

The background features a complex network graph. It consists of numerous circular nodes, some of which are semi-transparent green and others are semi-transparent yellow-green. These nodes are interconnected by a dense web of thin, yellow-green lines representing edges. The overall aesthetic is technical and digital, with a dark, almost black, background that makes the glowing nodes and edges stand out.

# Shortest Path Algorithms



# DIJKSTRA'S ALGORITHM

By Laksman Veeravagu and Luis Barrera

# THE AUTHOR: EDSGER WYBE DIJKSTRA

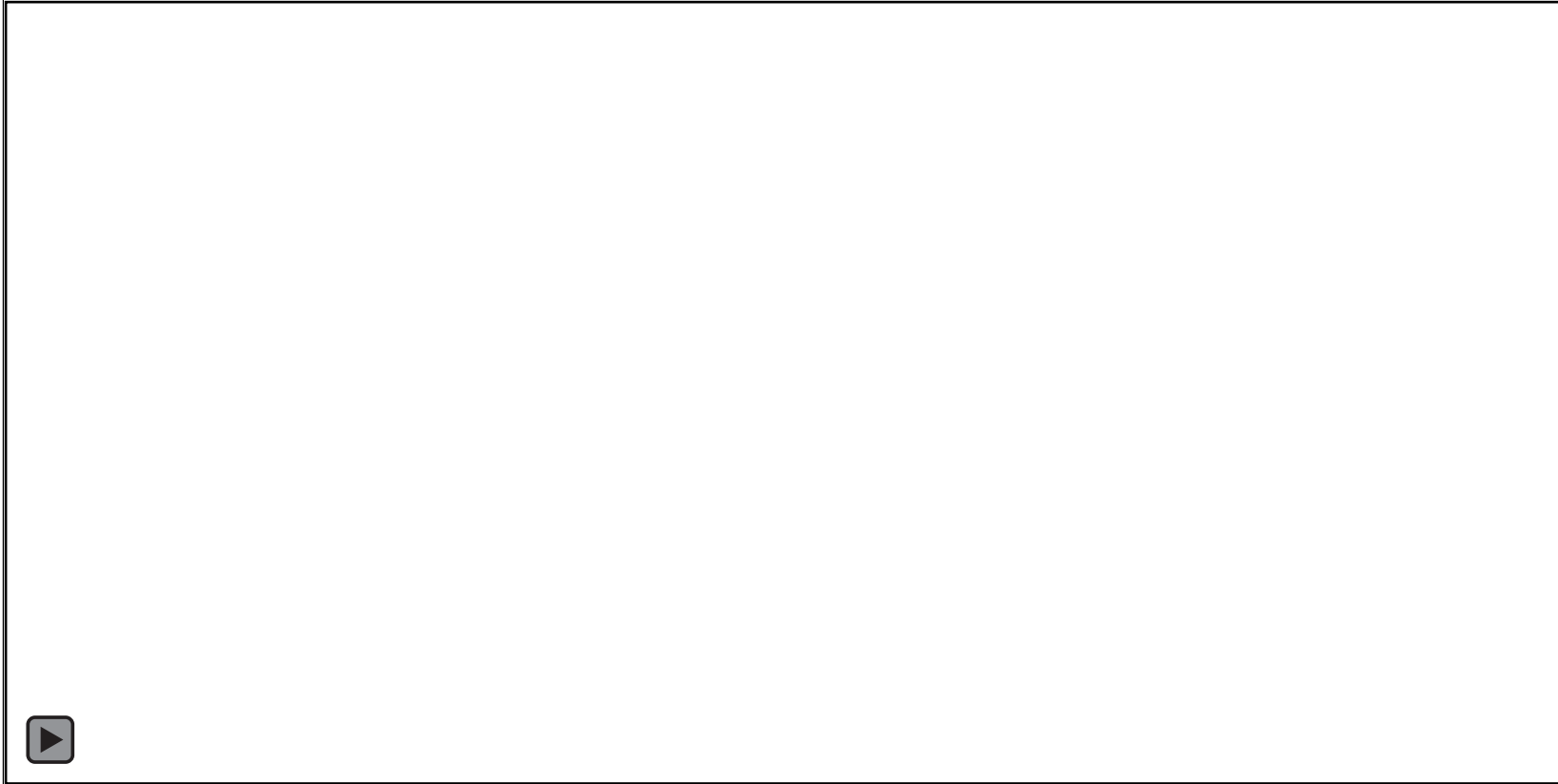


"Computer Science is no more about computers than astronomy is about telescopes."

<http://www.cs.utexas.edu/~EWD/>



# A\* VISUALIZATION



Visualization of A\*

*Vis Credit : <https://qiao.github.io/PathFinding.js/visual/>*



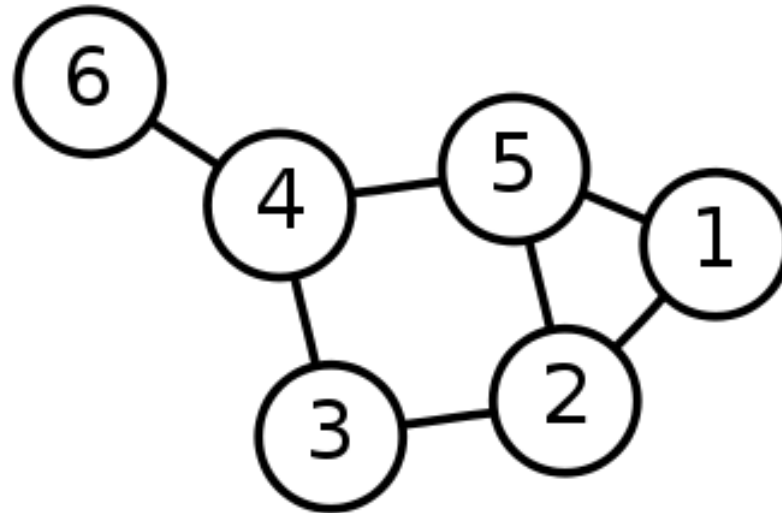
# EDSGER WYBE DIJKSTRA

- May 11, 1930 – August 6, 2002
- Received the 1972 A. M. Turing Award, widely considered the most prestigious award in computer science.
- The Schlumberger Centennial Chair of Computer Sciences at The University of Texas at Austin from 1984 until 2000
- Made a strong case against use of the GOTO statement in programming languages and helped lead to its deprecation.
- Known for his many essays on programming.



# SINGLE-SOURCE SHORTEST PATH PROBLEM

**Single-Source Shortest Path Problem** - The problem of finding shortest paths from a source vertex  $v$  to all other vertices in the graph.



# PRELIMINARY CONCEPTS

- Shortest Path (If exists)

$$\text{Shortest Path} = \min\{w(\pi) \mid \text{paths } \pi \text{ } s \rightarrow t\}$$

- DAG Relaxation

This maintains an upper estimate  $d(s, v)$  estimates upperbound and then gradually lowered until equal to  $\delta(s, v)$

- If  $d(s, v) > d(s, u) + w(s, v)$ , then “relax” by changing  $d(s, v)$  to  $d(s, u) + w(s, v)$

- Triangle Inequality

If  $\delta(u, v)$  is the shortest path length between  $u$  and  $v$ ,

$$\delta(u, v) \leq \delta(u, x) + \delta(x, v)$$



# DIJKSTRA'S ALGORITHM

Dijkstra's algorithm - is a solution to the single-source shortest path problem in graph theory.

Works on both directed and undirected graphs. However, all edges must have nonnegative weights.

**Approach:** Greedy

**Input:** Weighted graph  $G=\{E,V\}$  and source vertex  $v\in V$ , such that all edge weights are nonnegative

**Output:** Lengths of shortest paths (or the shortest paths themselves) from a given source vertex  $v\in V$  to all other vertices

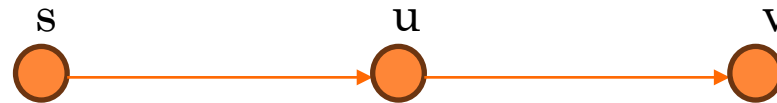




# DIJKSTRA

- If  $w \geq 0$ , then distance increases along shortest path

$$\delta(s, u) \leq \delta(s, v)$$



- Relax edges from vertices in increasing order of distance from s
- Find next vertex efficiently using a Data Structure
  - We will use a priority queue over here (priority is the d value)



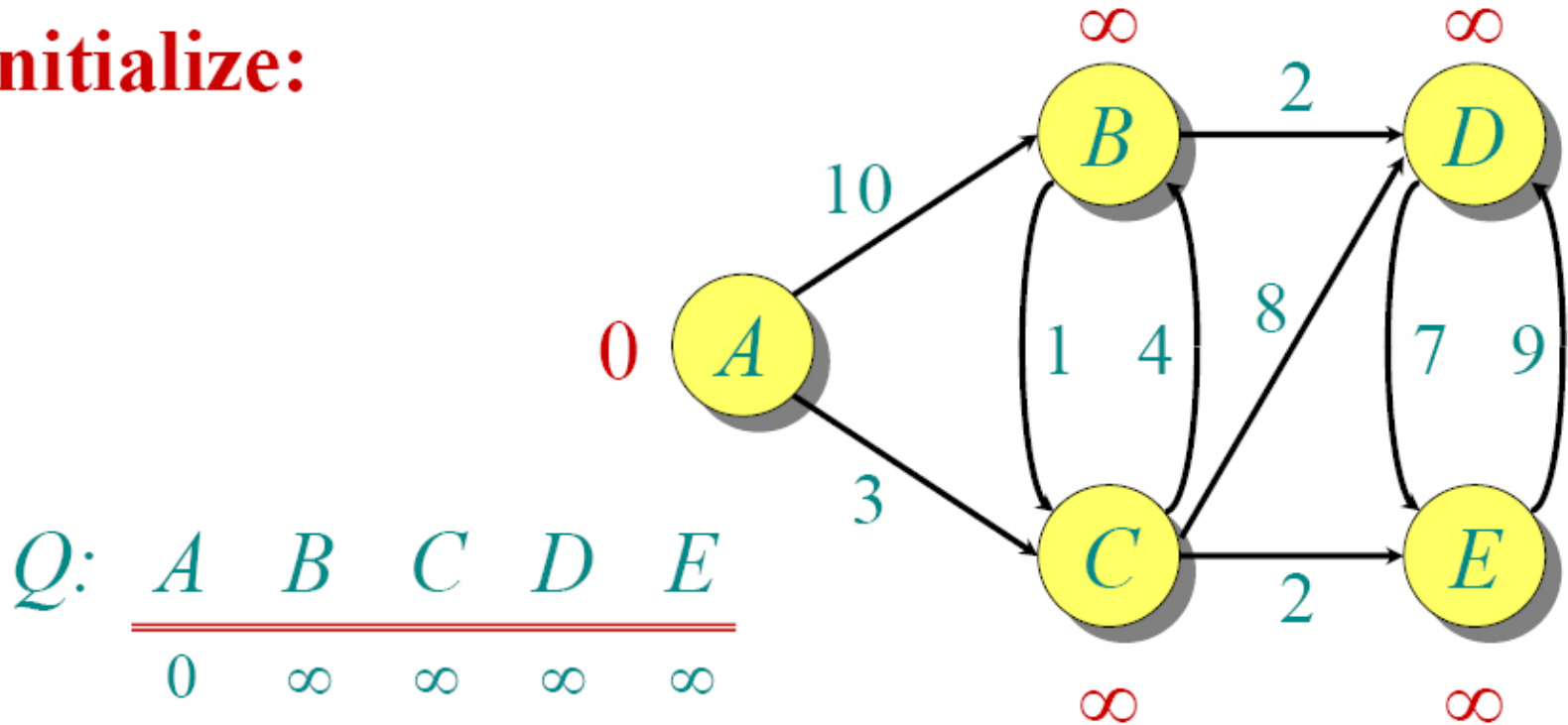
# DIJKSTRA'S ALGORITHM - PSEUDOCODE

```
dist[s] ← 0
for all v ∈ V - {s}
    do dist[v] ← ∞
S ← ∅
Q ← V
while Q ≠ ∅
do u ← mindistance(Q, dist)
   S ← S ∪ {u}
   for all v ∈ neighbors[u]
       Relax {u, v, w}
return dist
```



# DIJKSTRA ANIMATED EXAMPLE

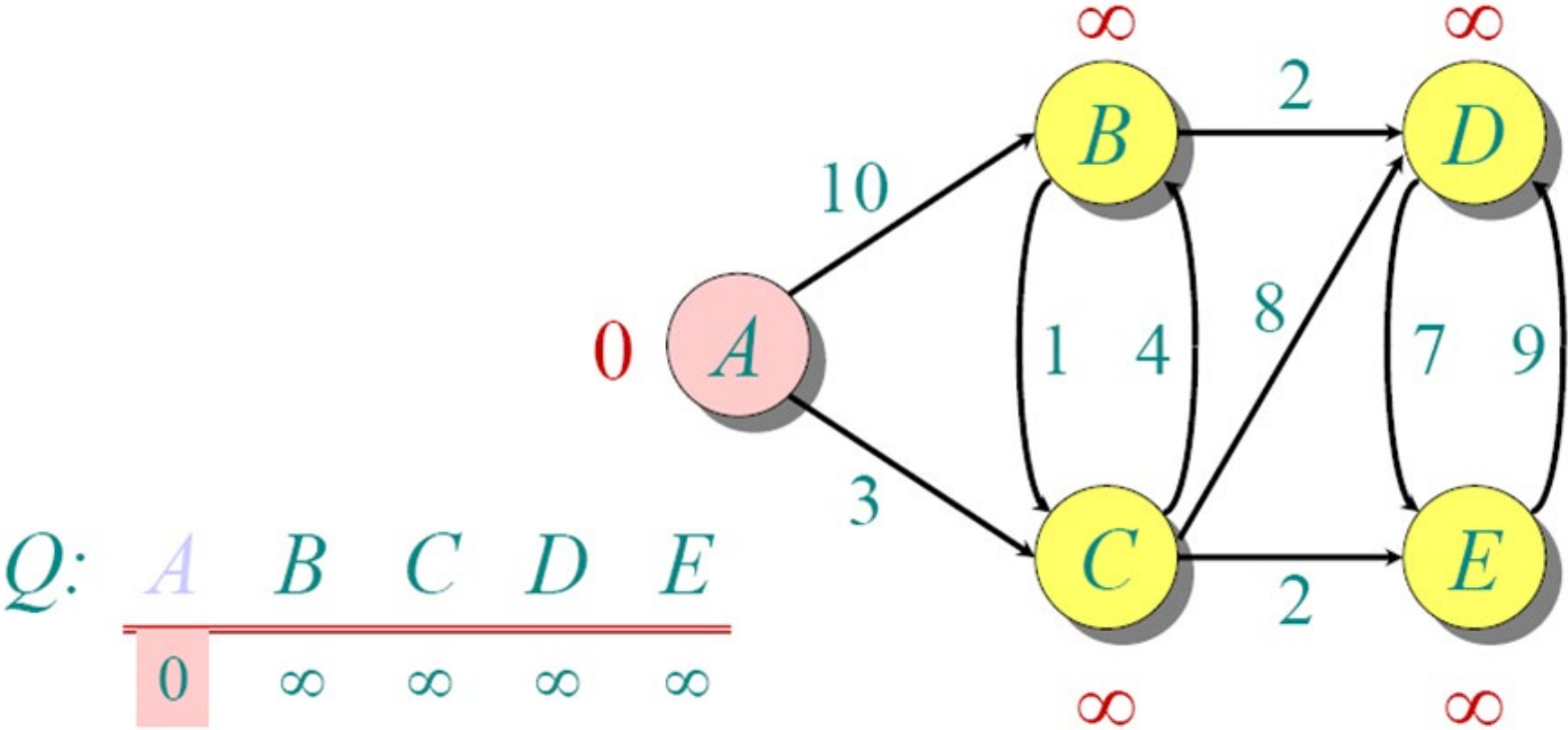
**Initialize:**



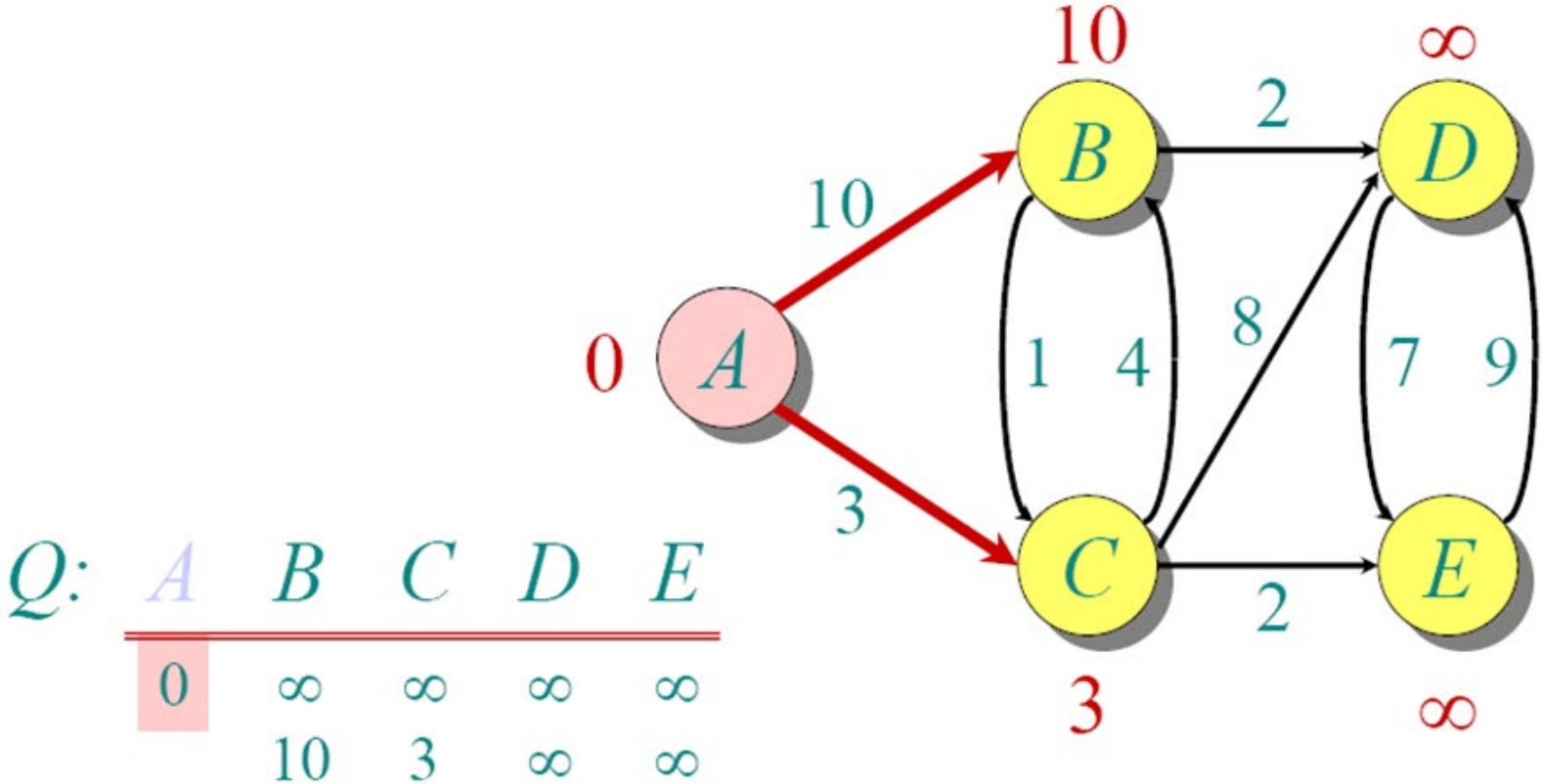
S: {}



# DIJKSTRA ANIMATED EXAMPLE



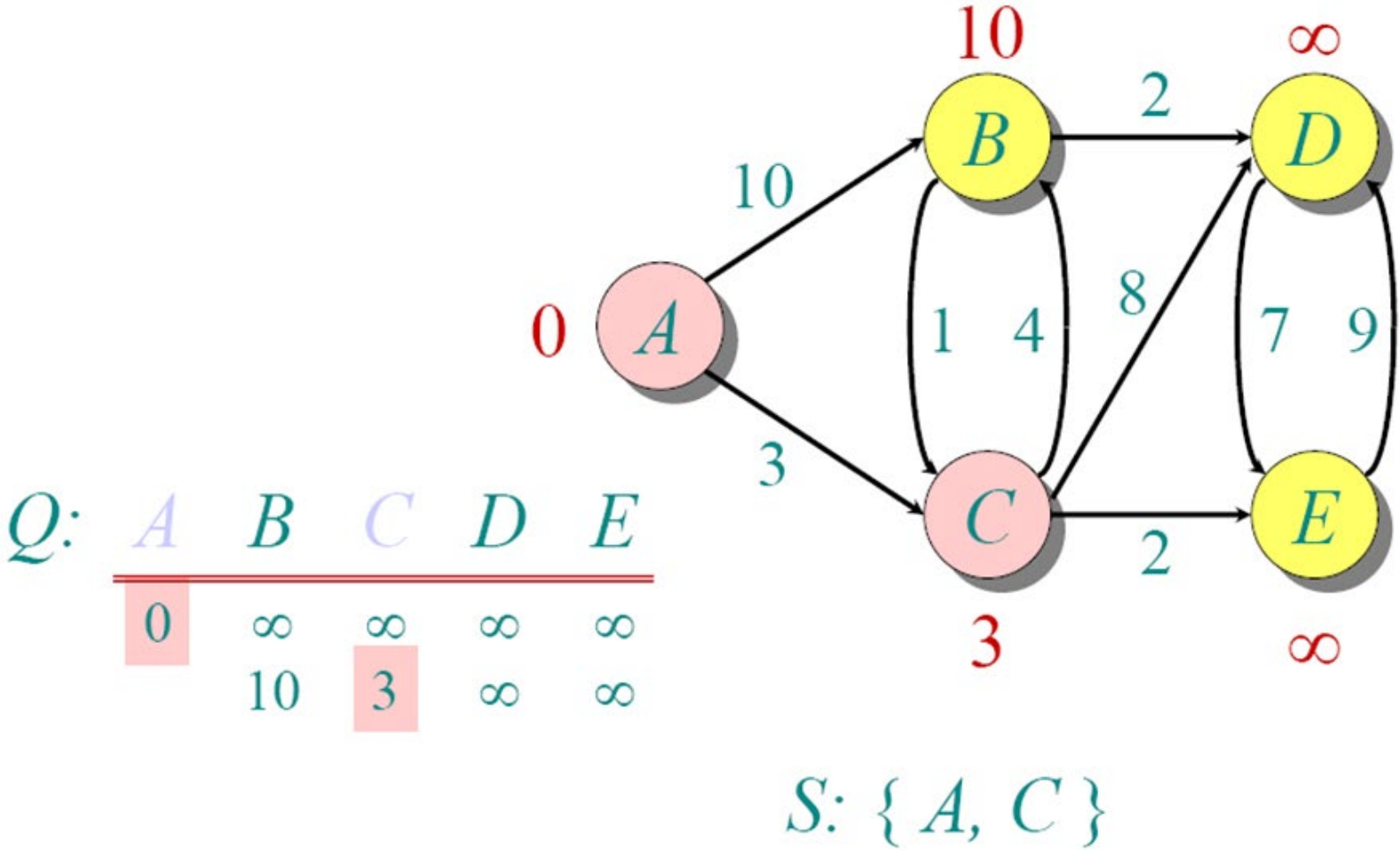
# DIJKSTRA ANIMATED EXAMPLE



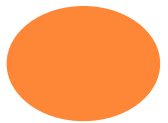
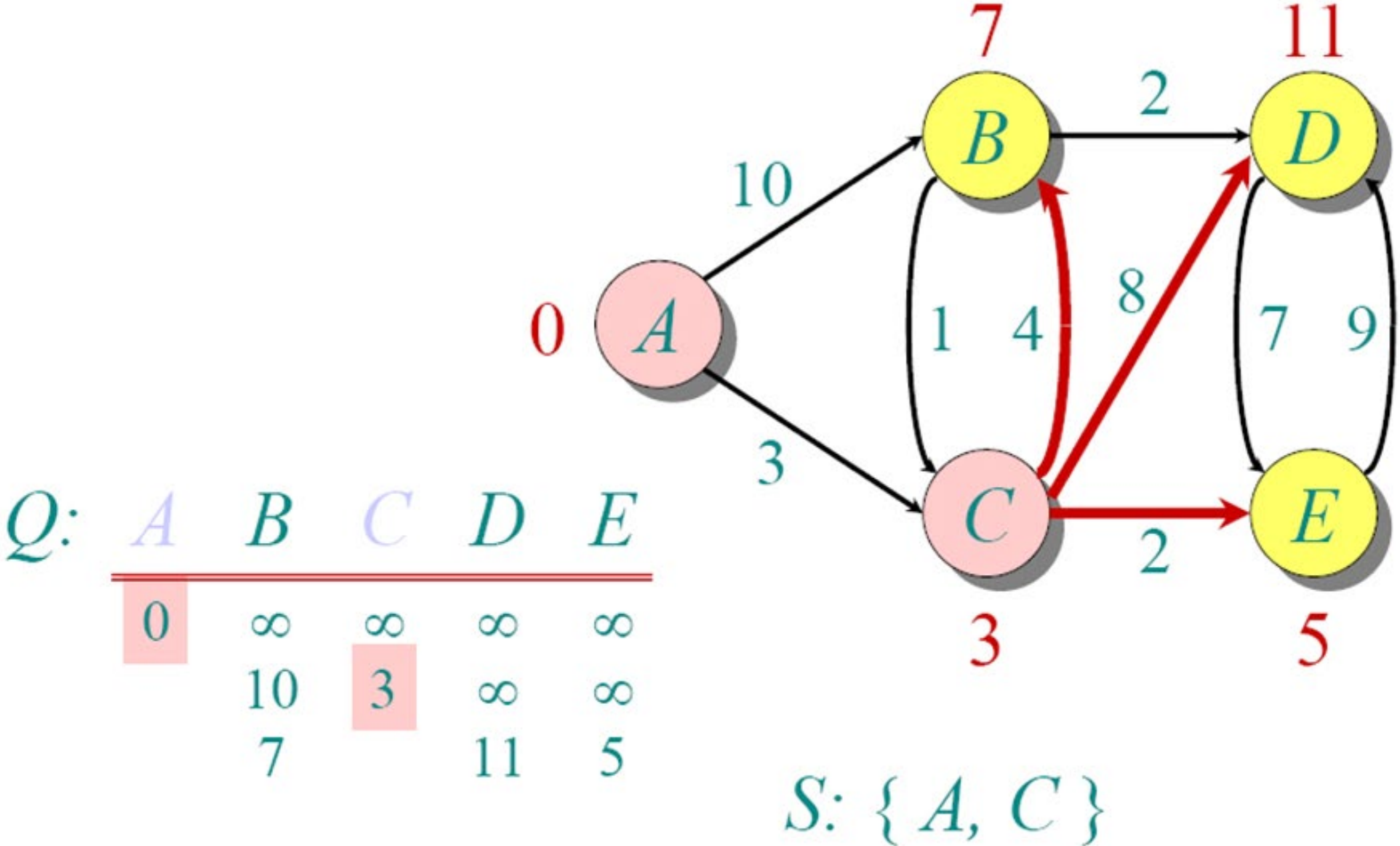
S: { A }



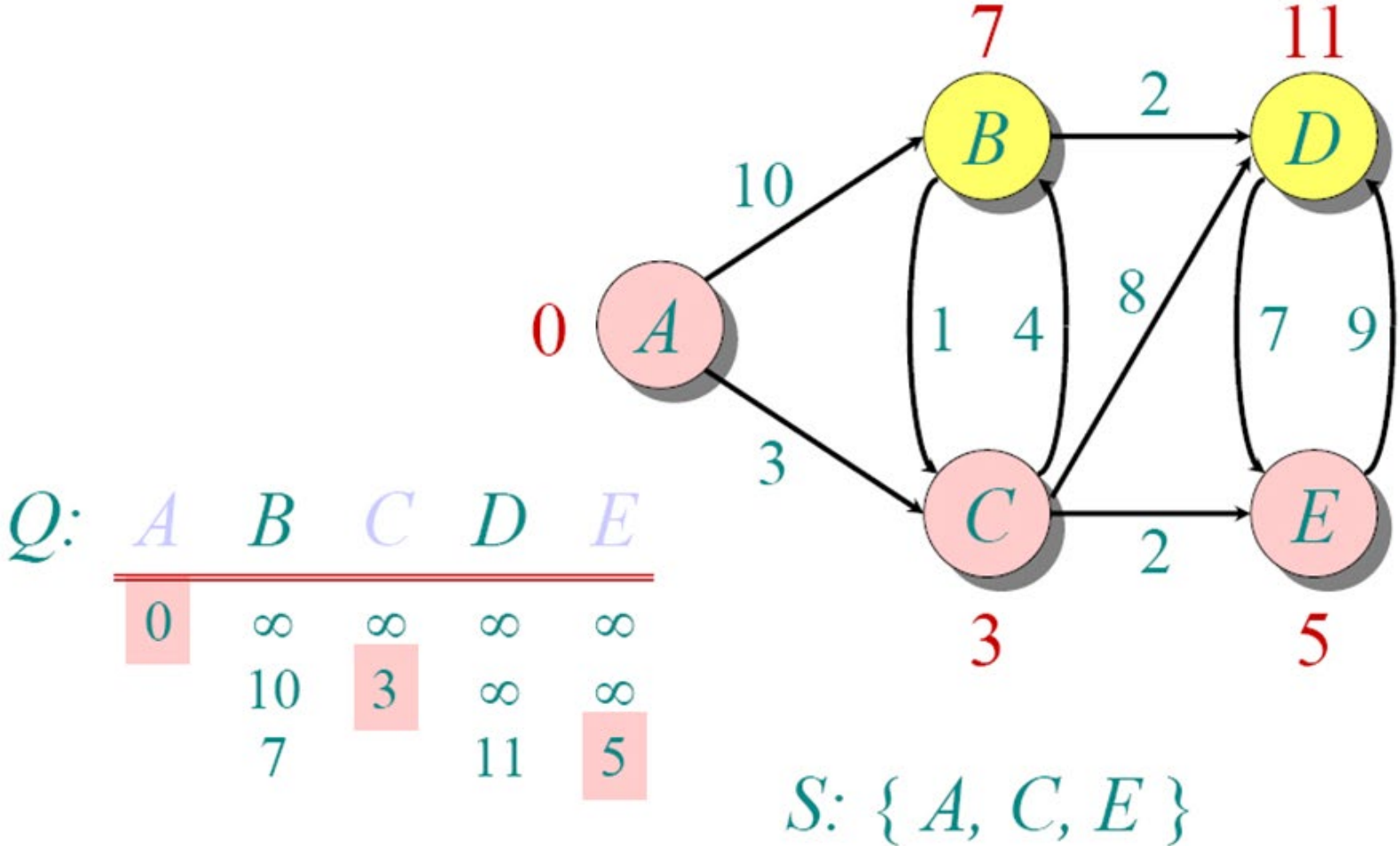
# DIJKSTRA ANIMATED EXAMPLE



# DIJKSTRA ANIMATED EXAMPLE

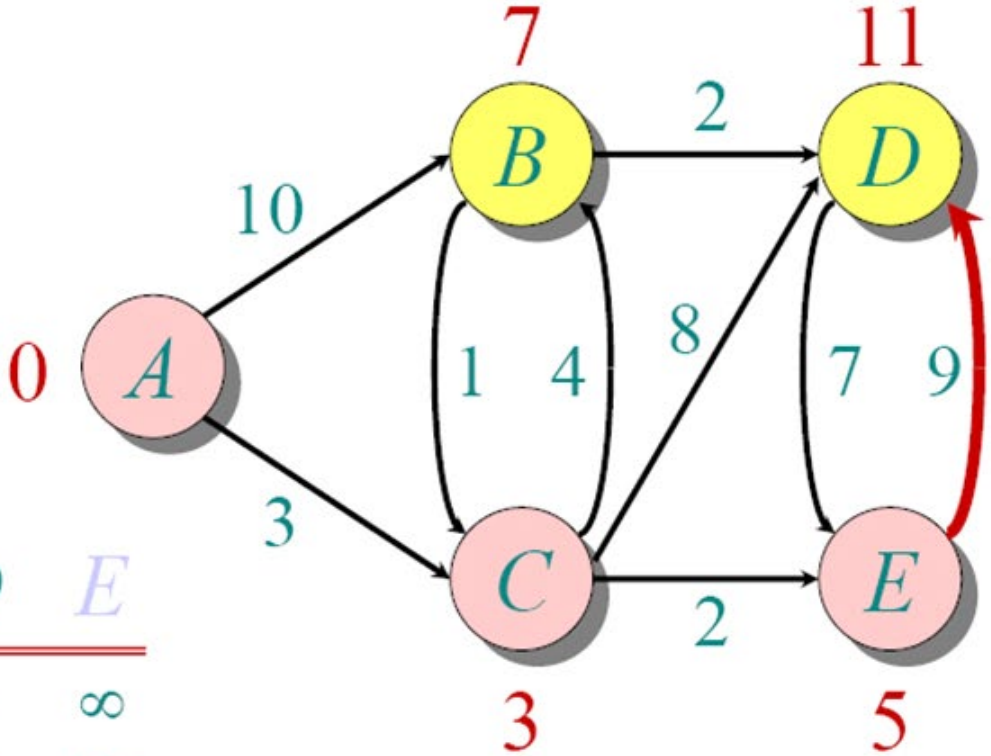


# DIJKSTRA ANIMATED EXAMPLE





# DIJKSTRA ANIMATED EXAMPLE



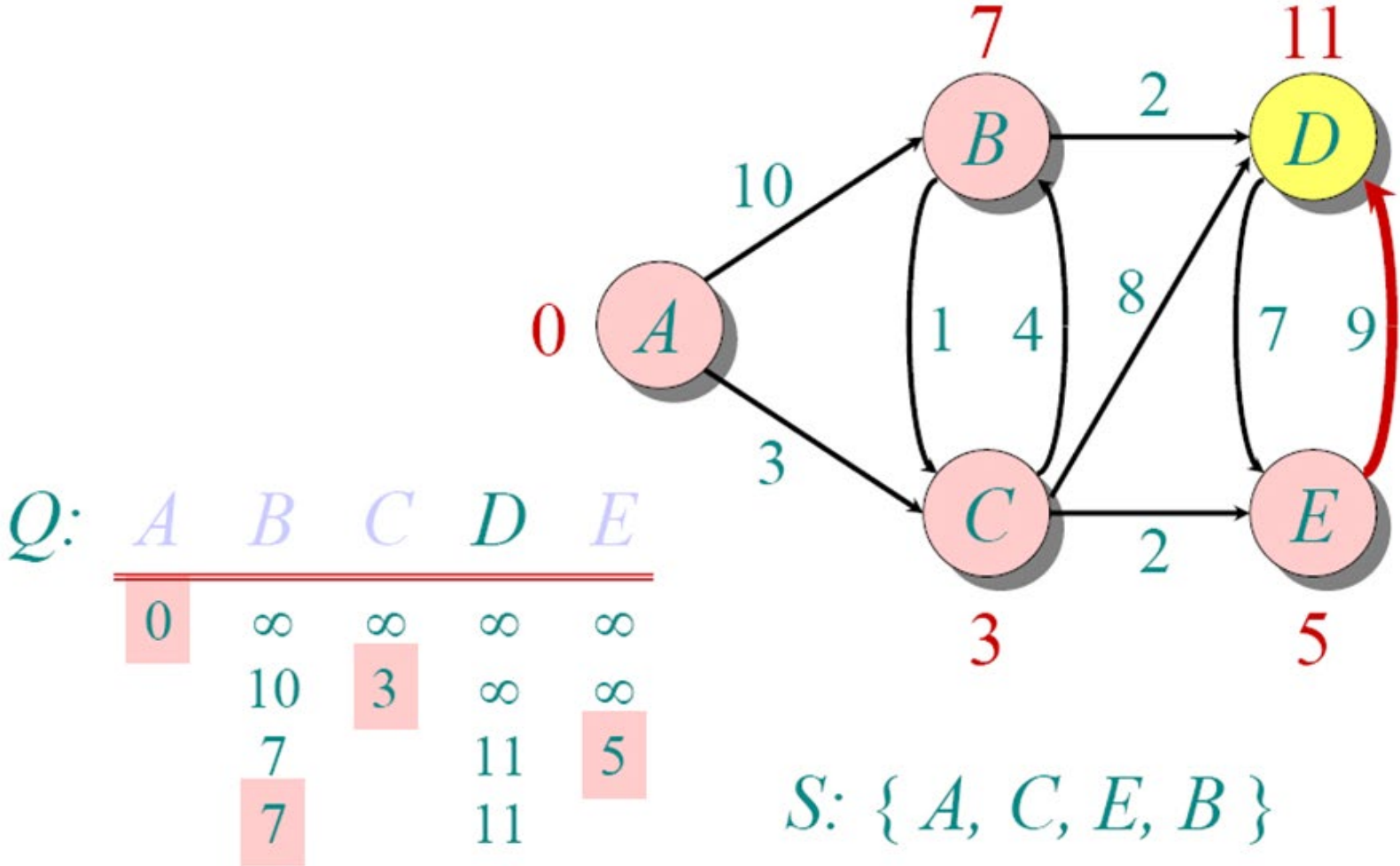
Q:

A	B	C	D	E
0	$\infty$	$\infty$	$\infty$	$\infty$
	10	3	$\infty$	$\infty$
	7		11	5
	7		11	

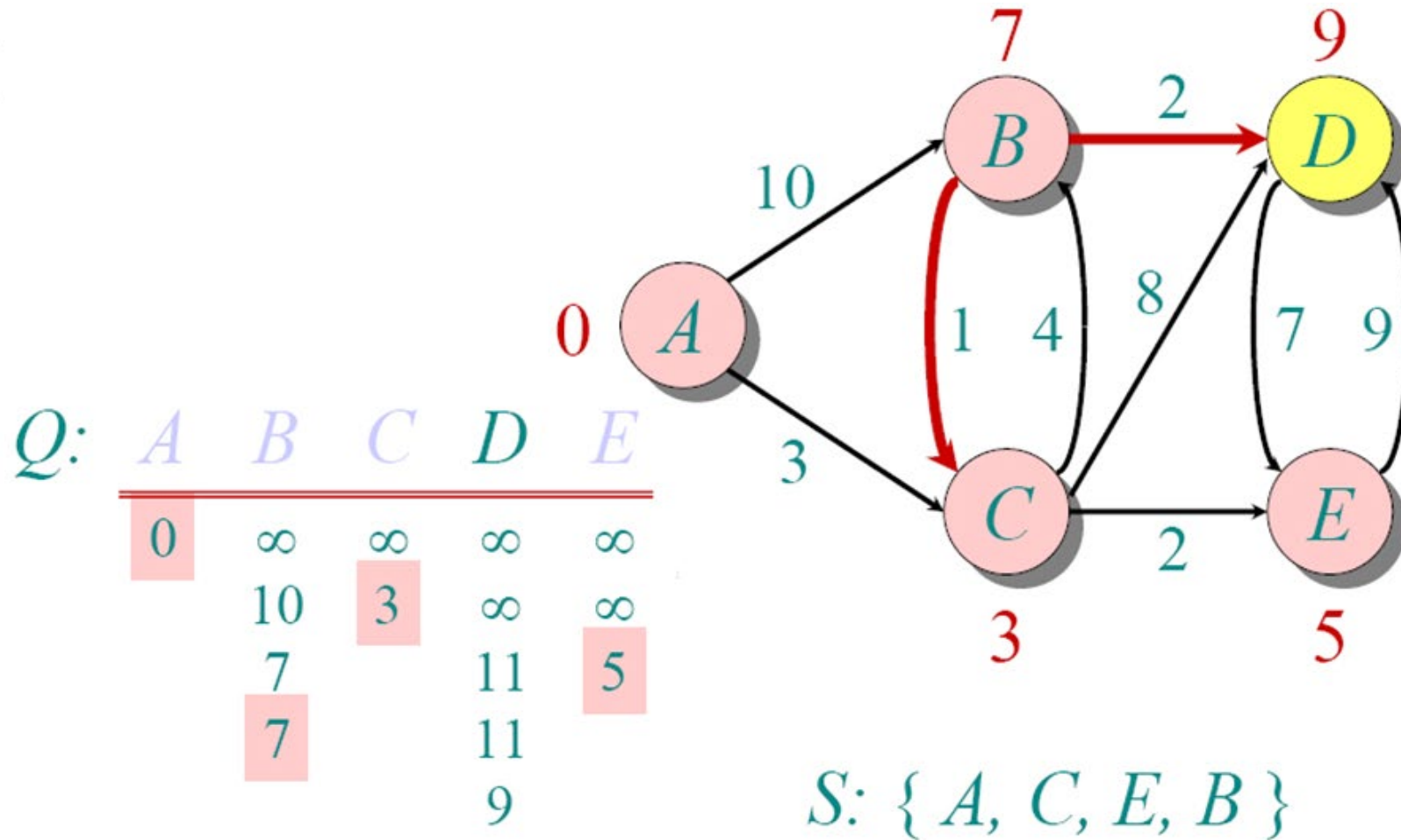
S: { A, C, E }



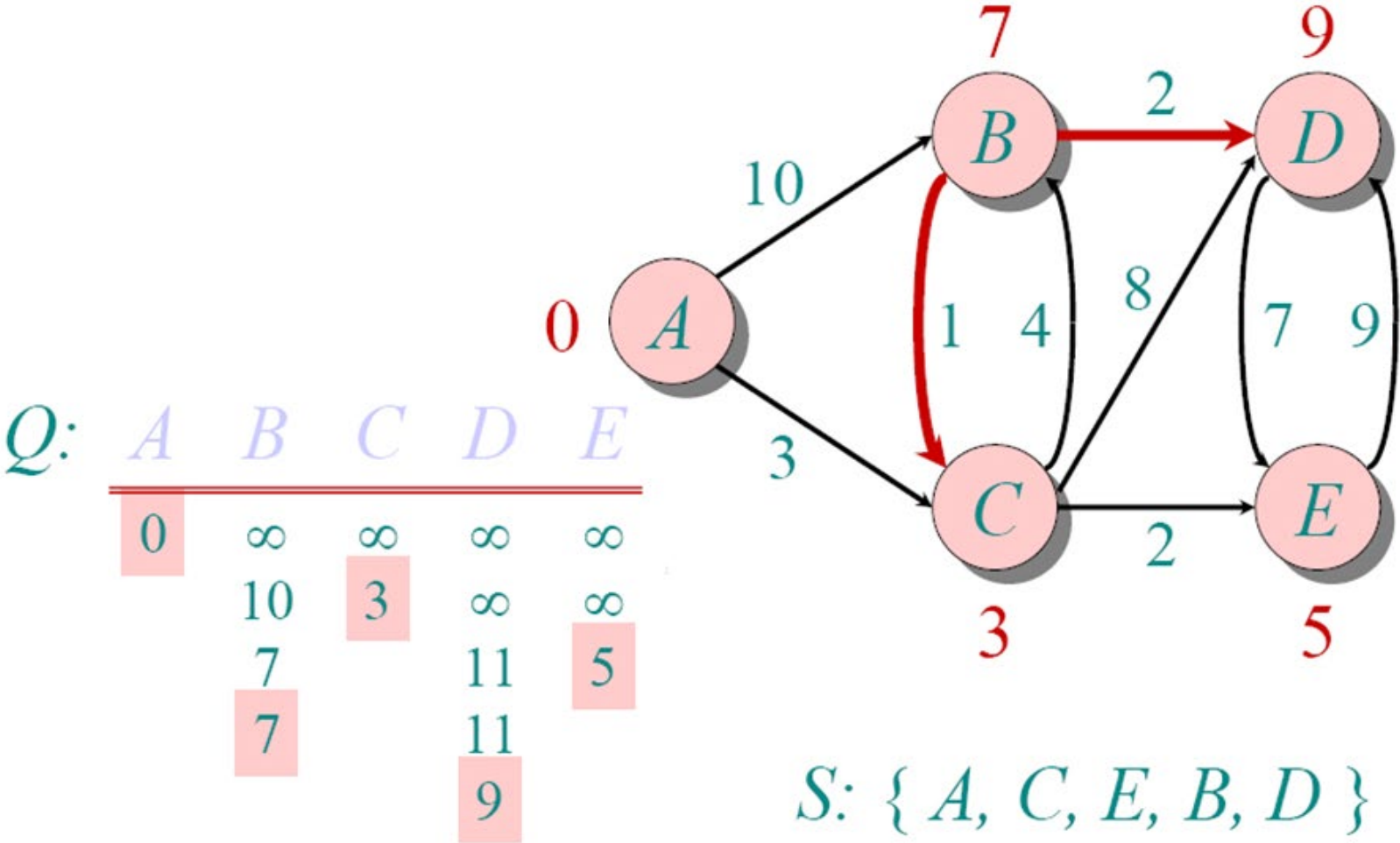
# DIJKSTRA ANIMATED EXAMPLE



# DIJKSTRA ANIMATED EXAMPLE



# DIJKSTRA ANIMATED EXAMPLE



# DIJKSTRA'S ALGORITHM - CORRECTNESS

- Because of time constraint we will not go deep into correctness proof
- The algorithm's correctness can be proved via induction
- Hints:
  - Optimal Substructure Property
    - Optimal solution can be constructed from optimal solutions of its subproblems
  - Relaxation is safe
  - Non-negative weights



# IMPLEMENTATIONS AND RUNNING TIMES

- For the given priority queue we are performing three different operation
  - Build
  - Extract minimum
  - Decrease key

Priority Queue $Q'$ on $n$ items	$Q$ Operations $O(\cdot)$			Dijkstra $O(\cdot)$
	$\text{build}(X)$	$\text{delete\_min}()$	$\text{decrease\_key}(id, k)$	$n =  V  = O( E )$
Array	$n$	$n$	1	$ V ^2$
Binary Heap	$n$	$\log n_{(a)}$	$\log n$	$ E  \log  V $
Fibonacci Heap	$n$	$\log n_{(a)}$	$1_{(a)}$	$ E  +  V  \log  V $



# DIJKSTRA VS A\*



Visualization of A\*



Visualization of Dijkstra

## Why Dijkstra?

**No Heuristic Dependency:** Dijkstra's algorithm does not require a heuristic function, making it more suitable when no good heuristic is available or when you need guaranteed correctness without the risk of an inadmissible/inconsistent heuristic.

**Guaranteed Exploration:** Dijkstra explores all reachable vertices, making it ideal for applications like finding the shortest path to all nodes (e.g., in network routing) rather than a single destination.

# THE BELLMAN-FORD SHORTEST PATH ALGORITHM



NEIL TANG  
03/11/2010



## SHORTEST SIMPLE PATH

- Shortest Path (If exists)

$$\text{Shortest Path} = \min\{w(\pi) \mid \text{paths } \pi \text{ } s \rightarrow t\}$$

- If  $\delta(s, v)$  is finite then there is a shortest path from s-v is simple
- **Simple path** in a graph is a path that does not revisit any vertex
  - At most  $|v|$  vertex
  - At most  $|v|-1$  edges
- $\delta_{|V|}(s, v) < \delta_{|V|-1}(s, v)$  then  $\delta(s, v) = -\infty$



# THE BELLMAN-FORD ALGORITHM

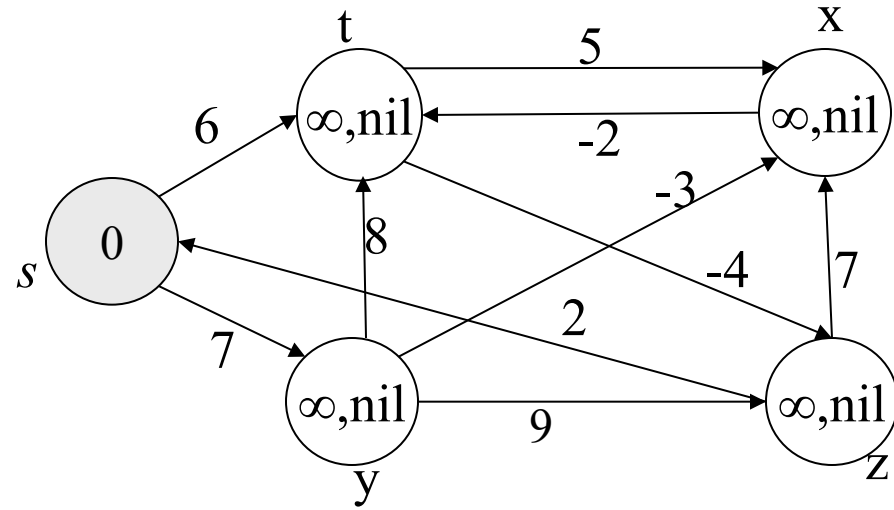
## Bellman-Ford( $G, w, s$ )

1. Initialize-Single-Source( $G, s$ )
2. **for**  $i := 1$  to  $|V| - 1$  **do**
3.     **for** each edge  $(u, v) \in E$  **do**
4.         Relax( $u, v, w$ )
5.     **for** each vertex  $v \in u.\text{adj}$  **do**
6.         **if**  $d[v] > d[u] + w(u, v)$
7.             **then return** False // there is a negative cycle
8.     **return** True

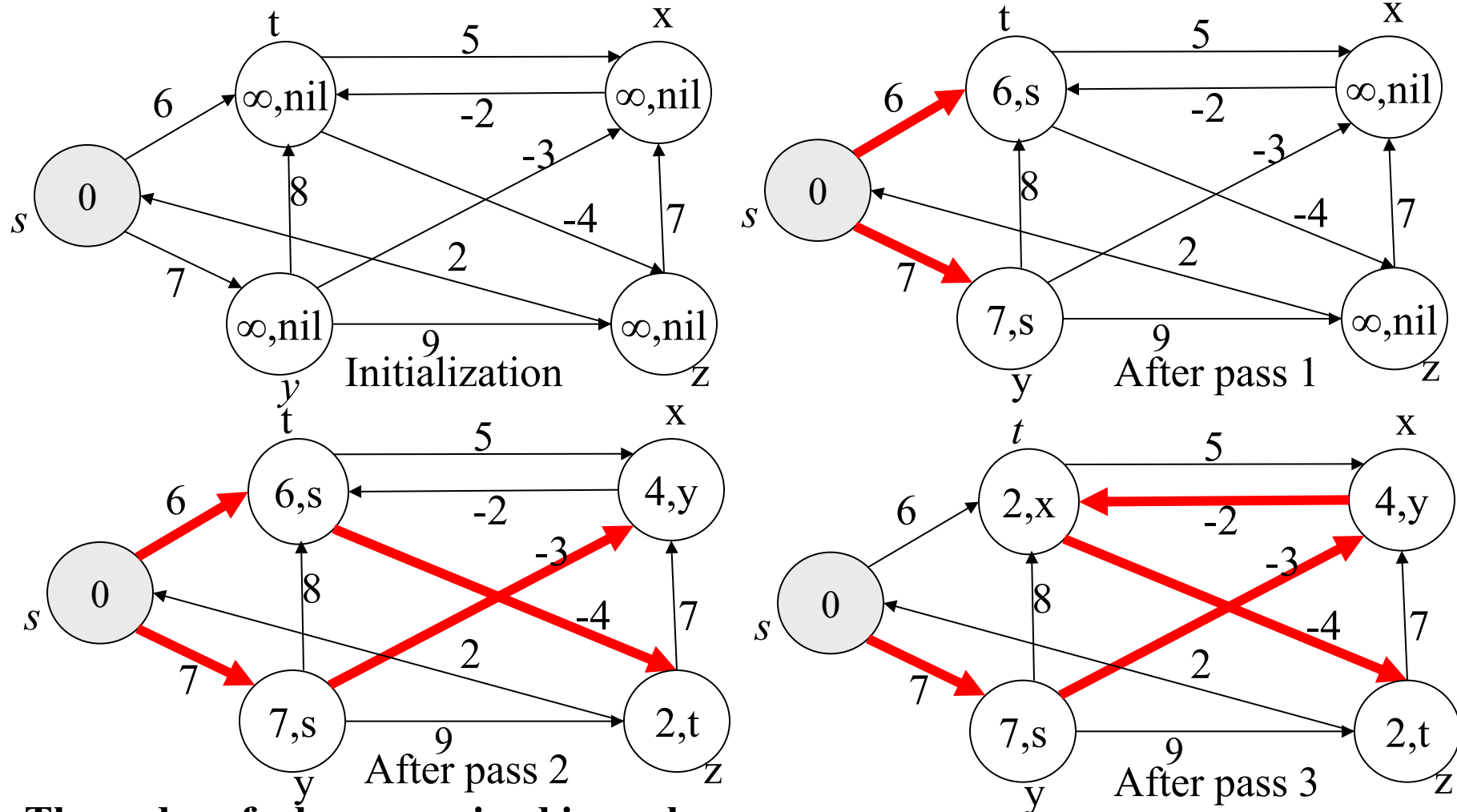
## Relax( $u, v, w$ )

- ```
if  $d[v] > d[u] + w(u, v)$   
  then  $d[v] := d[u] + w(u, v)$   
        $\text{parent}[v] := u$ 
```

# THE BELLMAN-FORD ALGORITHM



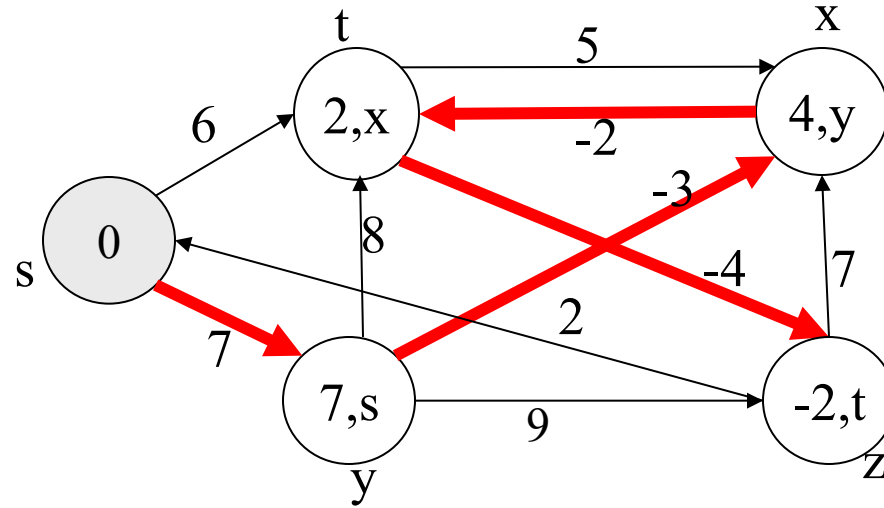
# GRAPH DUPLICATION



**The order of edges examined in each pass:**

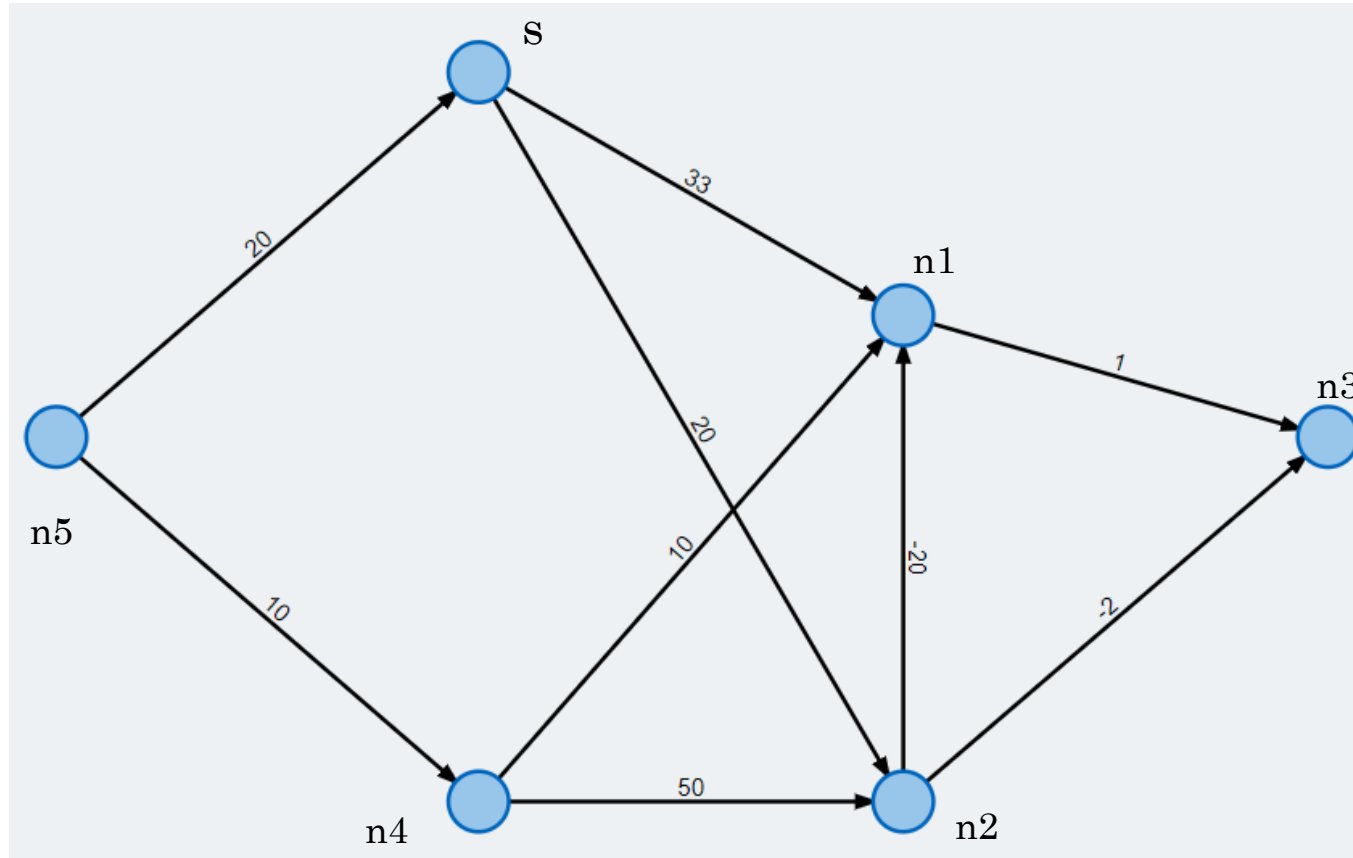
$(s, t), (s, y), (t, x), (t, z), (x, t), (y, x), (y, t), (y, z), (z, x), (z, s),$

# THE BELLMAN-FORD ALGORITHM



After pass 4

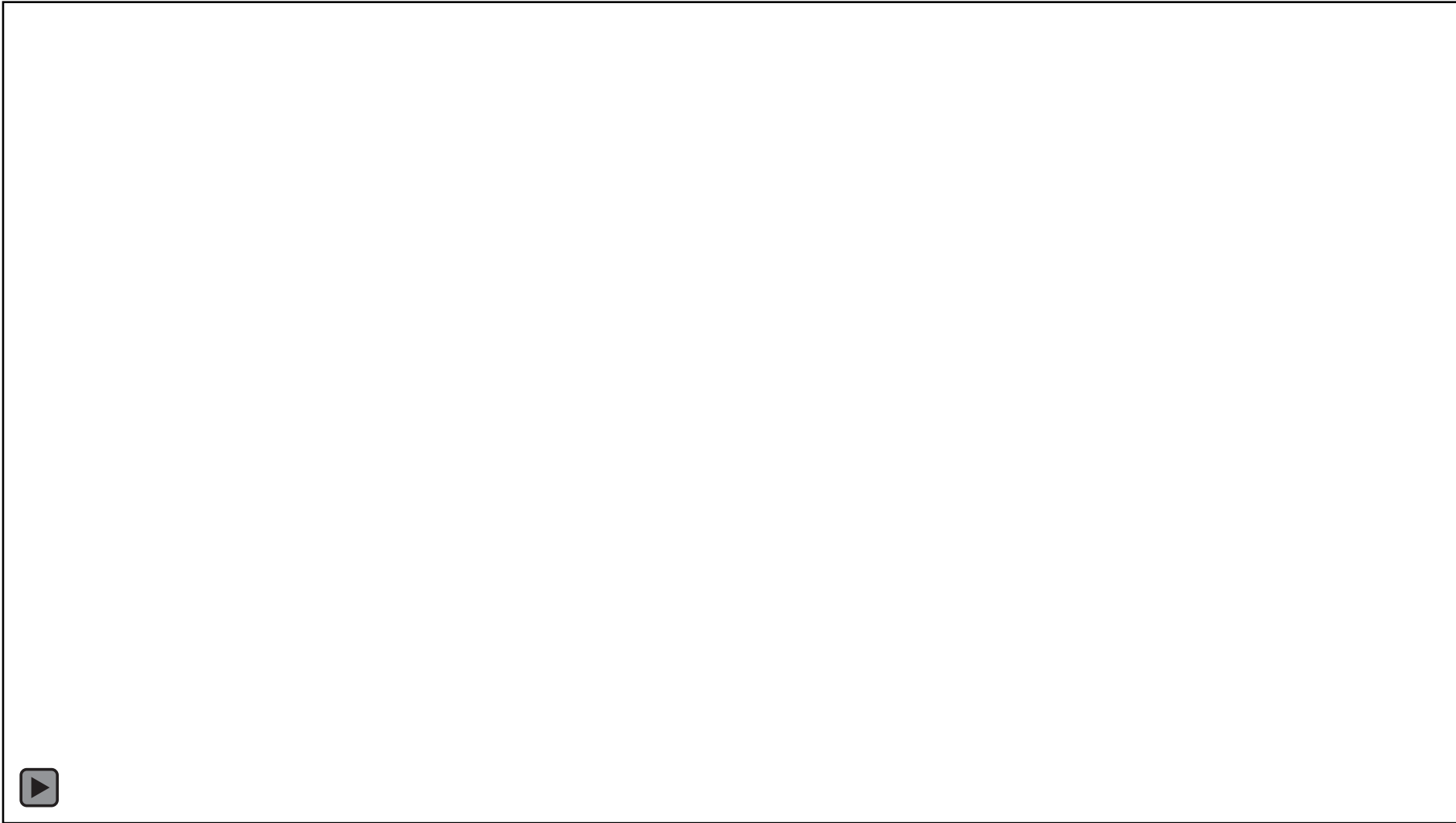
# THE BELLMAN-FORD ALGORITHM VISUALIZATION



**The order of edges examined in each pass:**

(s, n1), (n2, n3), (n2, n1), (n1, n3), (s, n2), (n4, n1), (n4, n2), (n5, s), (n5, n4)

# THE BELLMAN-FORD ALGORITHM VISUALIZATION

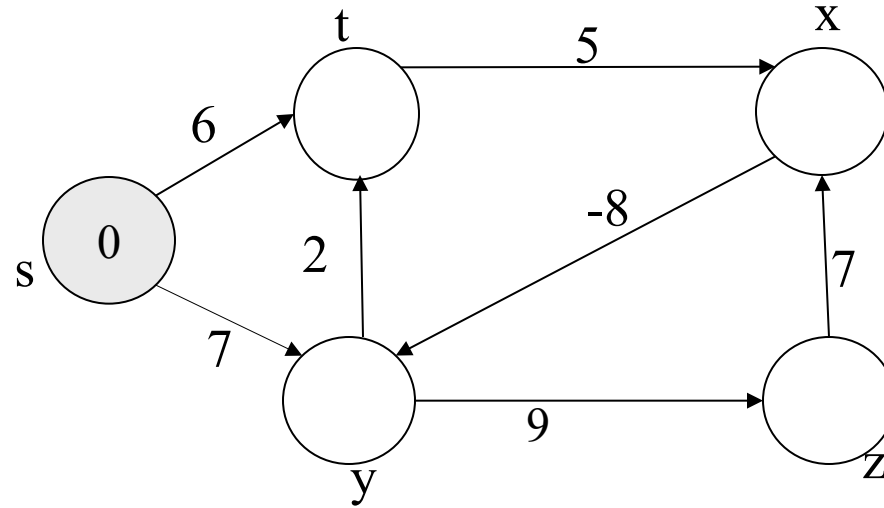


**The order of edges examined in each pass:**

$(s, n1), (n2, n3), (n2, n1), (n1, n3), (s, n2), (n4, n1), (n4, n2), (n5, s), (n5, n4)$

*Vis Credit : TUM Bellman Ford : [https://algorithms.discrete.ma.tum.de/graph-algorithms/spp-bellman-ford/index\\_en.html](https://algorithms.discrete.ma.tum.de/graph-algorithms/spp-bellman-ford/index_en.html)*

# THE BELLMAN-FORD ALGORITHM



**The order of edges examined in each pass:**

(s, t), (s, y), (t, x), (x, y), (y, t), (y, z), (z, x),

(s, t), (s, y), (t, x), (x, y), (y, t), (y, z), (z, x),

(s, t), (s, y), (t, x), (x, y), (y, t), (y, z), (z, x),

(s, t), (s, y), (t, x), (x, y), (y, t), (y, z), (z, x),



# TIME COMPLEXITY

## Bellman-Ford( $G, w, s$ )

1. Initialize-Single-Source( $G, s$ )  $\longrightarrow$   $O(|V|)$
2. **for**  $i := 1$  to  $|V| - 1$  **do**
3.     **for** each edge  $(u, v) \in E$  **do**  $\longrightarrow$   $O(|V||E|)$
4.         Relax( $u, v, w$ )  $\longrightarrow$   $O(|V||E|)$
5.     **for** each vertex  $v \in u.\text{adj}$  **do**  $\longrightarrow$   $O(|E|)$
6.         if  $d[v] > d[u] + w(u, v)$
7.             **then return** False // there is a negative cycle
8.     **return** True

Time complexity:  $O(|V||E|)$

# DIFFERENCES

- Negative link weight: The Bellman-Ford algorithm works; Dijkstra's algorithm doesn't.
- Time complexity: The Bellman-Ford algorithm is higher than Dijkstra's algorithm.

# REFERENCES

- Dijkstra's original paper:  
E. W. Dijkstra. (1959) *A Note on Two Problems in Connection with Graphs*. *Numerische Mathematik*, 1. 269-271.
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- Meyers, L.A. (2007) Contact network epidemiology: Bond percolation applied to infectious disease prediction and control. *Bulletin of the American Mathematical Society* 44: 63-86.
- Department of Mathematics, University of Melbourne. *Dijkstra's Algorithm*.  
<<http://www.ms.unimelb.edu.au/~moshe/620-261/dijkstra/dijkstra.html> > Accessed 4/25/09

