

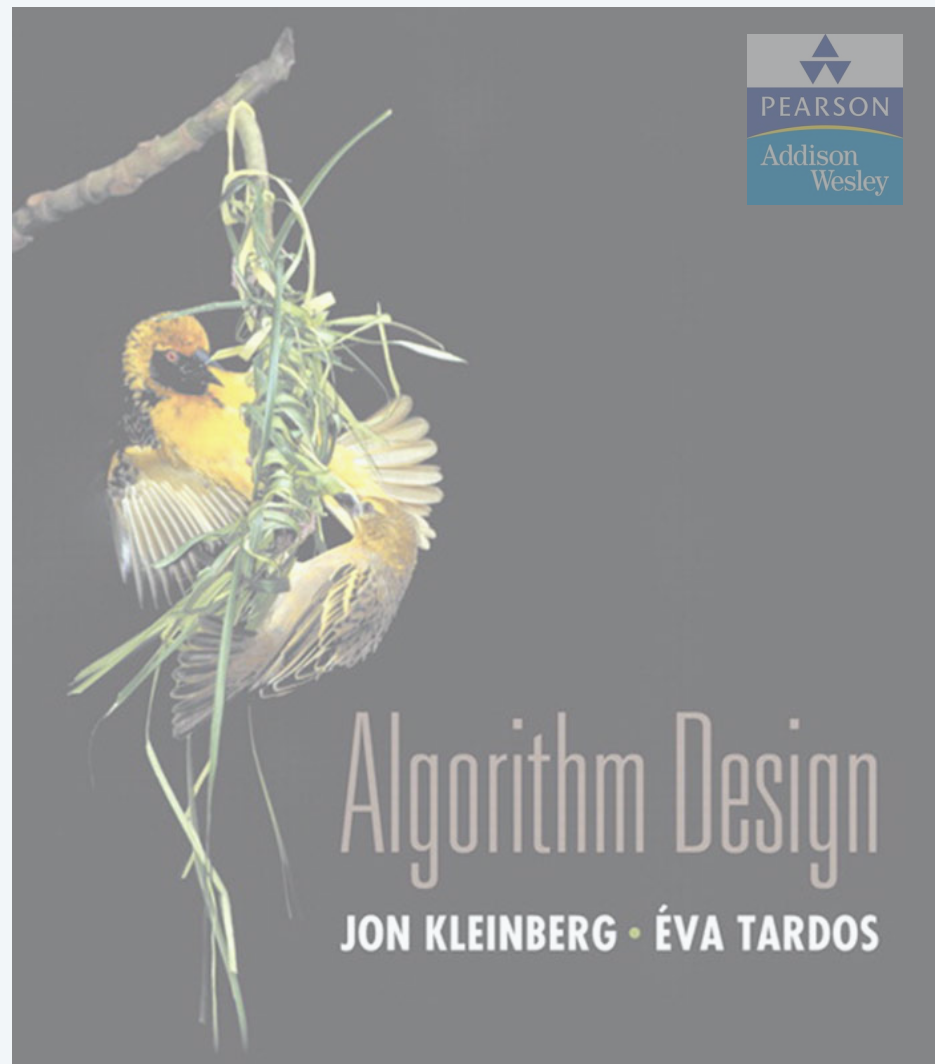
6. DYNAMIC PROGRAMMING II

- ▶ *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>



SECTION 6.6

6. DYNAMIC PROGRAMMING II

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

String similarity

Q. How similar are two strings?

Ex. occurrence and occurrence.



6 mismatches, 1 gap



1 mismatch, 1 gap



0 mismatches, 3 gaps

Edit distance

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

- Gap penalty δ ; mismatch penalty α_{pq} .
- Cost = sum of gap and mismatch penalties.

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

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

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

BLOSUM matrix for proteins

	A	R	N	D	C	Q	E	G	H	I	L	K	M	F	P	S	T	W	Y	V
A	7	-3	-3	-3	-1	-2	-2	0	-3	-3	-3	-1	-2	-4	-1	2	0	-5	-4	-1
R	-3	9	-1	-3	-6	1	-1	-4	0	-5	-4	3	-3	-5	-3	-2	-2	-5	-4	-4
N	-3	-1	9	2	-5	0	-1	-1	1	-6	-6	0	-4	-6	-4	1	0	-7	-4	-5
D	-3	-3	2	10	-7	-1	2	-3	-2	-7	-7	-2	-6	-6	-3	-1	-2	-8	-6	-6
C	-1	-6	-5	-7	13	-5	-7	-6	-7	-2	-3	-6	-3	-4	-6	-2	-2	-5	-5	-2
Q	-2	1	0	-1	-5	9	3	-4	1	-5	-4	2	-1	-5	-3	-1	-1	-4	-3	-4
E	-2	-1	-1	2	-7	3	8	-4	0	-6	-6	1	-4	-6	-2	-1	-2	-6	-5	-4
G	0	-4	-1	-3	-6	-4	-4	9	-4	-7	-7	-3	-5	-6	-5	-1	-3	-6	-6	-6
H	-3	0	1	-2	-7	1	0	-4	12	-6	-5	-1	-4	-2	-4	-2	-3	-4	3	-5
I	-3	-5	-6	-7	-2	-5	-6	-7	-6	7	2	-5	2	-1	-5	-4	-2	-5	-3	4
L	-3	-4	-6	-7	-3	-4	-6	-7	-5	2	6	-4	3	0	-5	-4	-3	-4	-2	1
K	-1	3	0	-2	-6	2	1	-3	-1	-5	-4	8	-3	-5	-2	-1	-1	-6	-4	-4
M	-2	-3	-4	-6	-3	-1	-4	-5	-4	2	3	-3	9	0	-4	-3	-1	-3	-3	1
F	-4	-5	-6	-6	-4	-5	-6	-6	-2	-1	0	-5	0	10	-6	-4	-4	0	4	-2
P	-1	-3	-4	-3	-6	-3	-2	-5	-4	-5	-5	-2	-4	-6	12	-2	-3	-7	-6	-4
S	2	-2	1	-1	-2	-1	-1	-1	-2	-4	-4	-1	-3	-4	-2	7	2	-6	-3	-3
T	0	-2	0	-2	-2	-1	-2	-3	-3	-2	-3	-1	-1	-4	-3	2	8	-5	-3	0
W	-5	-5	-7	-8	-5	-4	-6	-6	-4	-5	-4	-6	-3	0	-7	-6	-5	16	3	-5
Y	-4	-4	-4	-6	-5	-3	-5	-6	3	-3	-2	-4	-3	4	-6	-3	-3	3	11	-3
V	-1	-4	-5	-6	-2	-4	-4	-6	-5	4	1	-4	1	-2	-4	-3	0	-5	-3	7



What is edit distance between these two strings?

P A L E T T E

P A L A T E

Assume gap penalty = 2 and mismatch penalty = 1.

- A. 1
- B. 2
- C. 3
- D. 4
- E. 5

Sequence alignment

Goal. Given two strings $x_1 x_2 \dots x_m$ and $y_1 y_2 \dots y_n$, find a min-cost alignment.

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

$x_i - y_j$ and $x_{i'} - y_{j'}$ cross if $i < i'$, but $j > j'$

Def. The **cost** of an alignment M is:

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

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

an alignment of CTACCG and TACATG

$$M = \{ x_2 - y_1, x_3 - y_2, x_4 - y_3, x_5 - y_4, x_6 - y_6 \}$$

Sequence alignment: problem structure

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

Goal. $OPT(m, n)$.

Case 1. $OPT(i, j)$ matches $x_i - y_j$.

Pay mismatch for $x_i - y_j$ + min cost of aligning $x_1 x_2 \dots x_{i-1}$ and $y_1 y_2 \dots y_{j-1}$.

Case 2a. $OPT(i, j)$ leaves x_i unmatched.

Pay gap for x_i + min cost of aligning $x_1 x_2 \dots x_{i-1}$ and $y_1 y_2 \dots y_j$.

Case 2b. $OPT(i, j)$ leaves y_j unmatched.

Pay gap for y_j + min cost of aligning $x_1 x_2 \dots x_i$ and $y_1 y_2 \dots y_{j-1}$.

optimal substructure property
(proof via exchange argument)

Bellman equation.

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

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

$M[i, 0] \leftarrow i\delta.$

FOR $j = 0$ TO n

$M[0, j] \leftarrow j\delta.$

FOR $i = 1$ TO m

FOR $j = 1$ TO n

$M[i, j] \leftarrow \min \{ \alpha_{x_i y_j} + M[i-1, j-1],$
 $\delta + M[i-1, j],$
 $\delta + M[i, j-1] \}.$

already
computed

RETURN $M[m, n].$

Sequence alignment: traceback

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



1 mismatch, 1 gap

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 $\delta > 0$.

which would disprove SETH
(strong exponential time hypothesis)



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

Arturs Backurs[†]
MIT

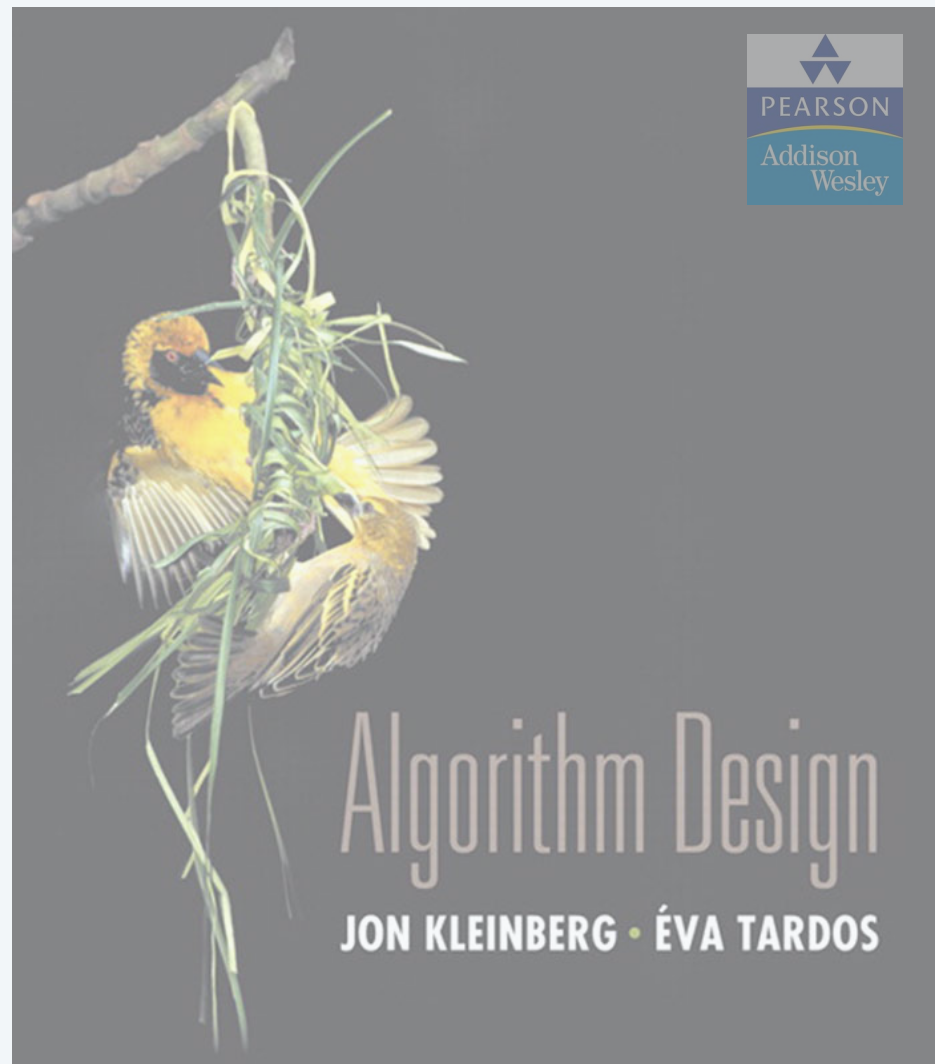
Piotr Indyk[‡]
MIT



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

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

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



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

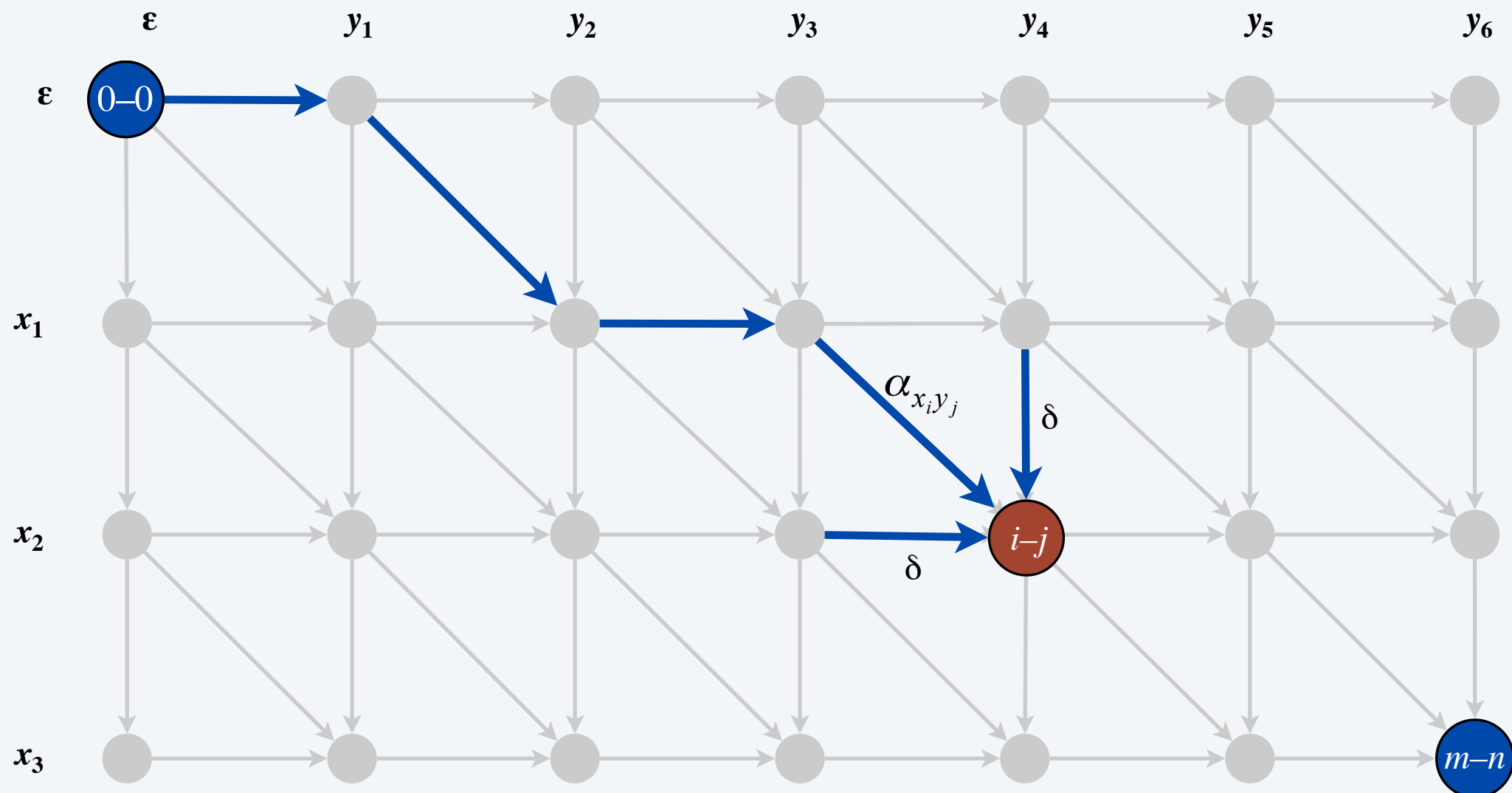
CR Categories: 3.63, 3.73, 3.79, 4.22, 5.25



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 .



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 .

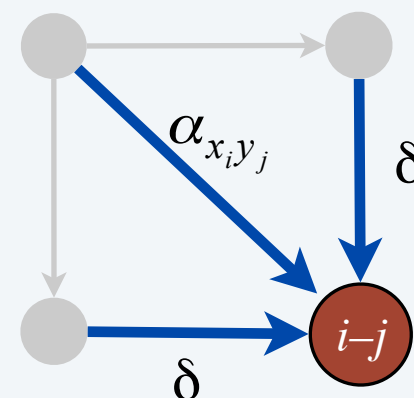
Pf of Lemma. [by strong induction on $i + j$]

- 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

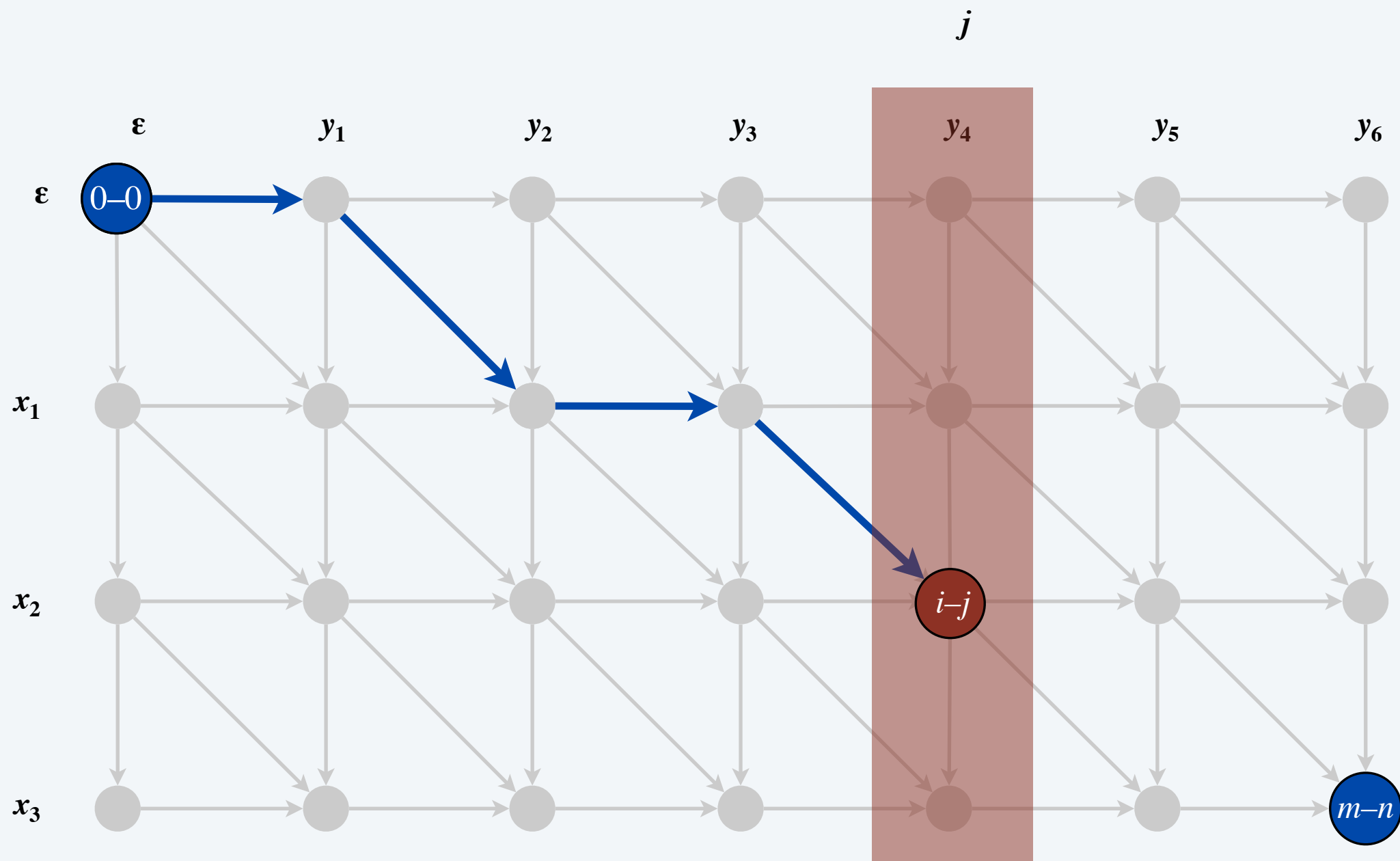
Bellman equation



Hirschberg's algorithm

Edit distance graph.

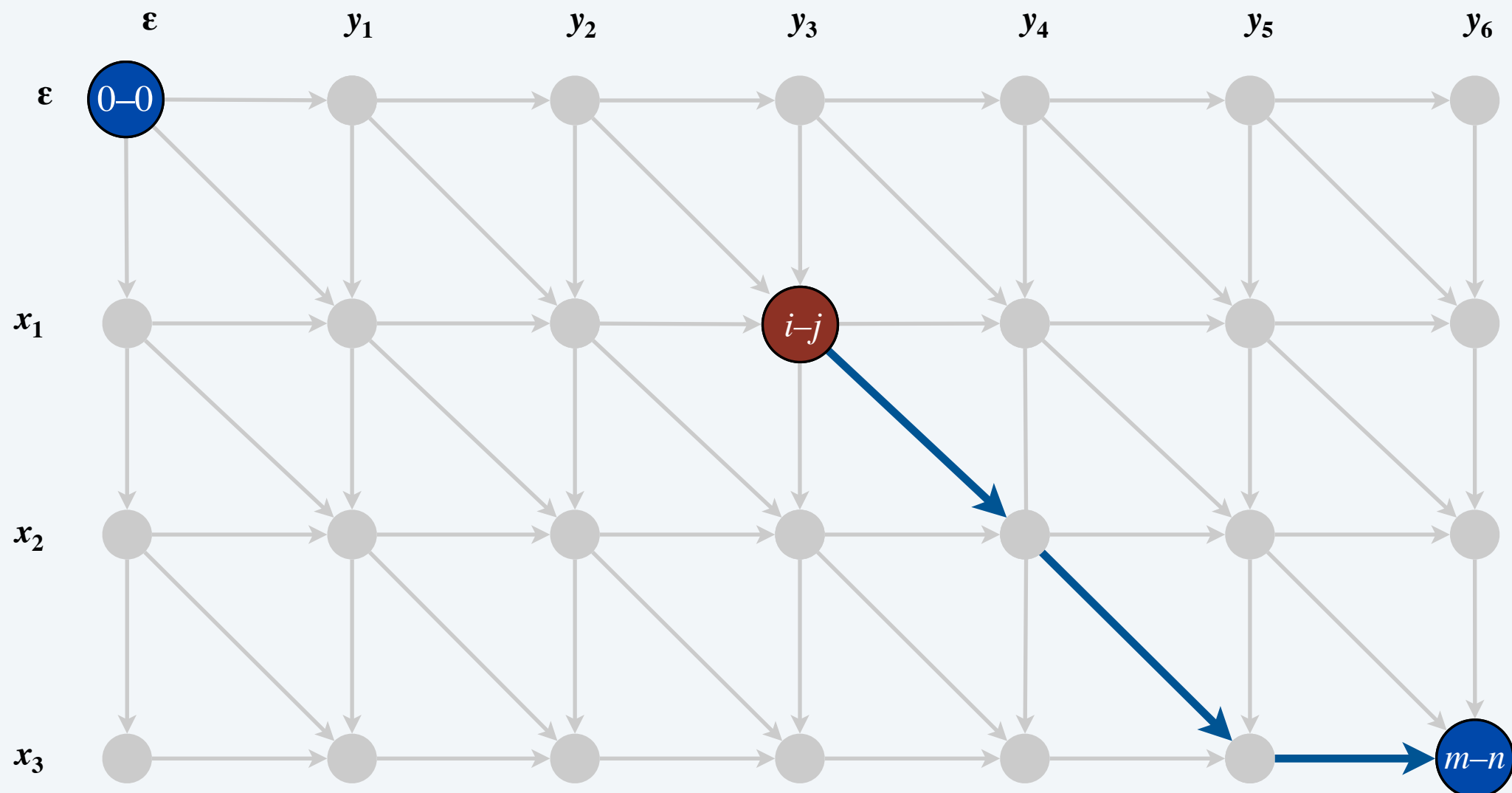
- Let $f(i, j)$ denote length of shortest path from $(0,0)$ to (i, j) .
- Lemma: $f(i, j) = OPT(i, j)$ for all i and j .
- Can compute $f(\cdot, j)$ for any j in $O(mn)$ time and $O(m + n)$ space.



Hirschberg's algorithm

Edit distance graph.

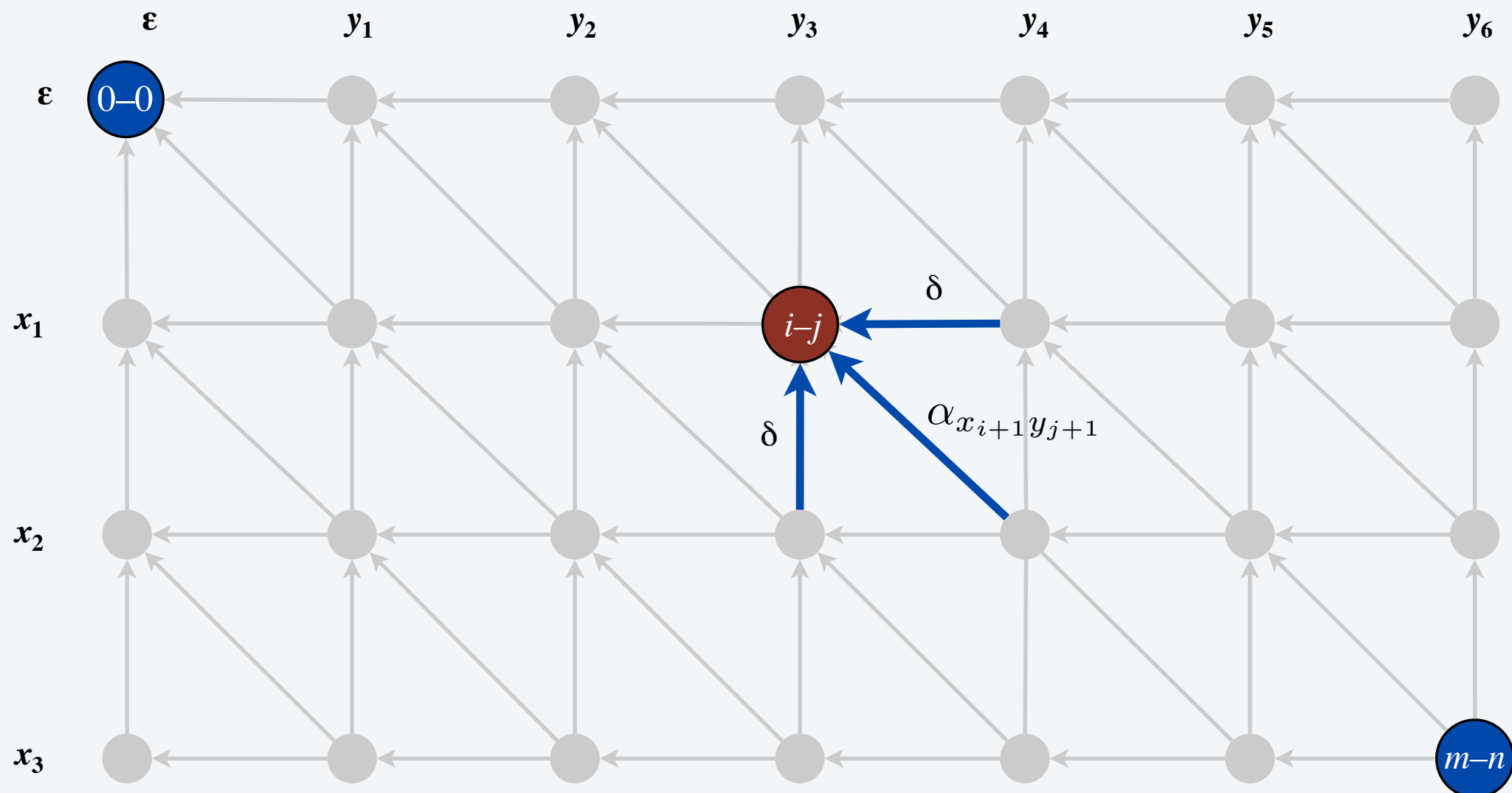
- Let $g(i, j)$ denote length of shortest path from (i, j) to (m, n) .



Hirschberg's algorithm

Edit distance graph.

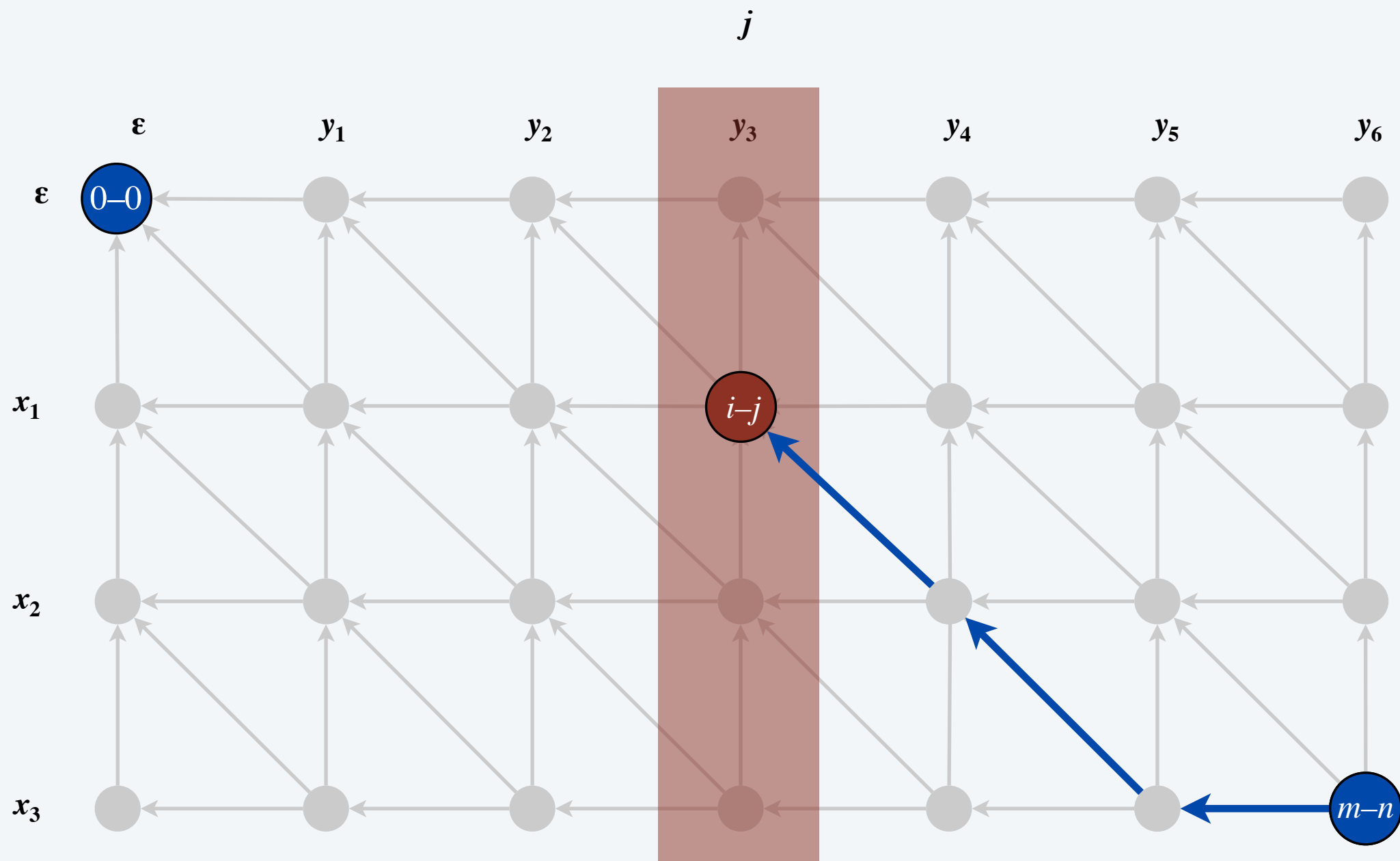
- Let $g(i, j)$ denote length of shortest path from (i, j) to (m, n) .
- Can compute $g(i, j)$ by reversing the edge orientations and inverting the roles of $(0, 0)$ and (m, n) .



Hirschberg's algorithm

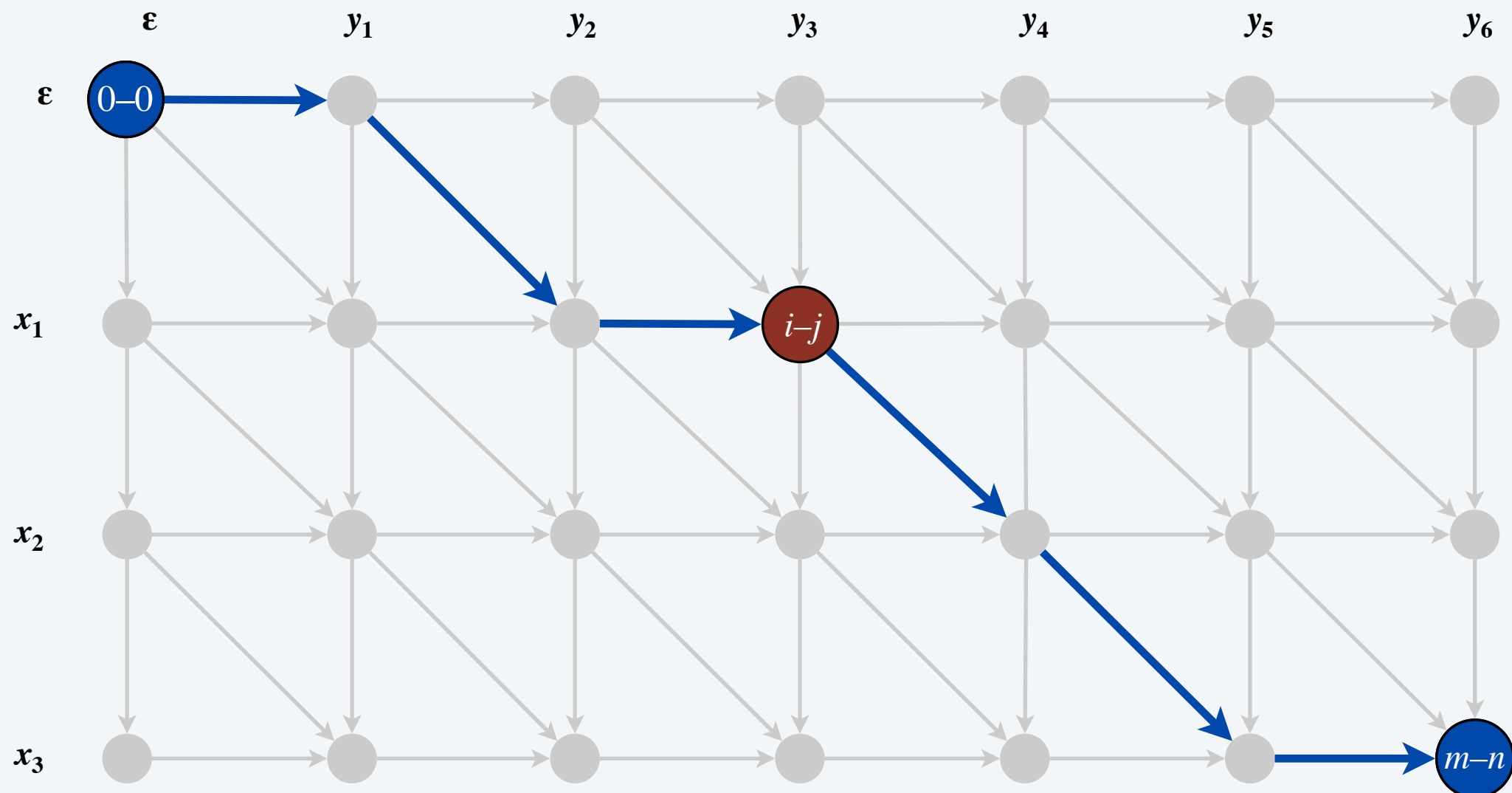
Edit distance graph.

- 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.



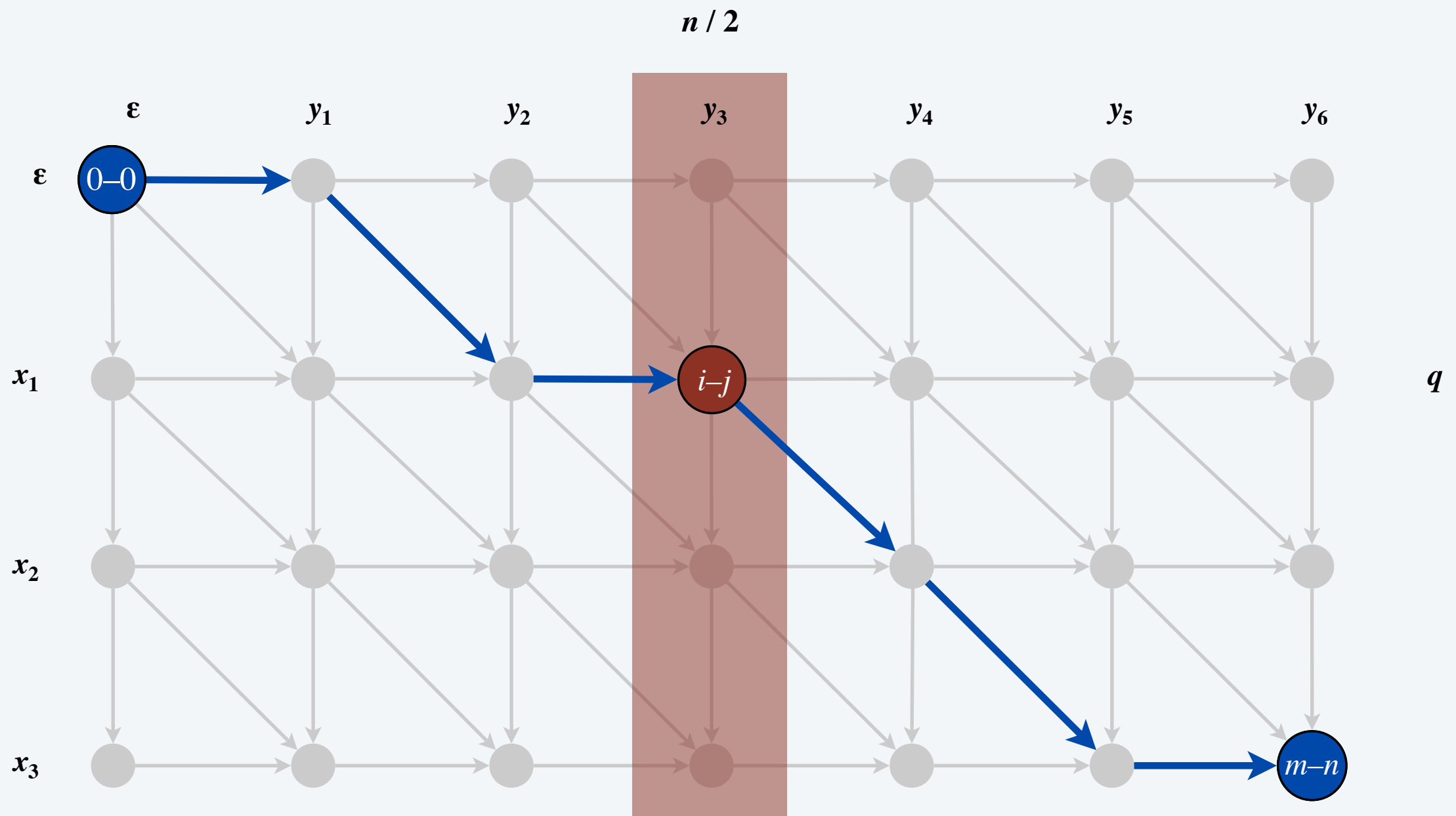
Hirschberg's algorithm

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



Hirschberg's algorithm

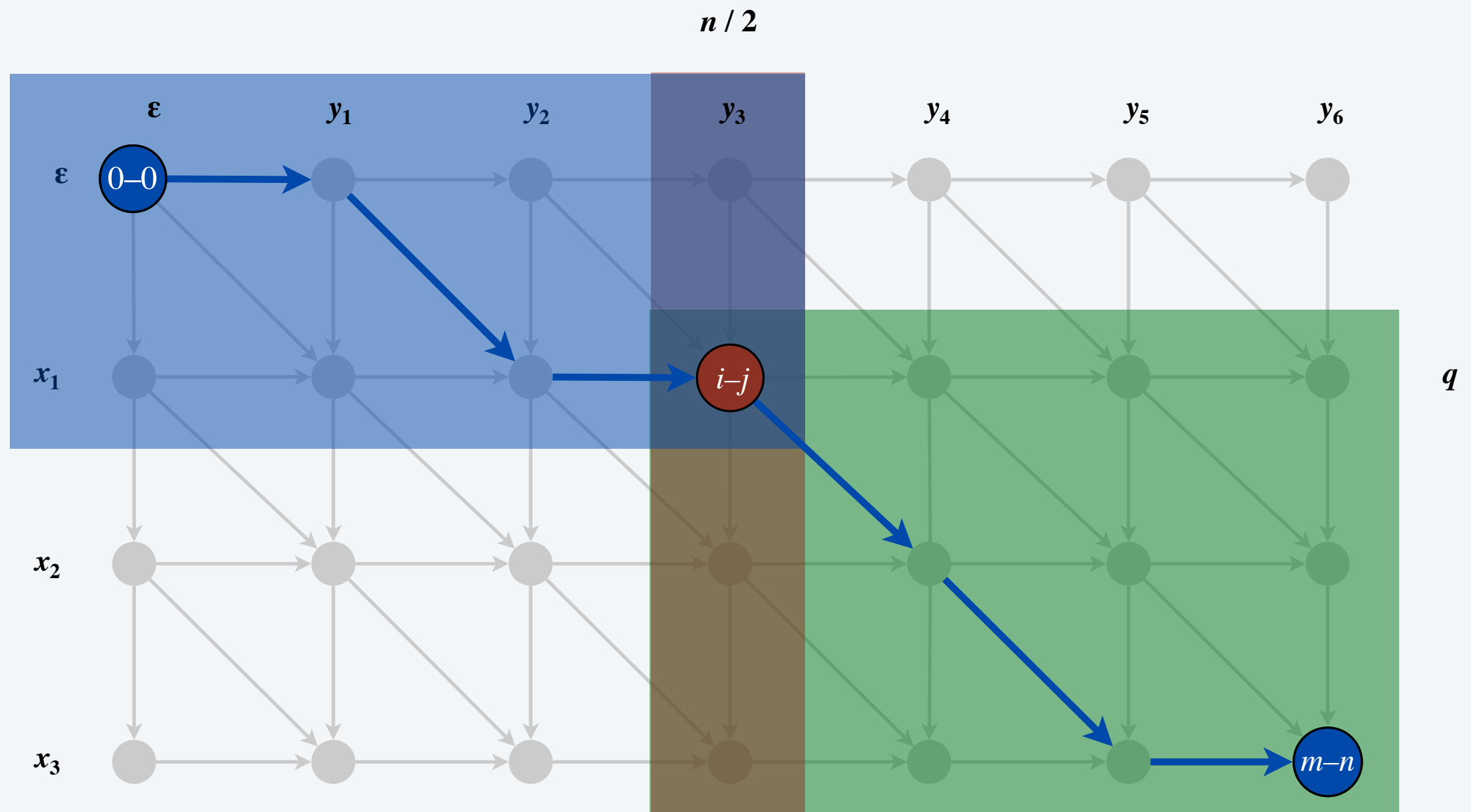
Observation 2. let q be an index that minimizes $f(q, n/2) + g(q, n/2)$.
Then, there exists a shortest path from $(0, 0)$ to (m, n) that uses $(q, n/2)$.



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.



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 $\leq n$. ■



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. $O(mn \log m \log n)$

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, $T(m, n) = O(m n \log n)$.

Pf.

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

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


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: $T(m', n') \leq 2cm'n'$ for all (m', n') with $m' + n' < m + n$.

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

inductive hypothesis 

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. $\text{LCS}(\text{GGCACACG}, \text{ACGGCGGATACG}) = \text{GGCAACG}$.

Applications. Unix diff, git, bioinformatics.

LONGEST COMMON SUBSEQUENCE



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 + length of LCS of $x_1 x_2 \dots x_{i-1}$ 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.

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

LONGEST COMMON SUBSEQUENCE



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.

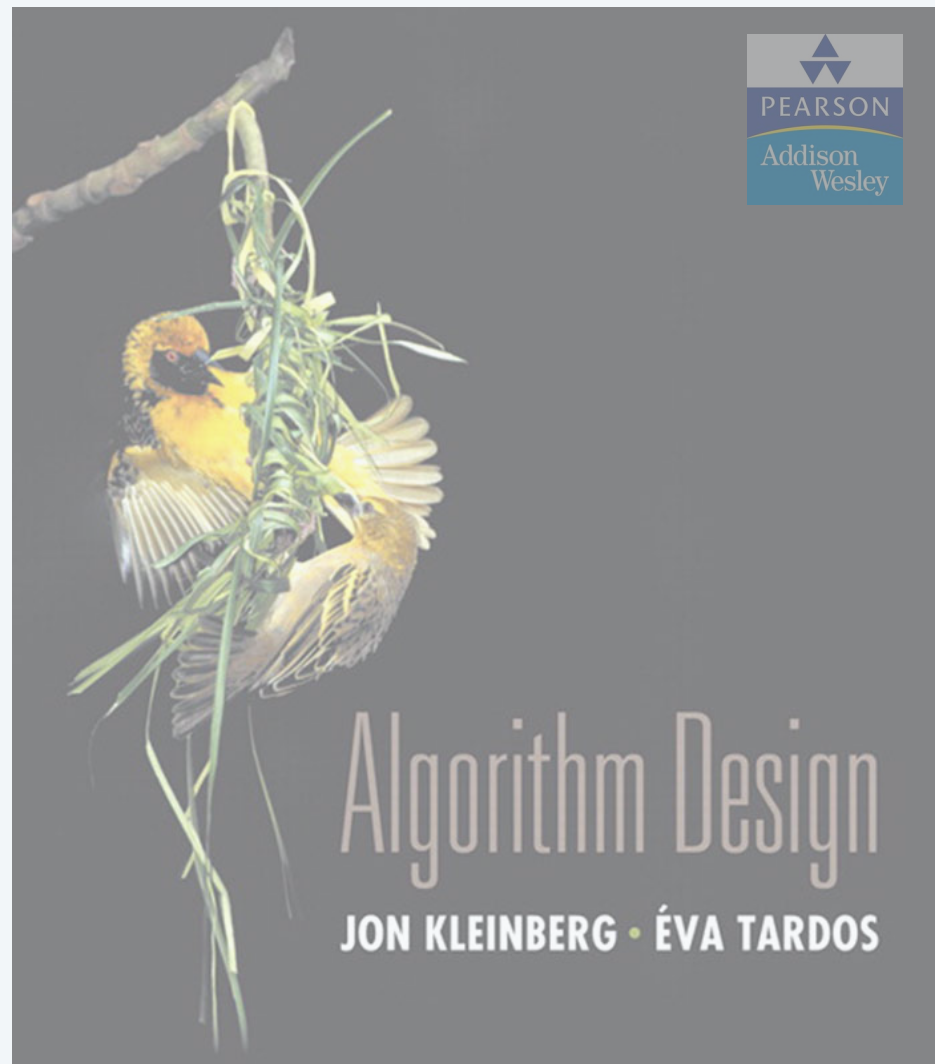
Lower bound. No $O(n^{2-\epsilon})$ algorithm unless SETH is false.

Tight Hardness Results for LCS and other
Sequence Similarity Measures

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SECTION 6.8

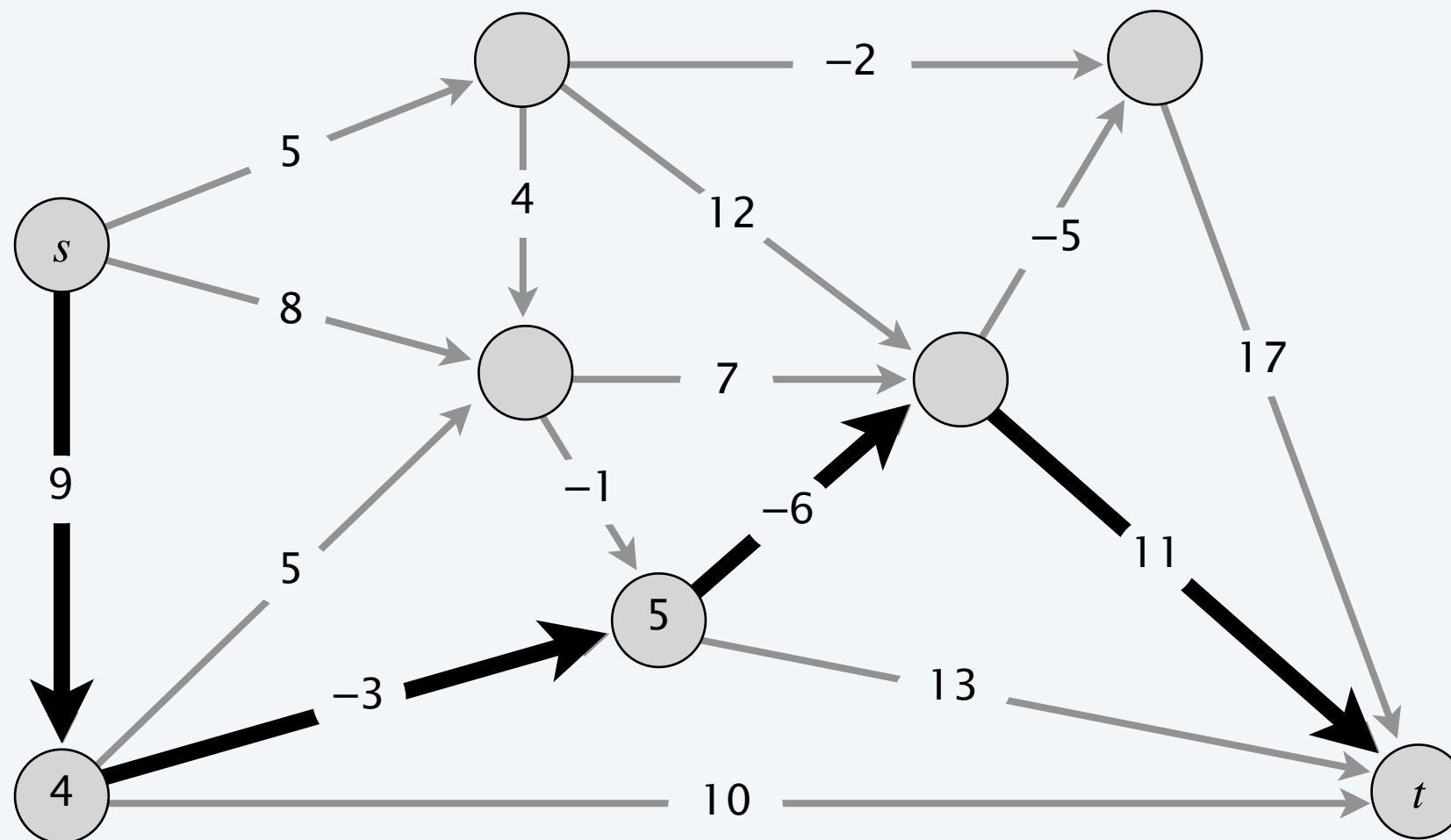
6. DYNAMIC PROGRAMMING II

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

Shortest paths with negative weights

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

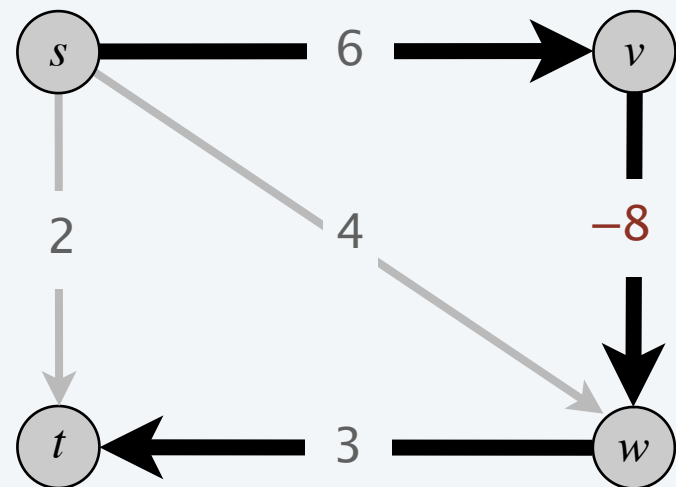
assume there exists a path from every node to t



length of shortest $s \rightsquigarrow t$ path = $9 - 3 - 6 + 11 = 11$

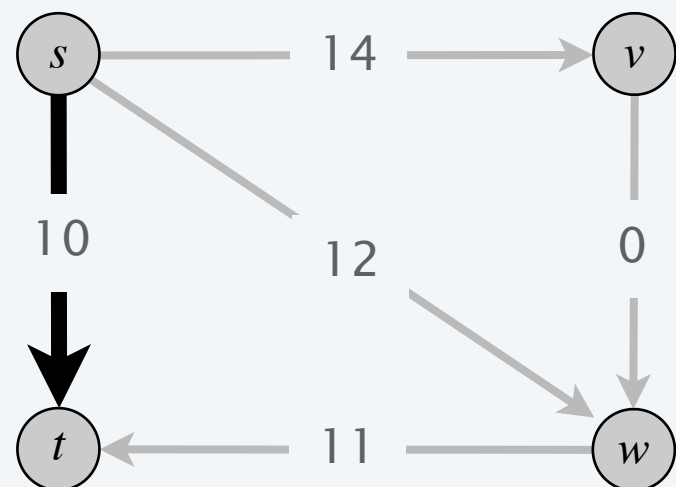
Shortest paths with negative weights: failed attempts

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 $s \rightarrow v \rightarrow w \rightarrow t$.

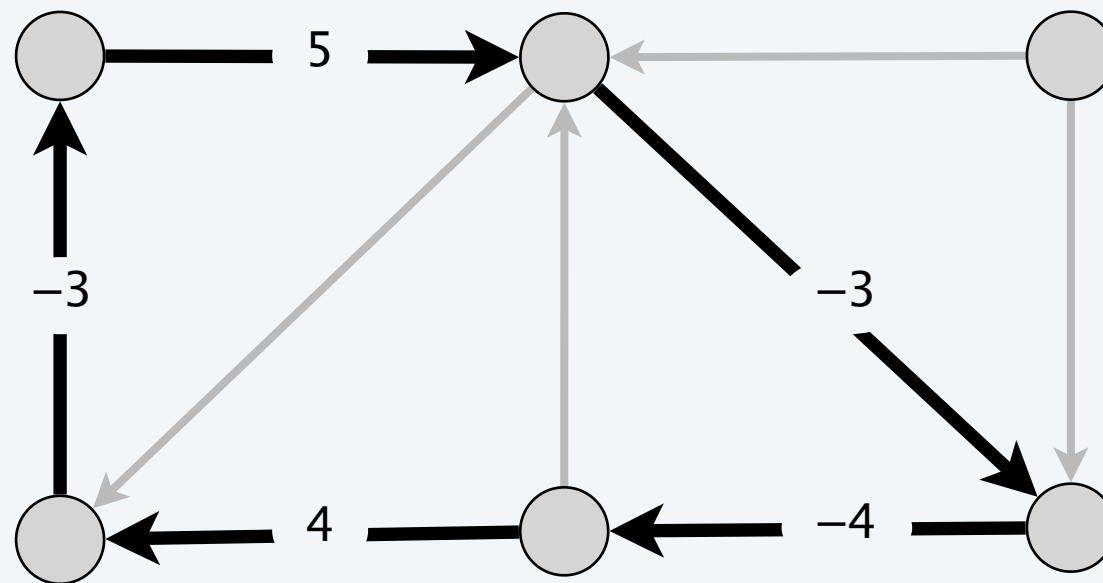
Reweighting. Adding a constant to every edge length does not necessarily make Dijkstra's algorithm produce shortest paths.



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

Negative cycles

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

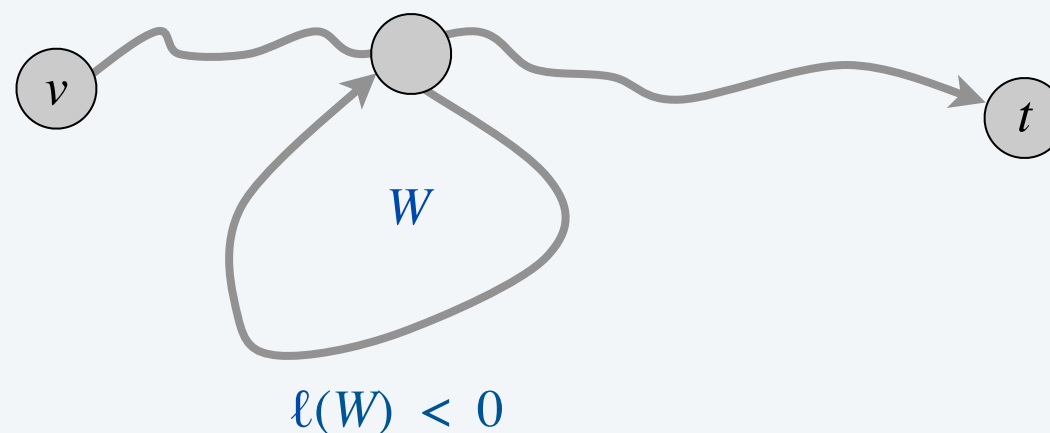


a negative cycle W : $l(W) = \sum_{e \in W} l_e < 0$

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. ■

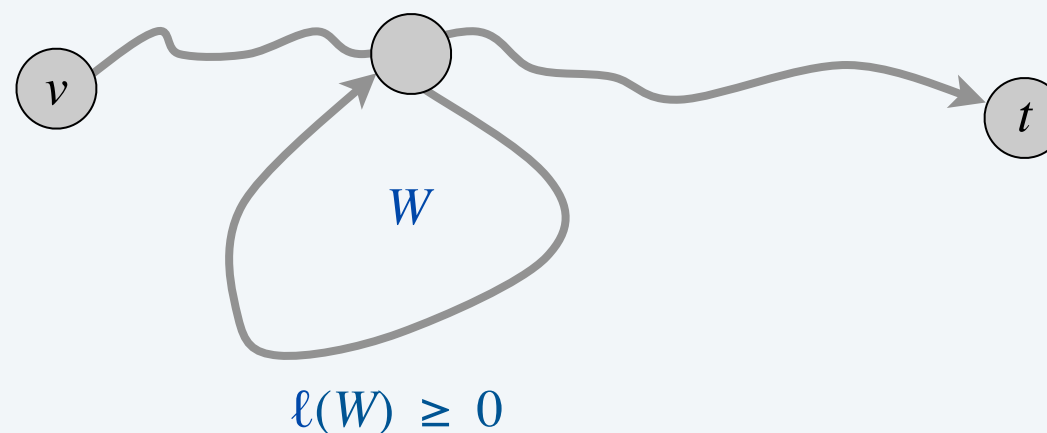


Shortest paths and negative cycles

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.

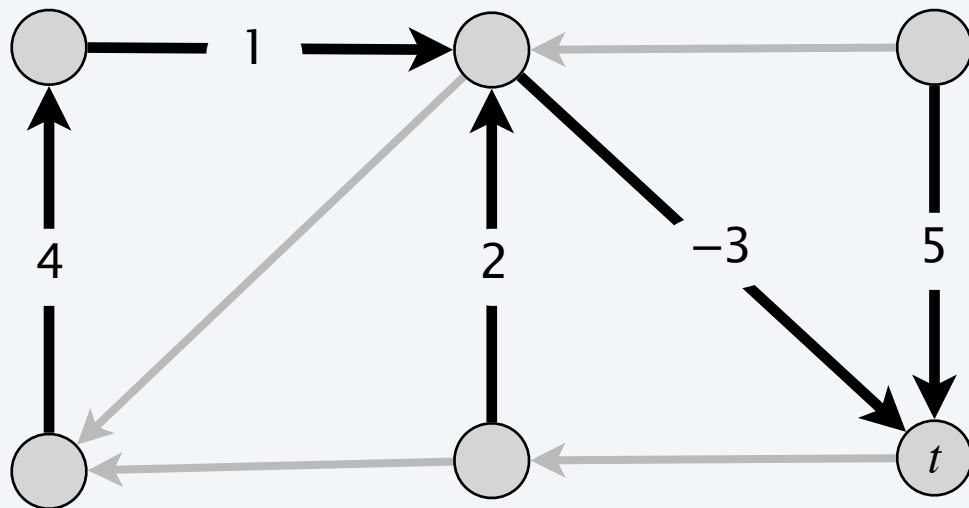
- Among all shortest $v \rightsquigarrow t$ paths, consider one that uses the fewest edges.
- If that path P contains a directed cycle W , can remove the portion of P corresponding to W without increasing its length. ■



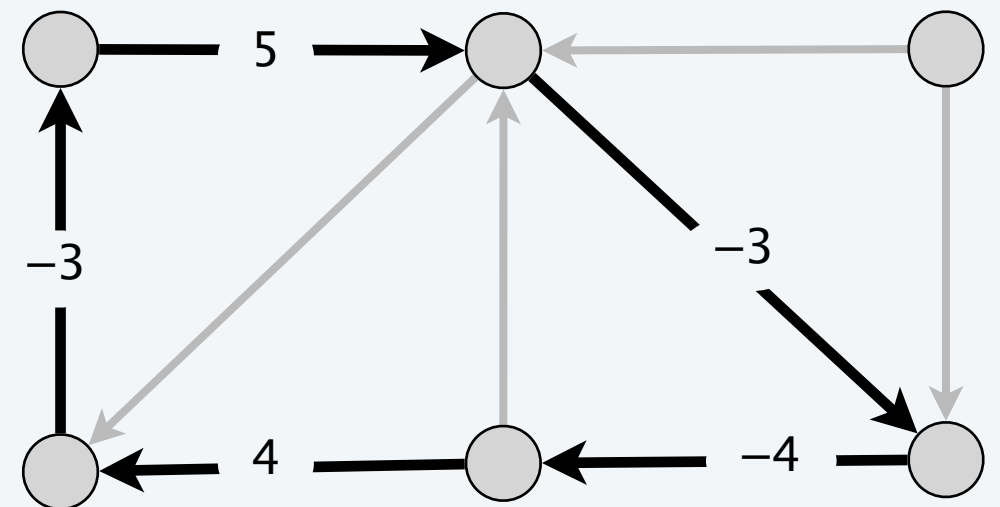
Shortest-paths and negative-cycle problems

Single-destination shortest-paths problem. Given a digraph $G = (V, E)$ with edge lengths ℓ_{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 ℓ_{vw} , find a negative cycle (if one exists).



shortest-paths tree



negative cycle



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 ℓ_{vw} .
- Then, select best $w \rightsquigarrow t$ path using $\leq i - 1$ edges.

Bellman equation.

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

Shortest paths with negative weights: implementation

SHORTEST-PATHS(V, E, ℓ, 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 $(v, w) \in E$:

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

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 $successor[i, v]$ 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.



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.

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.
- $successor[v]$ = 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 .

Bellman–Ford–Moore: efficient implementation

BELLMAN–FORD–MOORE(V, E, c, t)

FOREACH node $v \in V$:

$d[v] \leftarrow \infty$.

$successor[v] \leftarrow null$.

$d[t] \leftarrow 0$.

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 ($d[v] > d[w] + \ell_{vw}$)

$d[v] \leftarrow d[w] + \ell_{vw}$.

$successor[v] \leftarrow w$.

IF (no $d[\cdot]$ value changed in pass i) STOP.

pass i
 $O(m)$ time



Which properties must hold after pass i of Bellman–Ford–Moore?

- A.** $d[v]$ = length of a shortest $v \rightsquigarrow t$ path using $\leq i$ edges.
- B.** $d[v]$ = length of a shortest $v \rightsquigarrow t$ path using exactly i edges.
- C.** Both A and B.
- D.** Neither A nor B.

Bellman–Ford–Moore: analysis

Lemma 3. For each node v : $d[v]$ is the length of some $v \rightsquigarrow t$ path.

Lemma 4. For each node v : $d[v]$ is monotone non-increasing.

Lemma 5. After pass i , $d[v] \leq$ length of a shortest $v \rightsquigarrow t$ path using $\leq i$ edges.

Pf. [by induction on i]

- Base case: $i = 0$.
- Assume true after pass i .
- Let P be any $v \rightsquigarrow t$ path with $\leq i + 1$ edges.
- Let (v, w) be first edge in P and let P' be subpath from w to t .
- By inductive hypothesis, at the end of pass i , $d[w] \leq \ell(P')$ because P' is a $w \rightsquigarrow t$ path with $\leq i$ edges.
- After considering edge (v, w) in pass $i + 1$:

and by Lemma 4,
 $d[w]$ does not increase

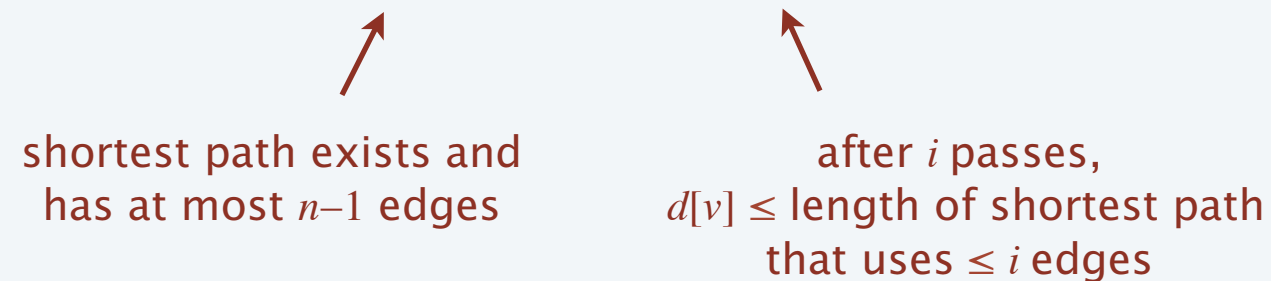
$$\begin{aligned} d[v] &\leq \ell_{vw} + d[w] \\ &\leq \ell_{vw} + \ell(P') \\ &= \ell(P) \quad \blacksquare \end{aligned}$$

and by Lemma 4,
 $d[v]$ does not increase

Bellman–Ford–Moore: analysis

Theorem 2. Assuming no negative cycles, Bellman–Ford–Moore computes the lengths of the shortest $v \rightsquigarrow t$ paths in $O(mn)$ time and $\Theta(n)$ extra space.

Pf. Lemma 2 + Lemma 5. ■



Remark. Bellman–Ford–Moore is typically faster in practice.

- Edge (v, w) considered in pass $i + 1$ only if $d[w]$ updated in pass i .
- If shortest path has k edges, then algorithm finds it after $\leq k$ passes.



Assuming no negative cycles, which properties must hold throughout Bellman–Ford–Moore?

- A. Following $successor[v]$ pointers gives a directed $v \rightsquigarrow t$ path.
- B. If following $successor[v]$ pointers gives a directed $v \rightsquigarrow t$ path, then the length of that $v \rightsquigarrow t$ path is $d[v]$.
- C. Both A and B.
- D. Neither A nor B.

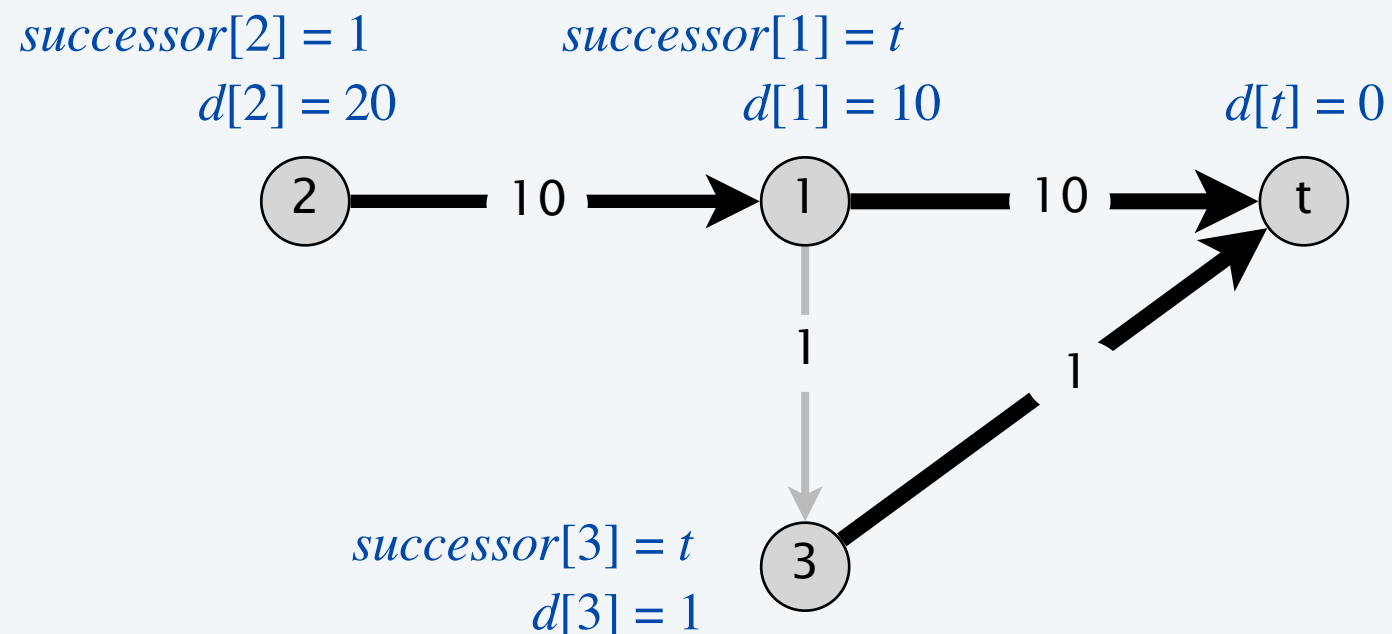
Bellman-Ford-Moore: analysis

Claim. ~~Throughout Bellman-Ford-Moore, following the $successor[v]$ pointers gives a directed path from v to t of length $d[v]$.~~

Counterexample. Claim is false!

- Length of successor $v \rightsquigarrow t$ path may be strictly shorter than $d[v]$.

consider nodes in order: $t, 1, 2, 3$



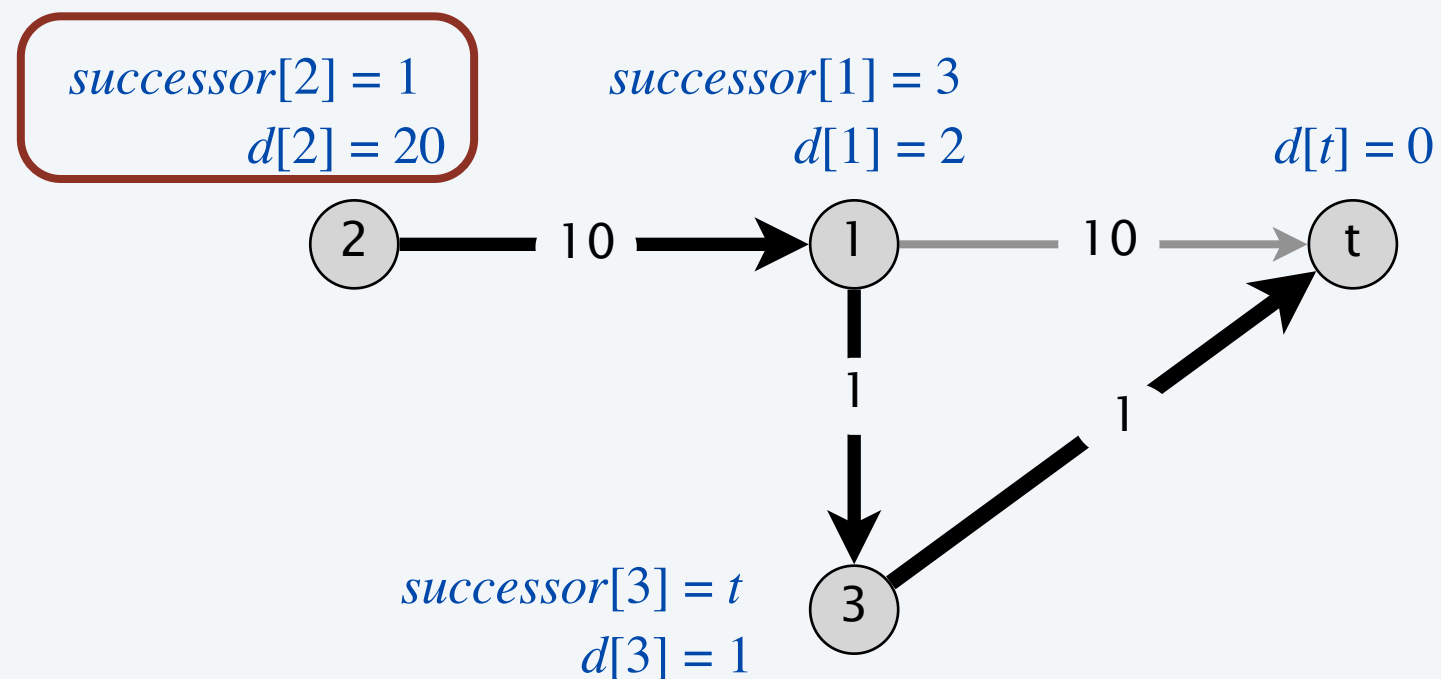
Bellman-Ford-Moore: analysis

Claim. ~~Throughout Bellman-Ford-Moore, following the $successor[v]$ pointers gives a directed path from v to t of length $d[v]$.~~

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consider nodes in order: $t, 1, 2, 3$



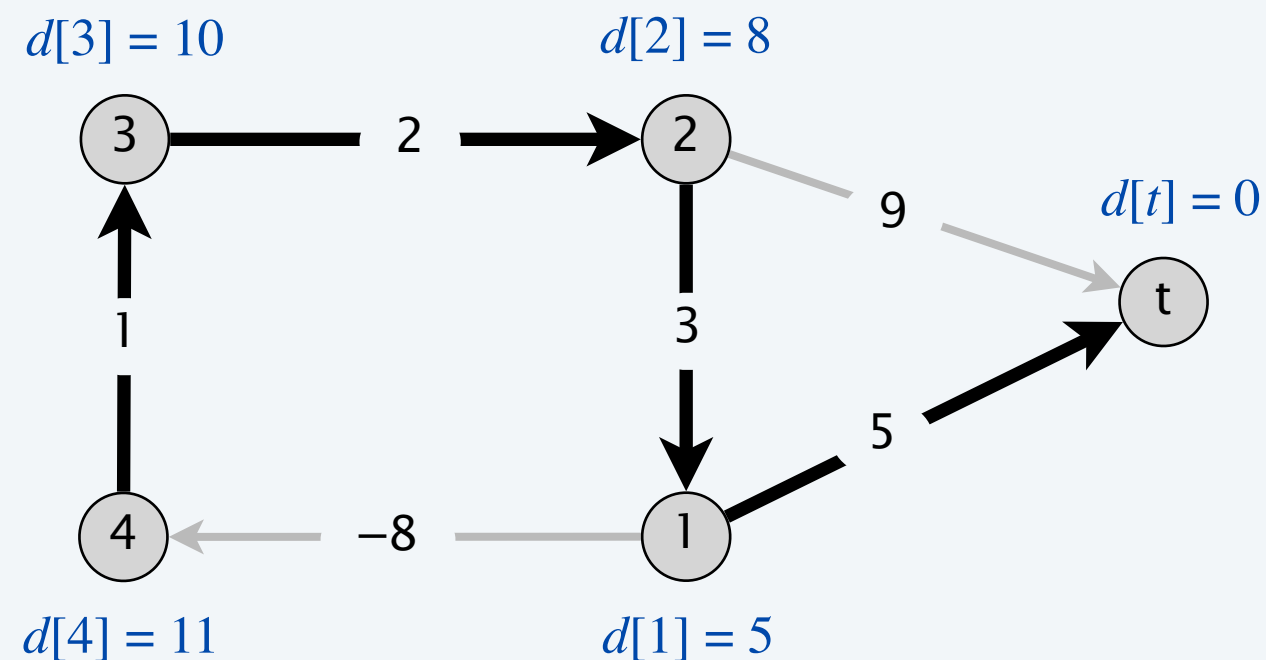
Bellman-Ford-Moore: analysis

Claim. ~~Throughout Bellman-Ford-Moore, following the $successor[v]$ pointers gives a directed path from v to t of length $d[v]$.~~

Counterexample. Claim is false!

- Length of successor $v \rightsquigarrow t$ path may be strictly shorter than $d[v]$.
- If negative cycle, successor graph may have directed cycles.

consider nodes in order: $t, 1, 2, 3, 4$



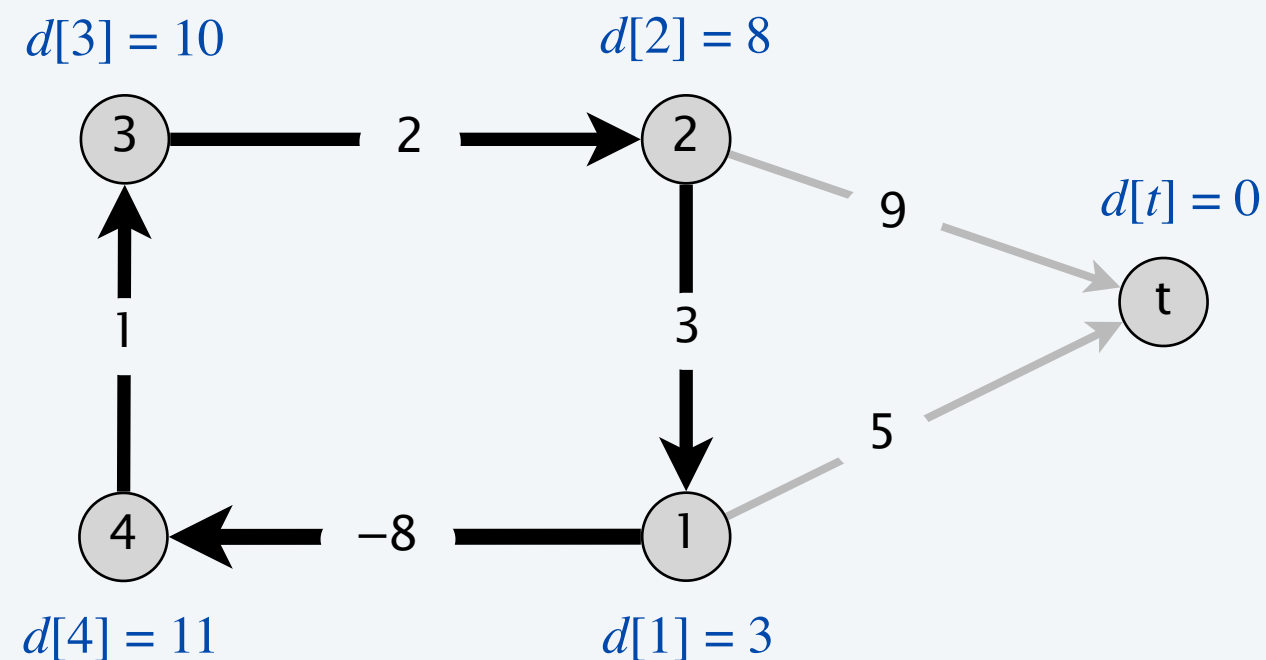
Bellman-Ford-Moore: analysis

Claim. ~~Throughout Bellman-Ford-Moore, following the $successor[v]$ pointers gives a directed path from v to t of length $d[v]$.~~

Counterexample. Claim is false!

- Length of successor $v \rightsquigarrow t$ path may be strictly shorter than $d[v]$.
- If negative cycle, successor graph may have directed cycles.

consider nodes in order: $t, 1, 2, 3, 4$



Bellman–Ford–Moore: finding the shortest paths

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

- If $successor[v] = w$, we must have $d[v] \geq d[w] + \ell_{vw}$.
(LHS and RHS are equal when $successor[v]$ is set; $d[w]$ can only decrease; $d[v]$ decreases only when $successor[v]$ 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:
$$\begin{array}{rcll} d[v_1] & \geq & d[v_2] & + \ell(v_1, v_2) \\ d[v_2] & \geq & d[v_3] & + \ell(v_2, v_3) \\ \vdots & & \vdots & \\ d[v_{k-1}] & \geq & d[v_k] & + \ell(v_{k-1}, v_k) \\ d[v_k] & > & d[v_1] & + \ell(v_k, v_1) \end{array}$$

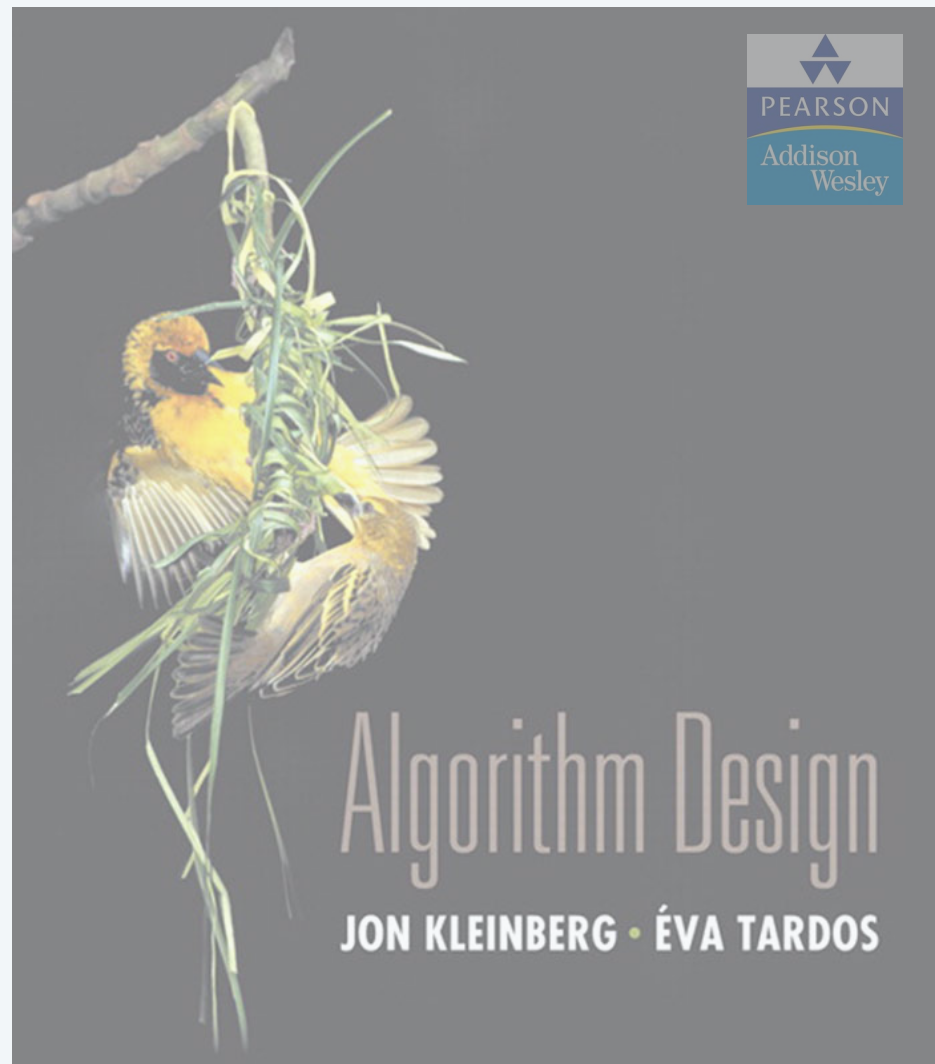
← holds with strict inequality
since we are updating $d[v_k]$
- Adding inequalities yields $\ell(v_1, v_2) + \ell(v_2, v_3) + \dots + \ell(v_{k-1}, v_k) + \ell(v_k, v_1) < 0$. ■

|—————|
 W is a negative cycle

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–Vladu
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*

Distance-vector routing protocols

Communication network.

- Node \approx router.
- Edge \approx direct communication link.
- Length of edge \approx latency of link.  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.

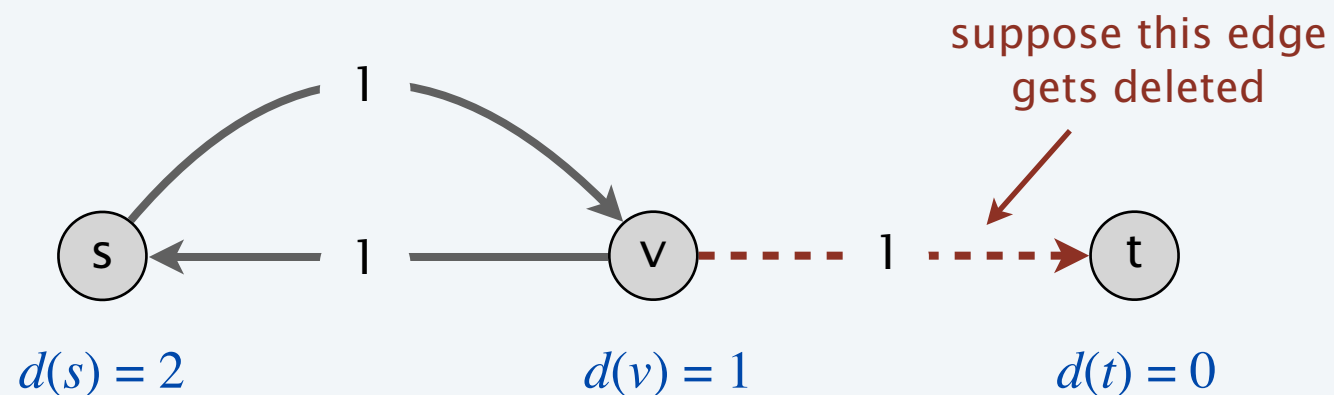
Distance-vector routing protocols

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).




“counting to infinity”

Path-vector routing protocols

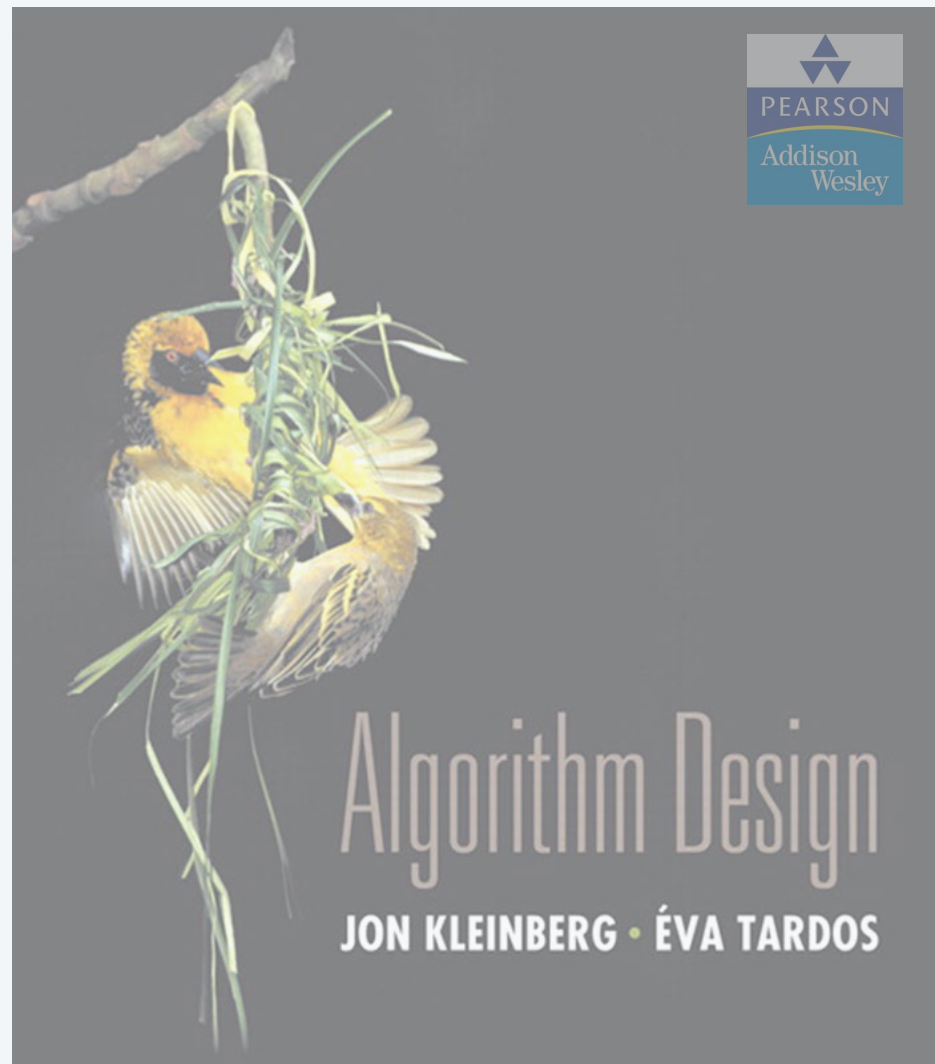
Link-state routing protocols.

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).



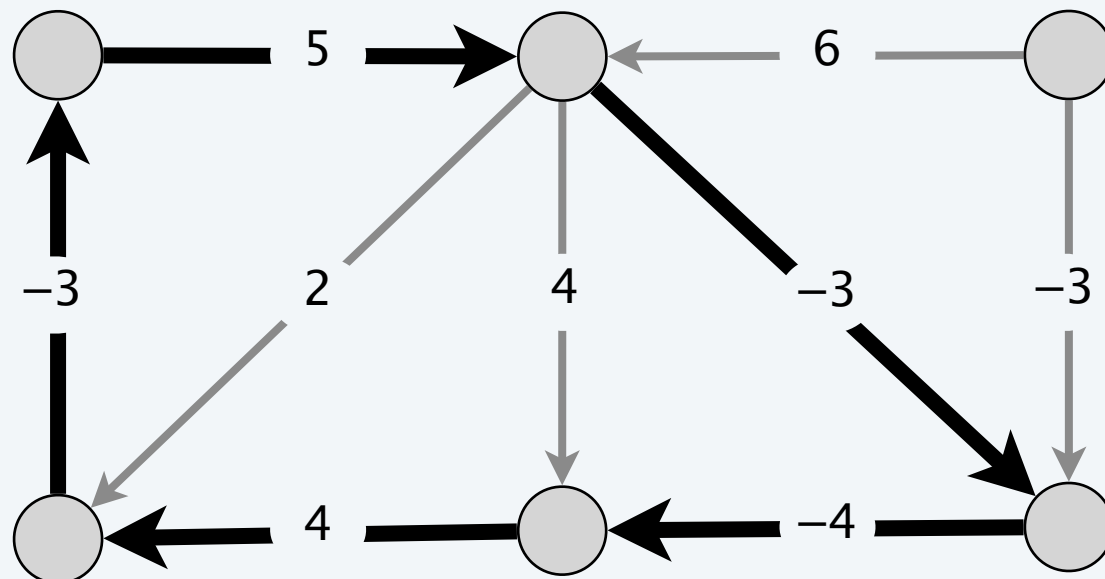
SECTION 6.10

6. DYNAMIC PROGRAMMING II

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

Detecting negative cycles

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

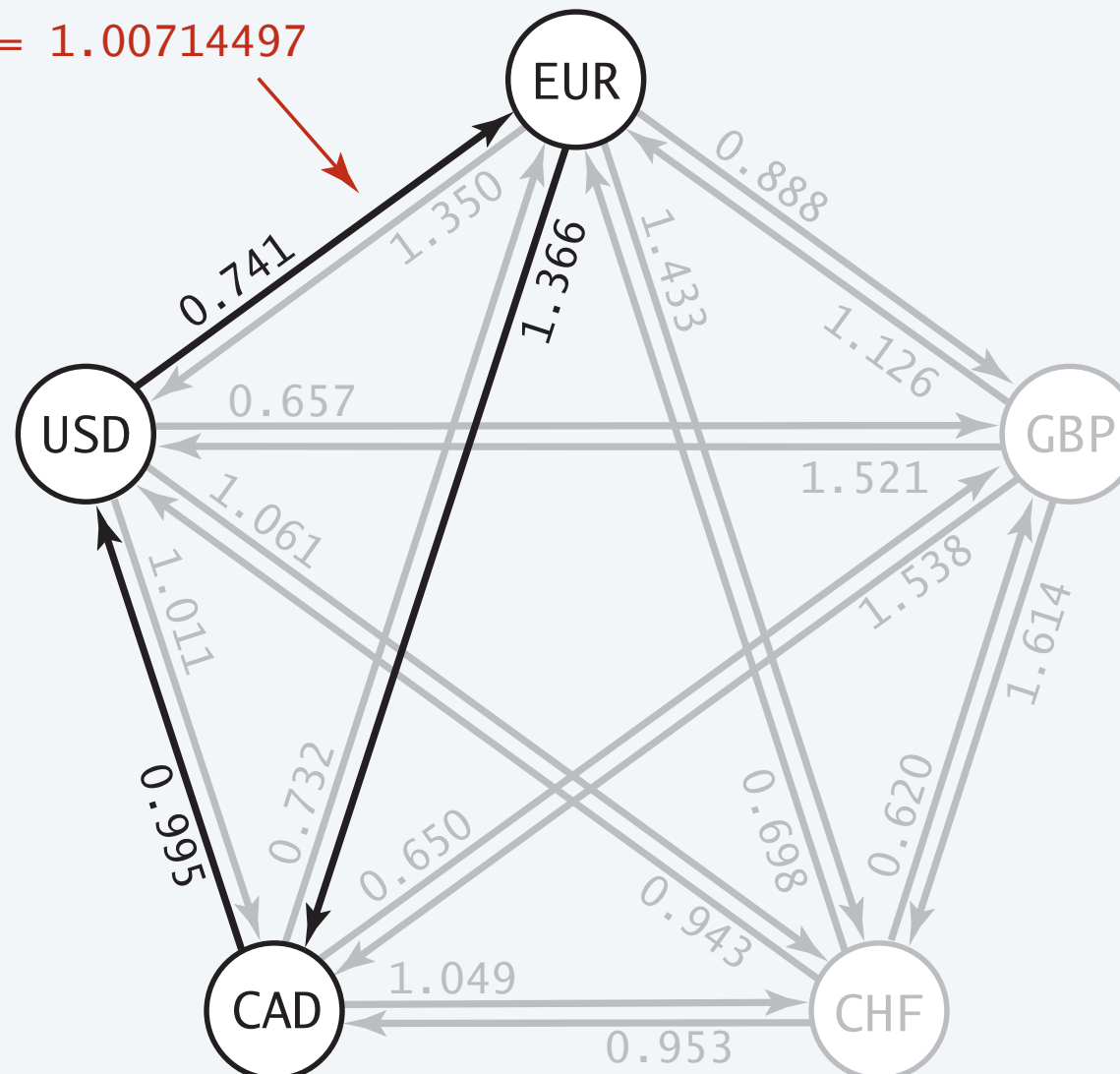


Detecting negative cycles: application

Currency conversion. Given n currencies and exchange rates between pairs of currencies, is there an arbitrage opportunity?

Remark. Fastest algorithm very valuable!

$$0.741 * 1.366 * .995 = 1.00714497$$



Detecting negative cycles

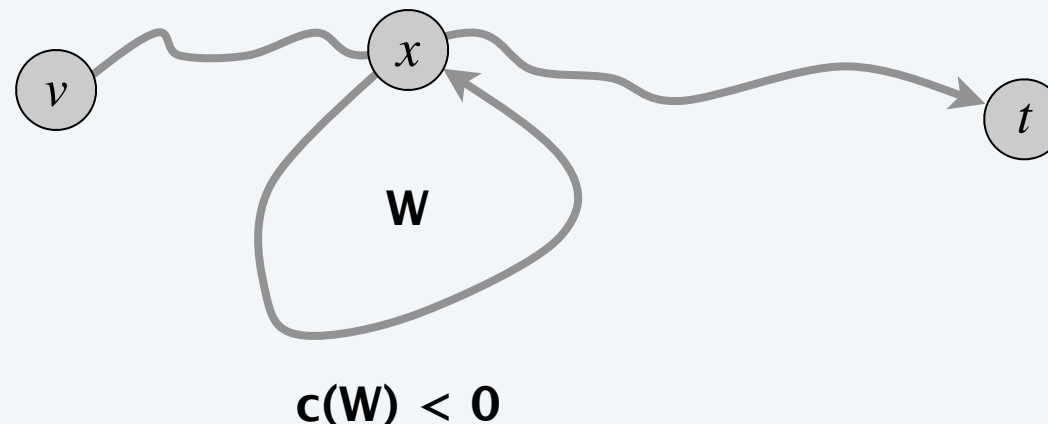
Lemma 7. If $OPT(n, v) = OPT(n - 1, v)$ for every node v , then no negative cycles.

Pf. The $OPT(n, v)$ values have converged \Rightarrow shortest $v \rightsquigarrow t$ path exists. ■

Lemma 8. If $OPT(n, v) < OPT(n - 1, v)$ for some node v , then (any) shortest $v \rightsquigarrow t$ path of length $\leq n$ contains a cycle W . Moreover W is a negative cycle.

Pf. [by contradiction]

- Since $OPT(n, v) < OPT(n - 1, v)$, we know that shortest $v \rightsquigarrow t$ path P has exactly n edges.
- By pigeonhole principle, the path P must contain a repeated node x .
- Let W be any cycle in P .
- Deleting W yields a $v \rightsquigarrow t$ path with $< n$ edges $\Rightarrow W$ is a negative cycle. ■



Detecting negative cycles

Theorem 4. Can find a negative cycle in $\Theta(mn)$ time and $\Theta(n^2)$ space.

Pf.

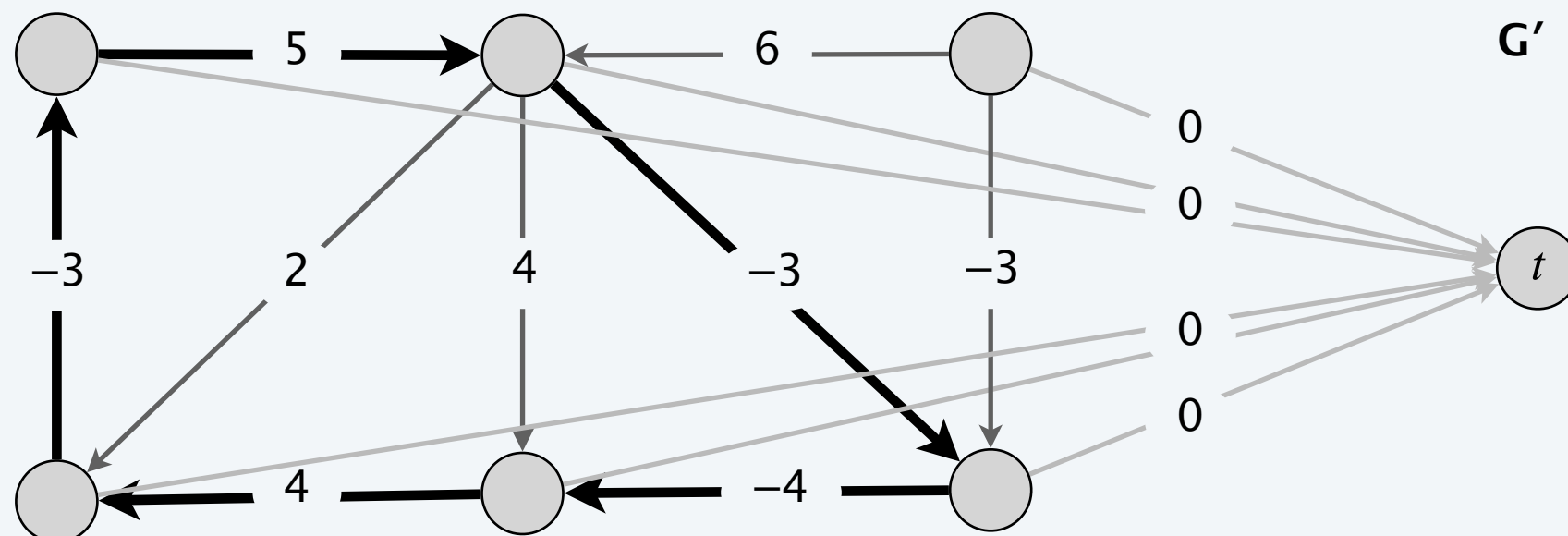
- Add new sink node t and connect all nodes to t with 0-length edge.
- G has a negative cycle iff G' has a negative cycle.
- Case 1. [$OPT(n, v) = OPT(n - 1, v)$ for every node v]

By Lemma 7, no negative cycles.

- Case 2. [$OPT(n, v) < OPT(n - 1, v)$ for some node v]

Using proof of Lemma 8, can extract negative cycle from $v \rightsquigarrow t$ path.

(cycle cannot contain t since no edge leaves t) ■



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 $pass(v) =$ last pass in which $d[v]$ was updated.
- Observe $pass(s) = n'$ and $pass(successor[v]) \geq pass(v) - 1$ 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)



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.