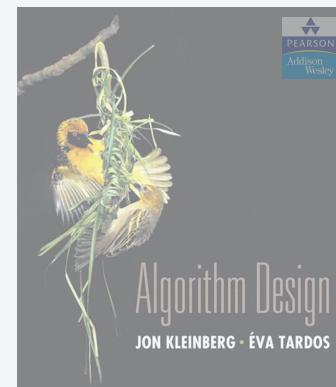
### Link-state routing protocols.

not just the distance and first hop

- Each router stores the whole path (or network topology).
- Based on Dijkstra's algorithm.
- Avoids “counting-to-infinity” problem and related difficulties.
- Requires significantly more storage.

**Ex.** Border Gateway Protocol (BGP), Open Shortest Path First (OSPF).

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SECTION 6.10

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## 6. DYNAMIC PROGRAMMING II

- ▶ sequence alignment
- ▶ Hirschberg's algorithm
- ▶ Bellman–Ford–Moore algorithm
- ▶ distance vector protocol
- ▶ negative cycles



## Detecting negative cycles

**Theorem 5.** Can find a negative cycle in  $O(mn)$  time and  $O(n)$  extra space.

Pf.

- Run Bellman–Ford–Moore on  $G'$  for  $n' = n + 1$  passes (instead of  $n' - 1$ ).
- If no  $d[v]$  values updated in pass  $n'$ , then no negative cycles.
- Otherwise, suppose  $d[s]$  updated in pass  $n'$ .
- Define  $\text{pass}(v) = \text{last pass in which } d[v] \text{ was updated}$ .
- Observe  $\text{pass}(s) = n'$  and  $\text{pass}(\text{successor}[v]) \geq \text{pass}(v) - 1$  for each  $v$ .
- Following successor pointers, we must eventually repeat a node.
- Lemma 6  $\Rightarrow$  the corresponding cycle is a negative cycle. ■

**Remark.** See p. 304 for improved version and early termination rule.

(Tarjan's subtree disassembly trick)

## Dynamic programming: quiz 9



**How difficult to find a negative cycle in an undirected graph?**

- A.  $O(m \log n)$
- B.  $O(mn)$
- C.  $O(mn + n^2 \log n)$
- D.  $O(n^{2.38})$
- E. No poly-time algorithm is known.