

# Roadmap

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## Part I: Shortest Paths

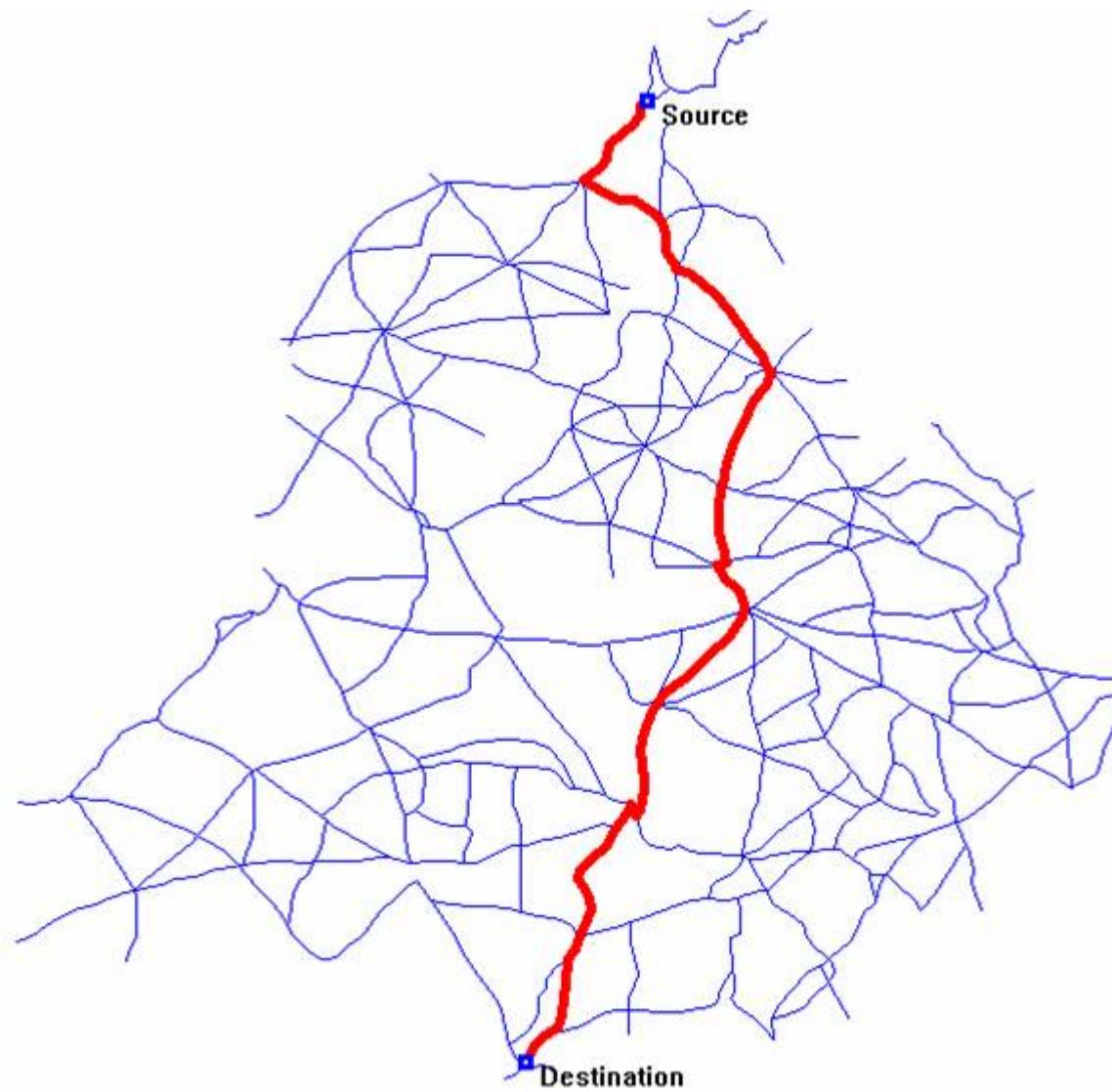
- Dijkstra

## Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

# SHORTEST PATHS

(ON WEIGHTED GRAPHS)



# Shortest Path Problem

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Basic question: **find the shortest path!**

- Source-to-destination: one vertex to another
- Single source: one vertex to every other
- All pairs: between all pairs of vertices

Variants:

- Edge weights: non-negative, arbitrary, Euclidean, ...
- Cycles: cyclic, acyclic, no negative cycles

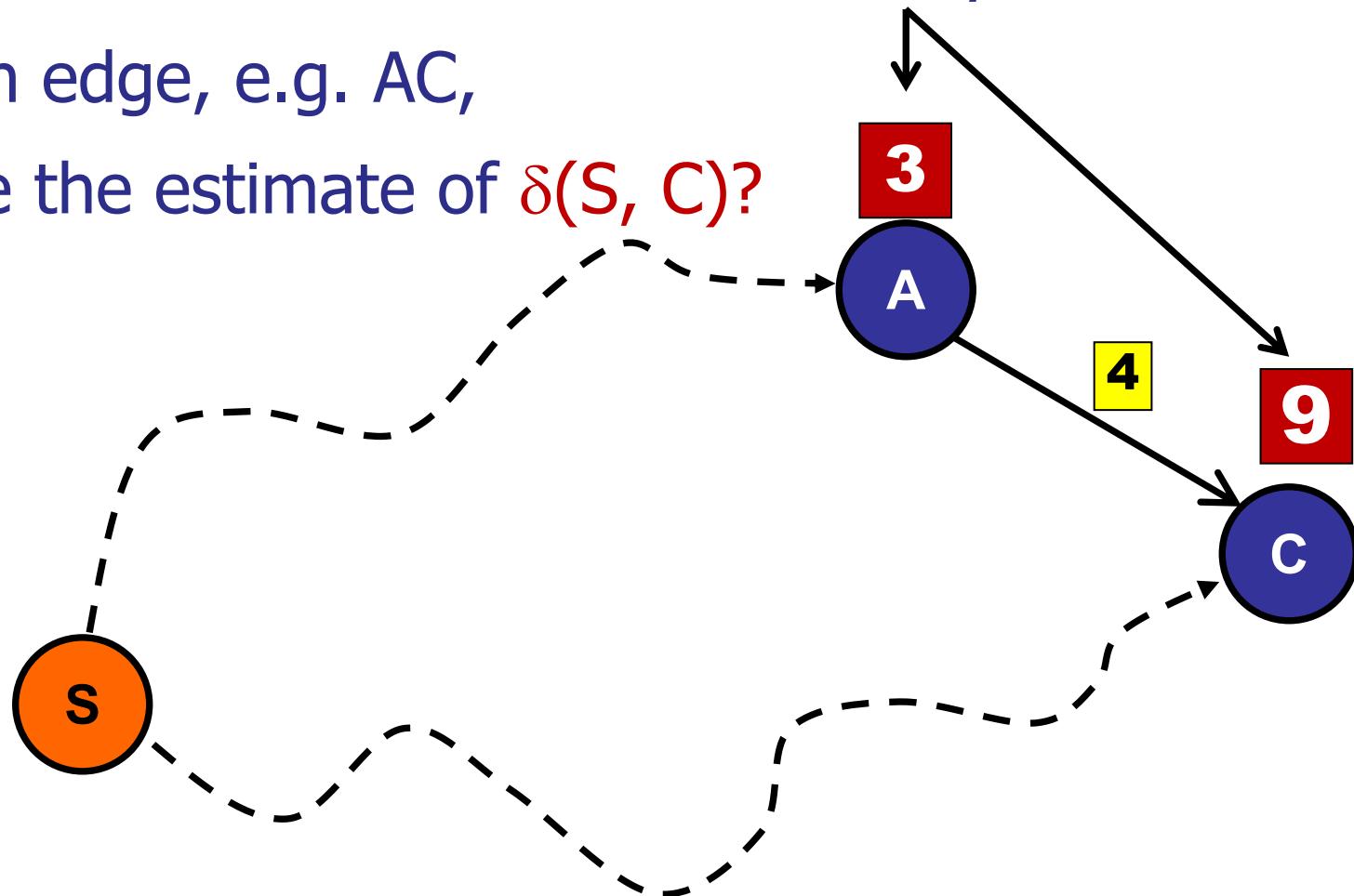
# Shortest Paths

Maintain estimate for each distance:

Lets say we have some estimate already

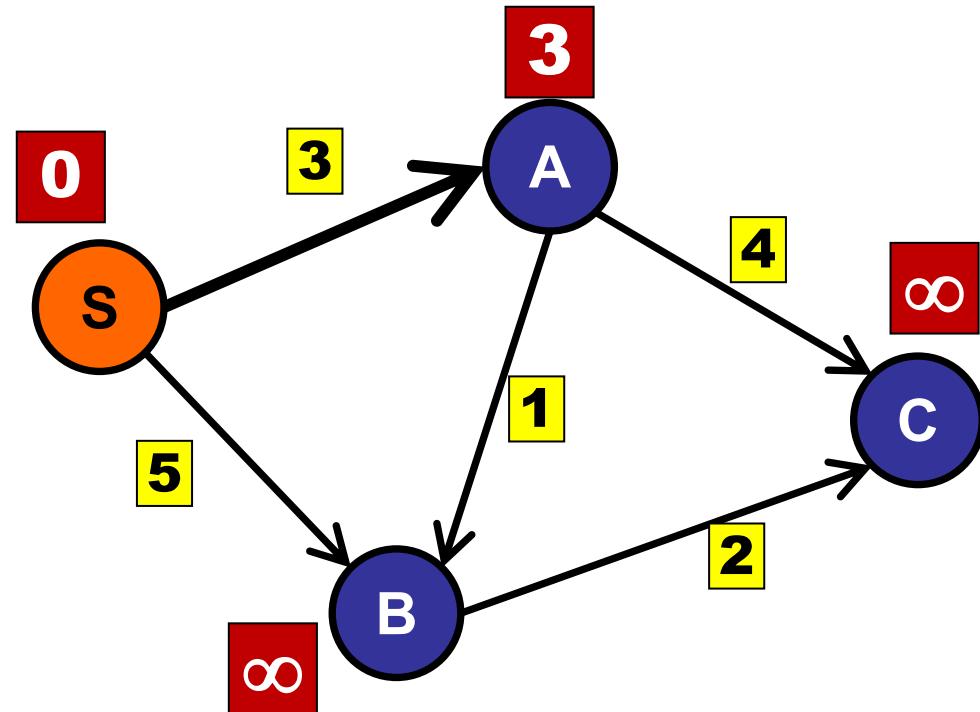
For each edge, e.g. AC,

Improve the estimate of  $\delta(S, C)$ ?



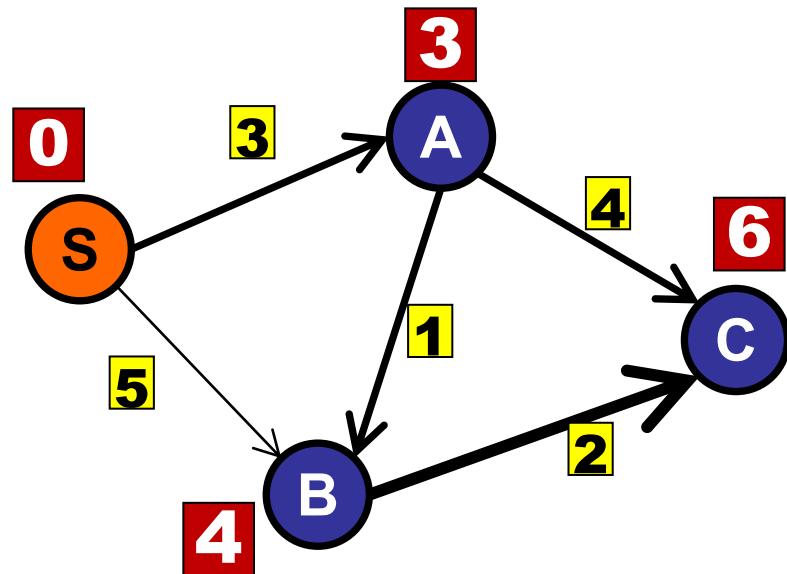
# Shortest Paths

```
relax(int u, int v) {  
    if (dist[v] > dist[u] + weight(u,v))  
        dist[v] = dist[u] + weight(u,v);  
}
```



# Bellman-Ford

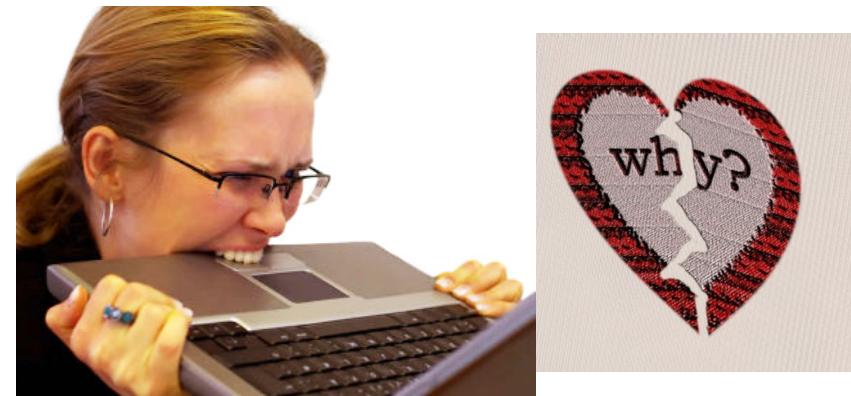
```
n = V.length;  
for (i=0; i<n; i++)  
    for (each edge e in the graph)  
        relax(e)
```



# Bellman-Ford Summary

Basic idea:

- Repeat  $|V|$  times: relax every edge
- Stop when “converges”.
- $O(VE)$  time.



Special issues:

- If negative weight-cycle: impossible.
- Use Bellman-Ford to **detect** negative weight cycle.
- If all weights are the same, use BFS.

# Today

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Key idea:

Relax the edges in the “right” order.

Only relax each edge once:

- $O(E)$  cost (for relaxation step).



# Edsger W. Dijkstra

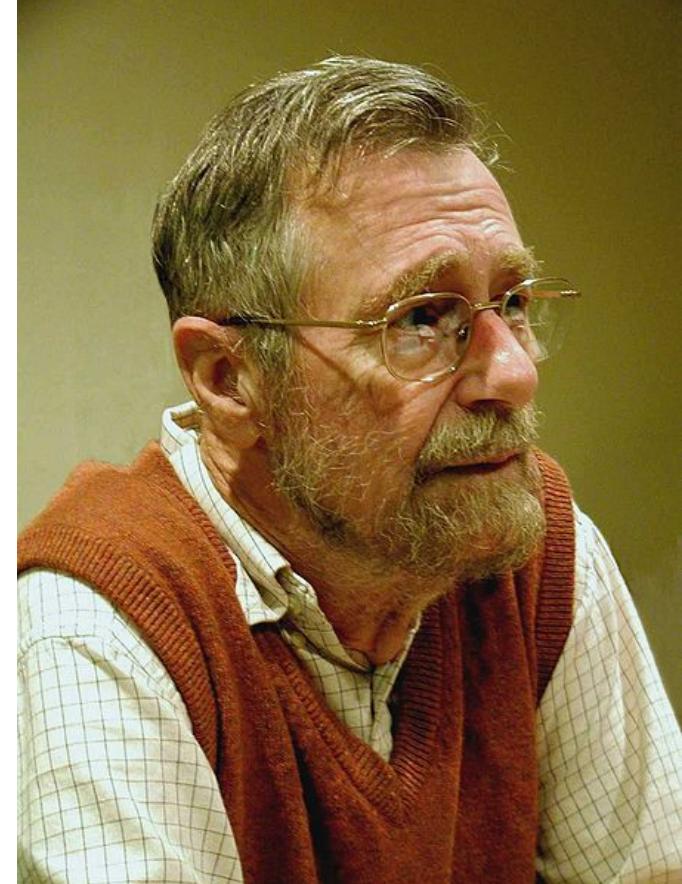
*“Computer science is no more about computers than astronomy is about telescopes.”*

*“The question of whether a computer can think is no more interesting than the question of whether a submarine can swim.”*

*“There should be no such thing as boring mathematics.”*

*“Elegance is not a dispensable luxury but a factor that decides between success and failure.”*

*“Simplicity is prerequisite for reliability.”*



1930-2002

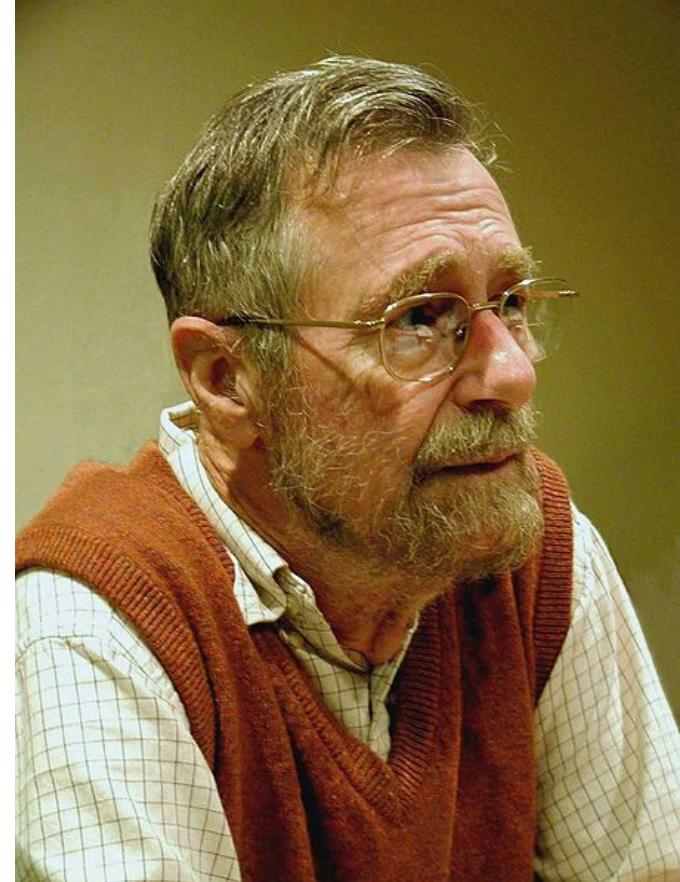
# Edsger W. Dijkstra

*“It is practically impossible to teach good programming to students that have had a prior exposure to BASIC: as potential programmers they are mentally mutilated beyond hope of regeneration.”*

*“The use of COBOL cripples the mind; its teaching should, therefore, be regarded as a criminal offense.”*

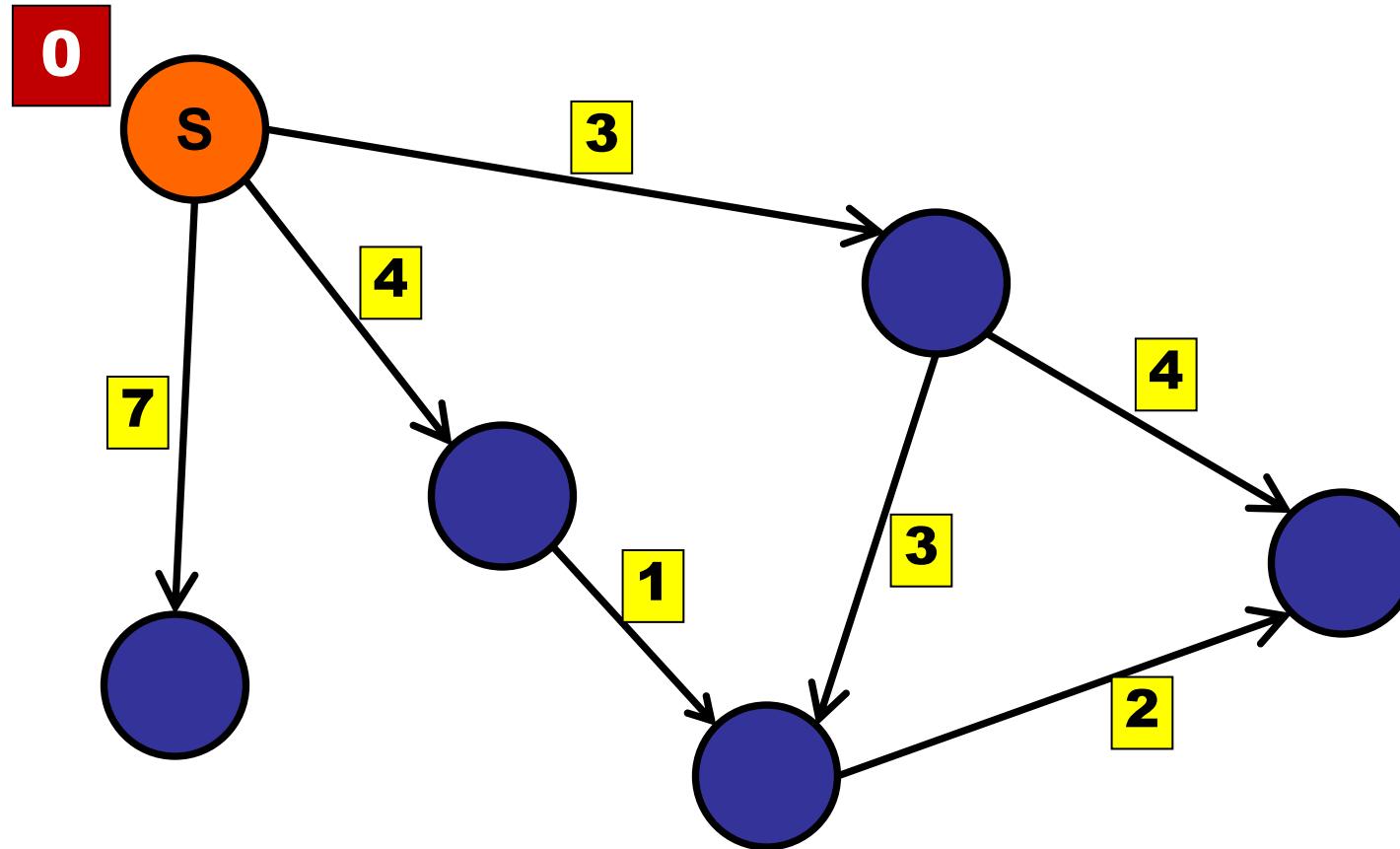
*“APL is a mistake, carried through to perfection. It is the language of the future for the programming techniques of the past: it creates a new generation of coding bums.”*

*“Object-oriented programming is an exceptionally bad idea which could only have originated in California.”*



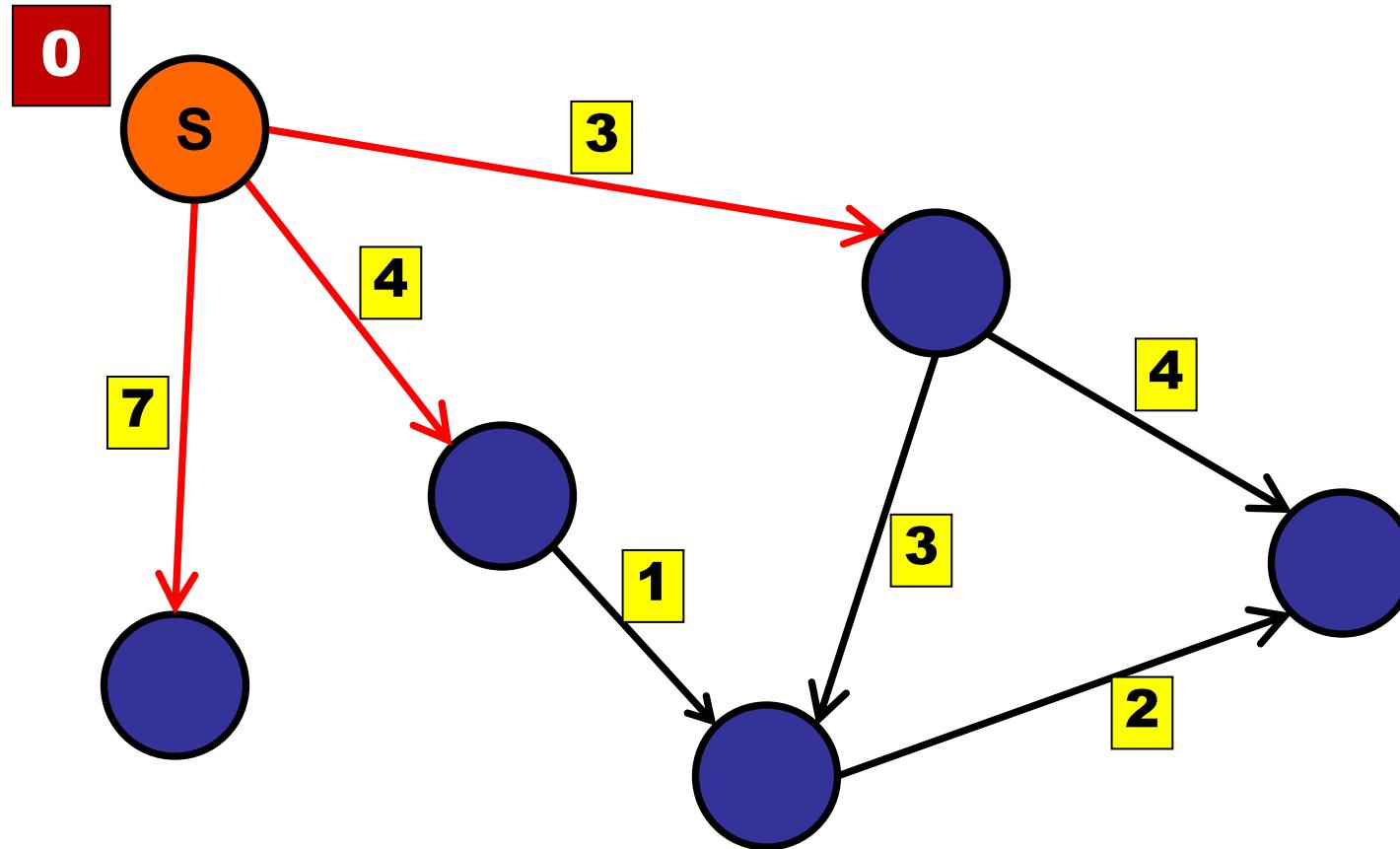
# Dijkstra's Algorithm (First Try)

Relax shortest edge first



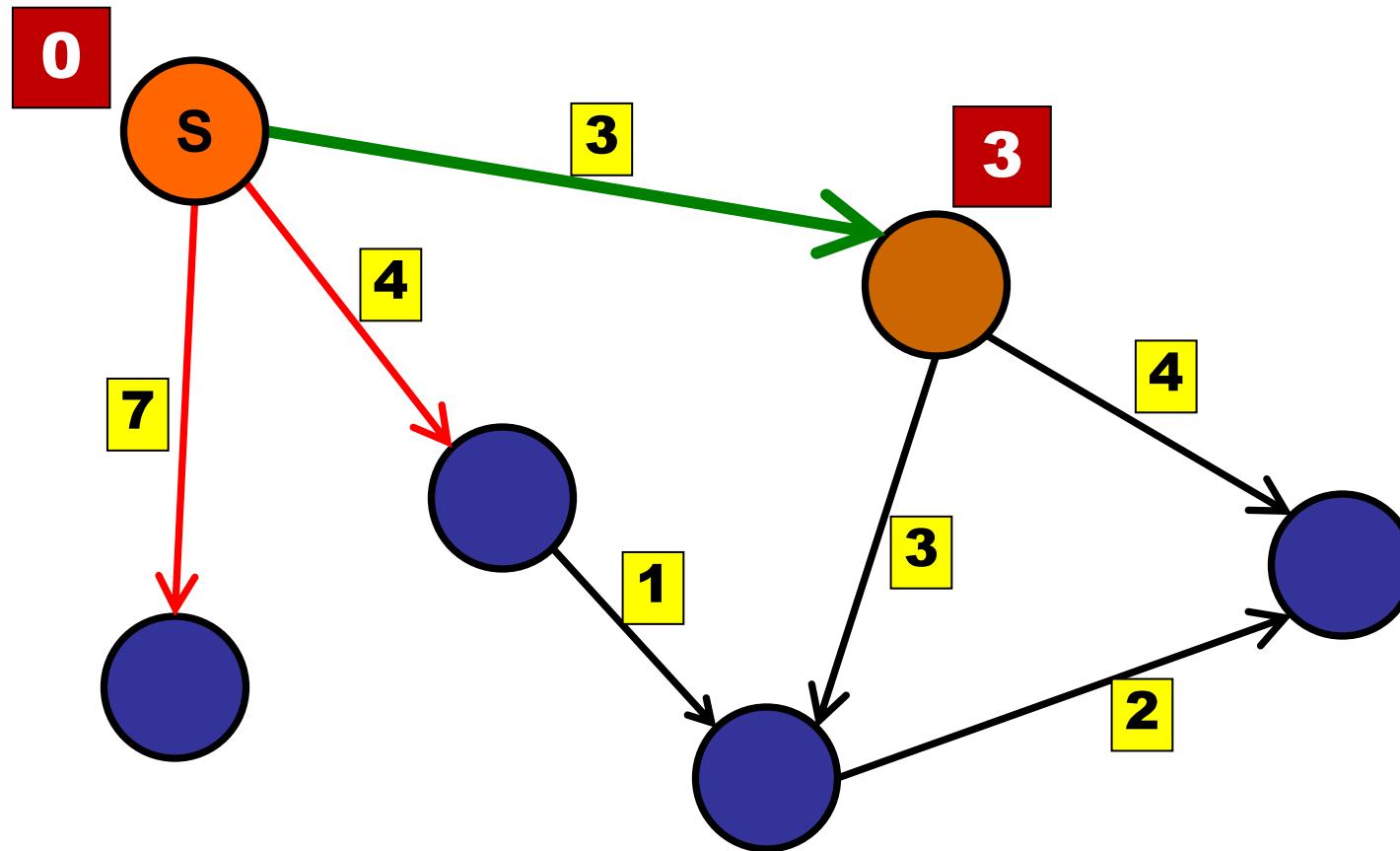
# Dijkstra's Algorithm (First Try)

Relax shortest edge first



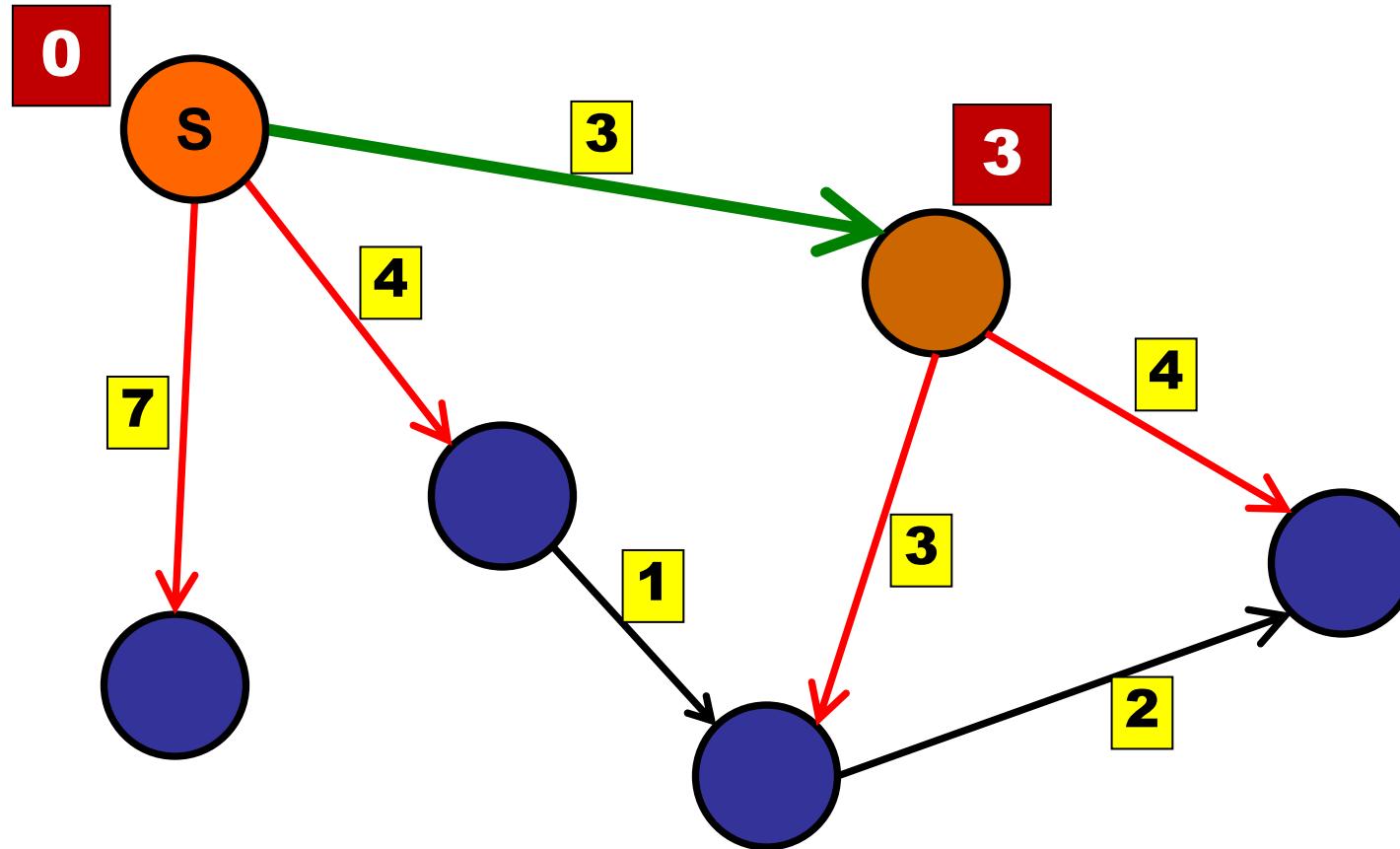
# Dijkstra's Algorithm (First Try)

Relax shortest edge first



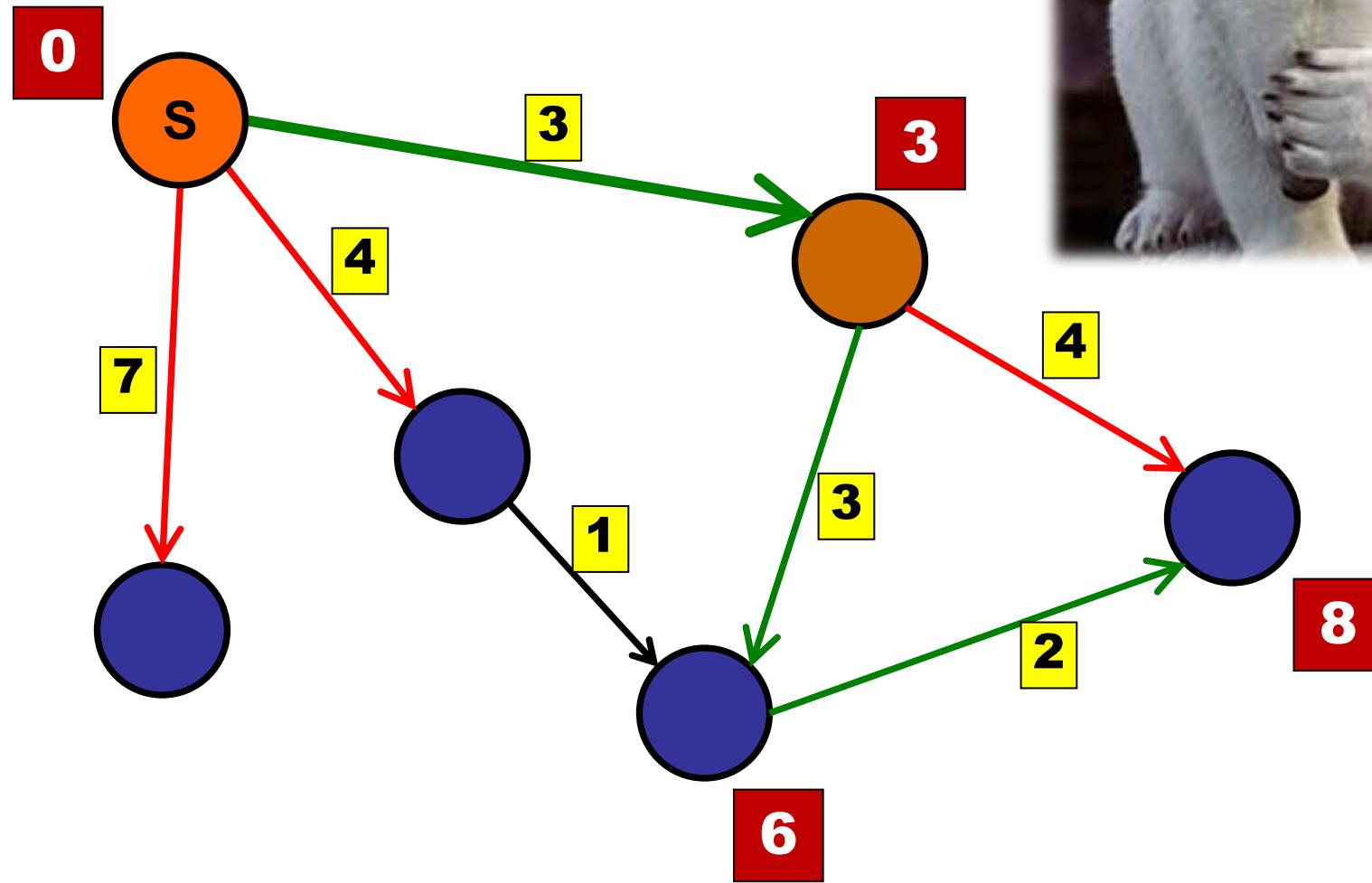
# Dijkstra's Algorithm (First Try)

Relax shortest edge first



# Dijkstra's Algorithm (Failed Try)

Oops....



# Dijkstra's Algorithm

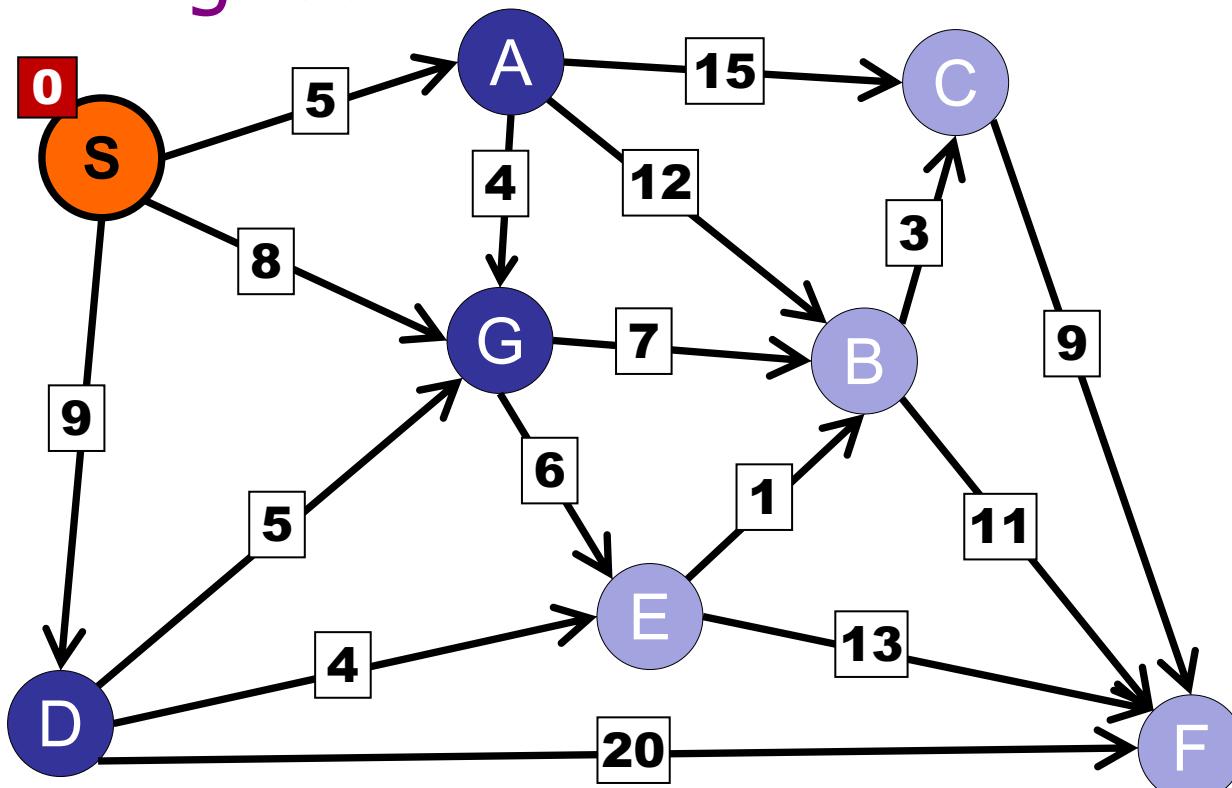
---

## Basic idea:

- Initialize:
  - Put all vertices into a priority queue
  - Set all priorities to estimated distances as infinity
  - Set the starting vertex estimated distance as 0
- Repeat until the priority is empty:
  - Extract the vertex  $v$  in the priority queue with the shortest estimated distance
  - Relax all the neighbors of  $v$  in the priority queue and update their estimated distance

# Dijkstra's Algorithm

Extract S and relax/update neighbors

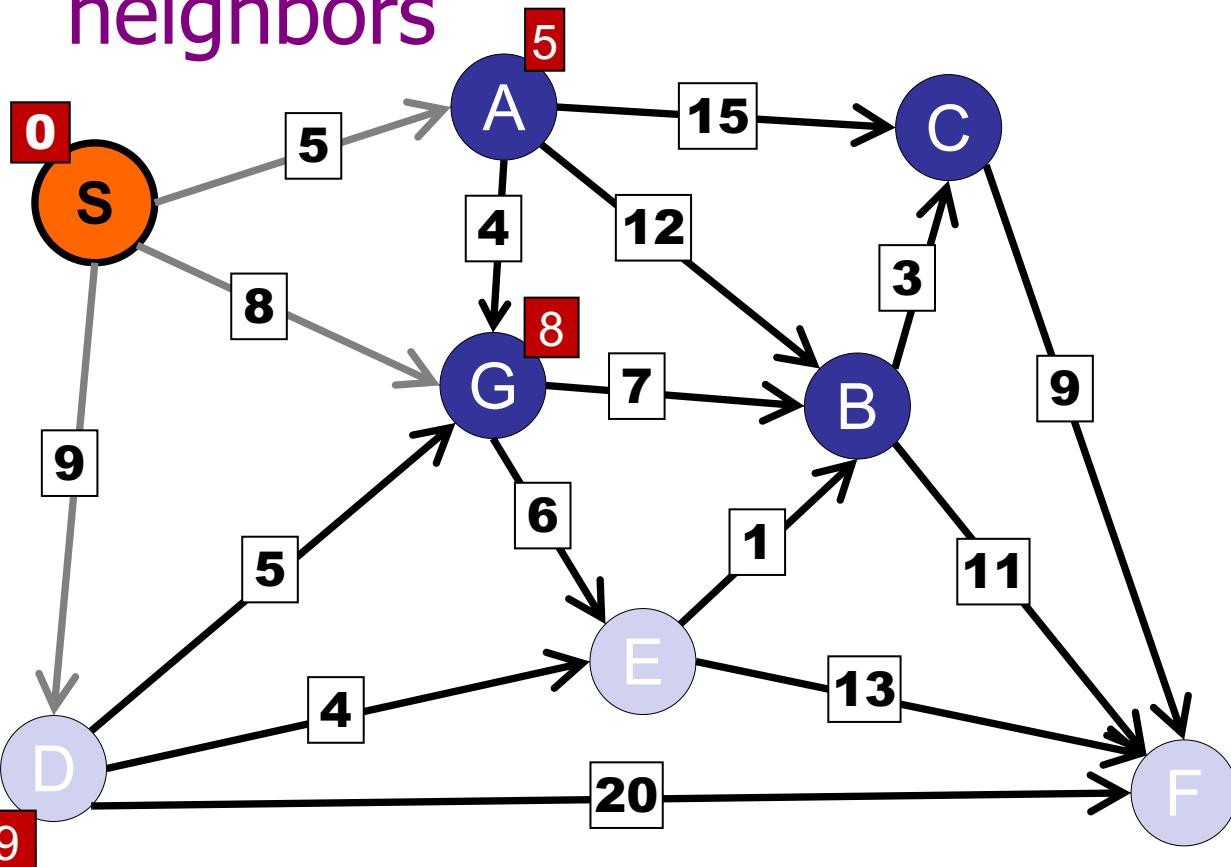


Vertex	Dist.
S	0

(Not showing vertices with distance = infinity)

# Dijkstra's Algorithm

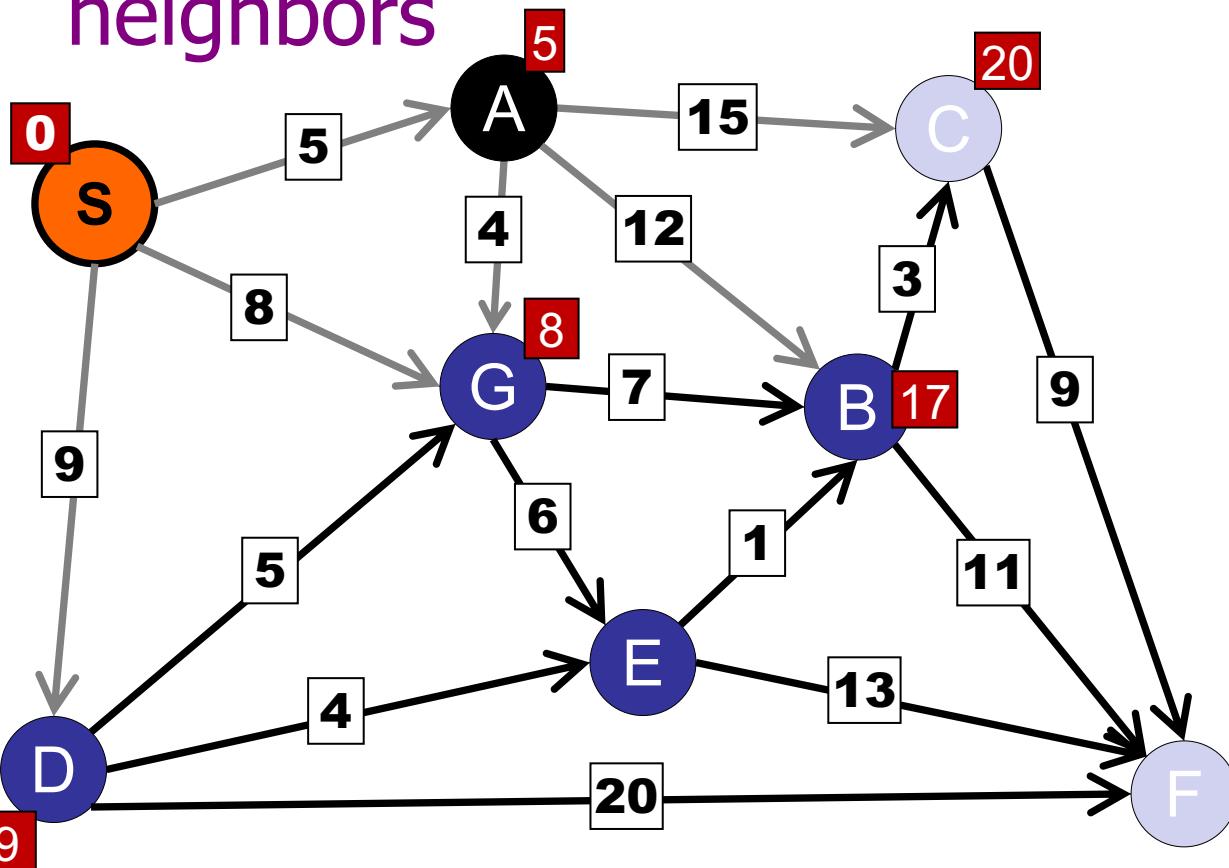
Extract A and relax/update neighbors



Vertex	Dist.
A	5
G	8
D	9

# Dijkstra's Algorithm

Extract G and relax/update neighbors

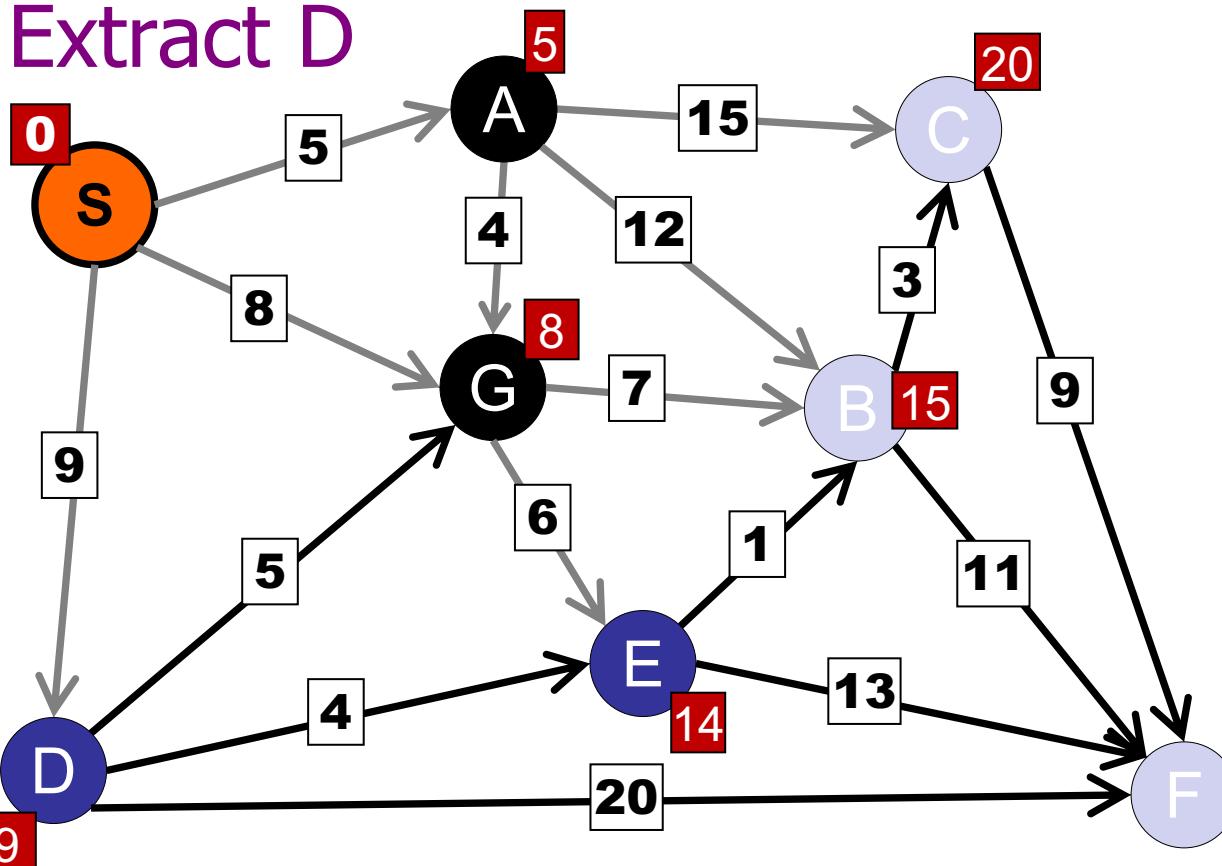


Vertex	Dist.
<b>G</b>	8
<b>D</b>	9
<b>B</b>	17
<b>C</b>	20

# Dijkstra's Algorithm

Dist. of B updated to 15

Extract D

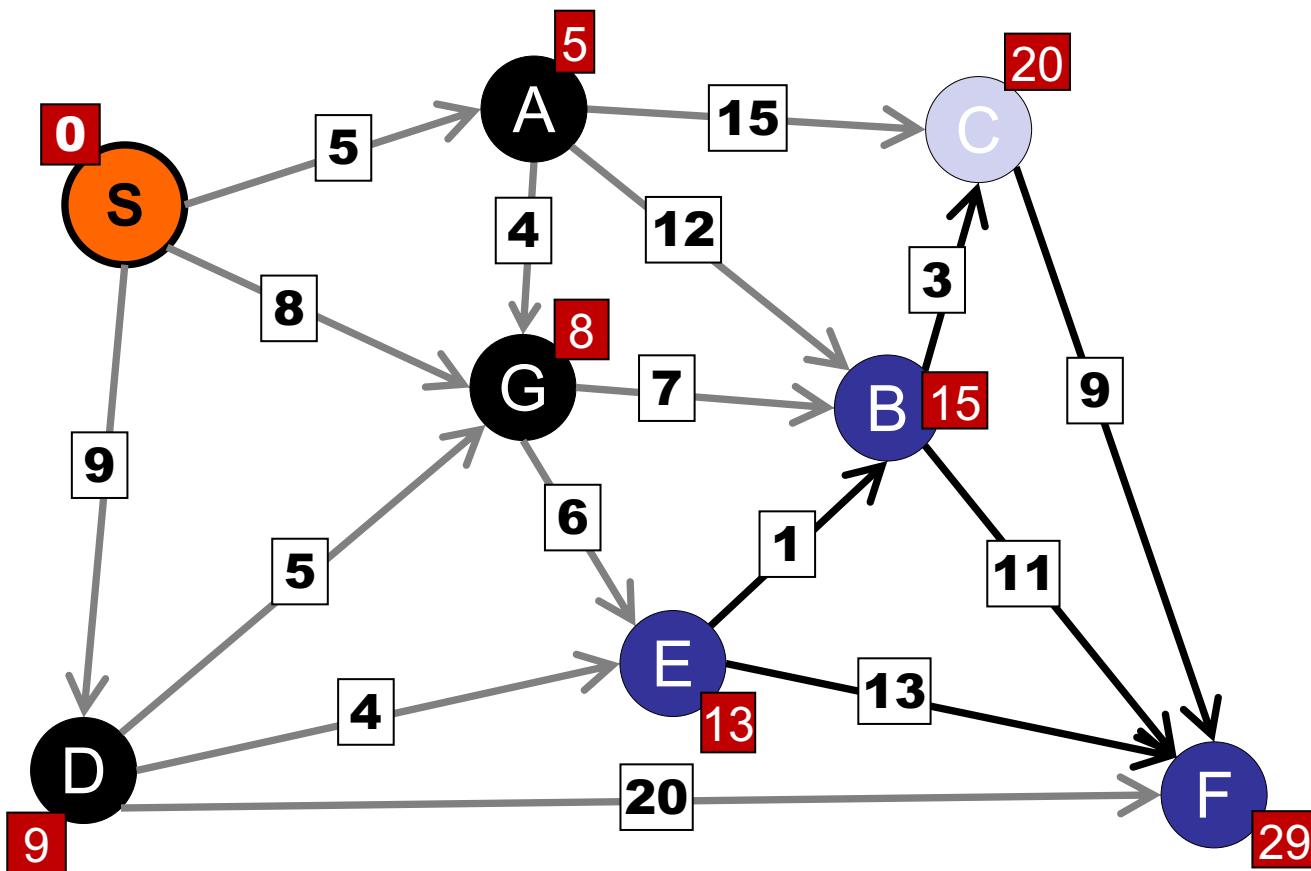


Vertex	Dist.
D	9
E	14
B	15
C	20

G is a neighbor of D. However, since G is already "dequeued", G won't be added back to the PQ anymore

# Dijkstra's Algorithm

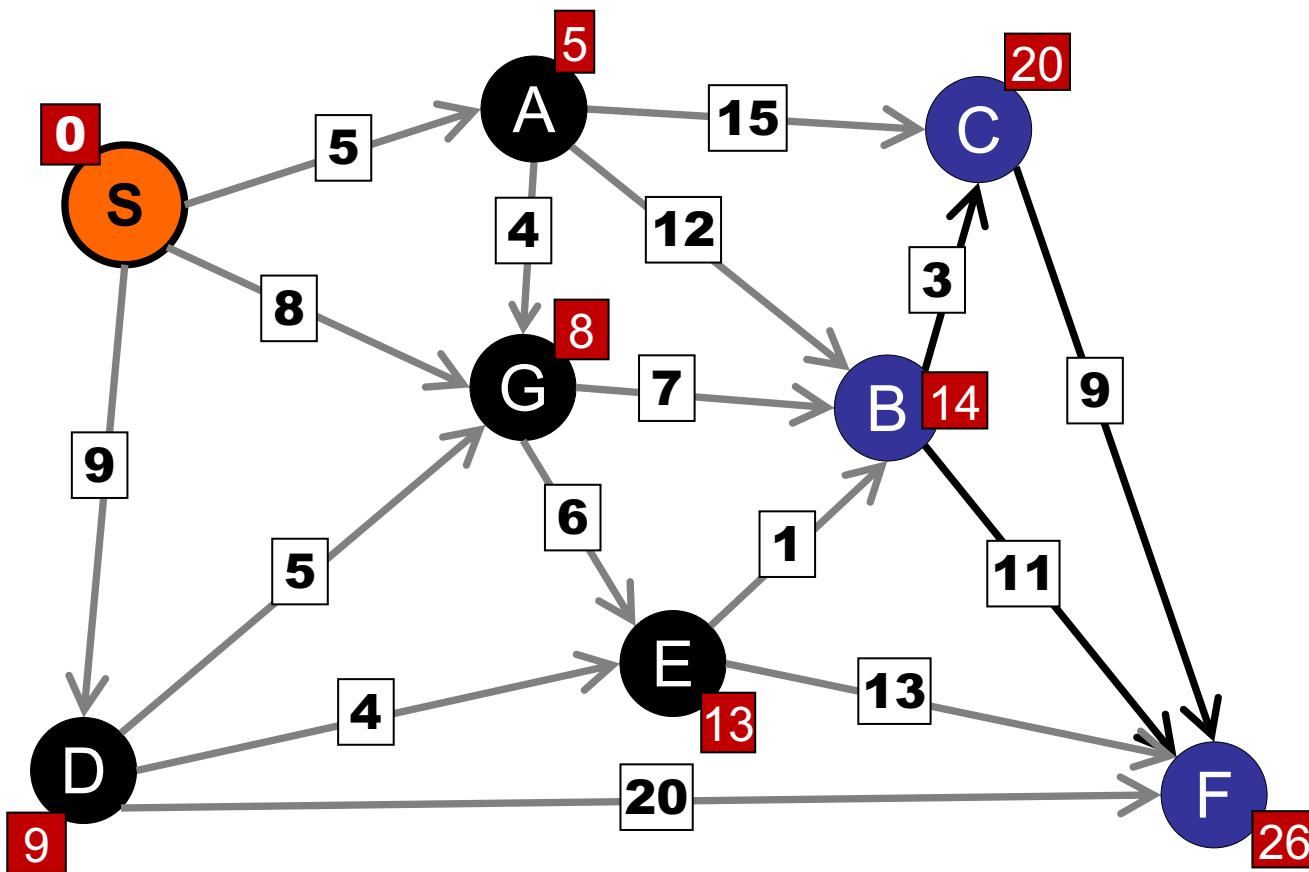
Extract E



Vertex	Dist.
E	13
B	15
C	20
F	29

# Dijkstra's Algorithm

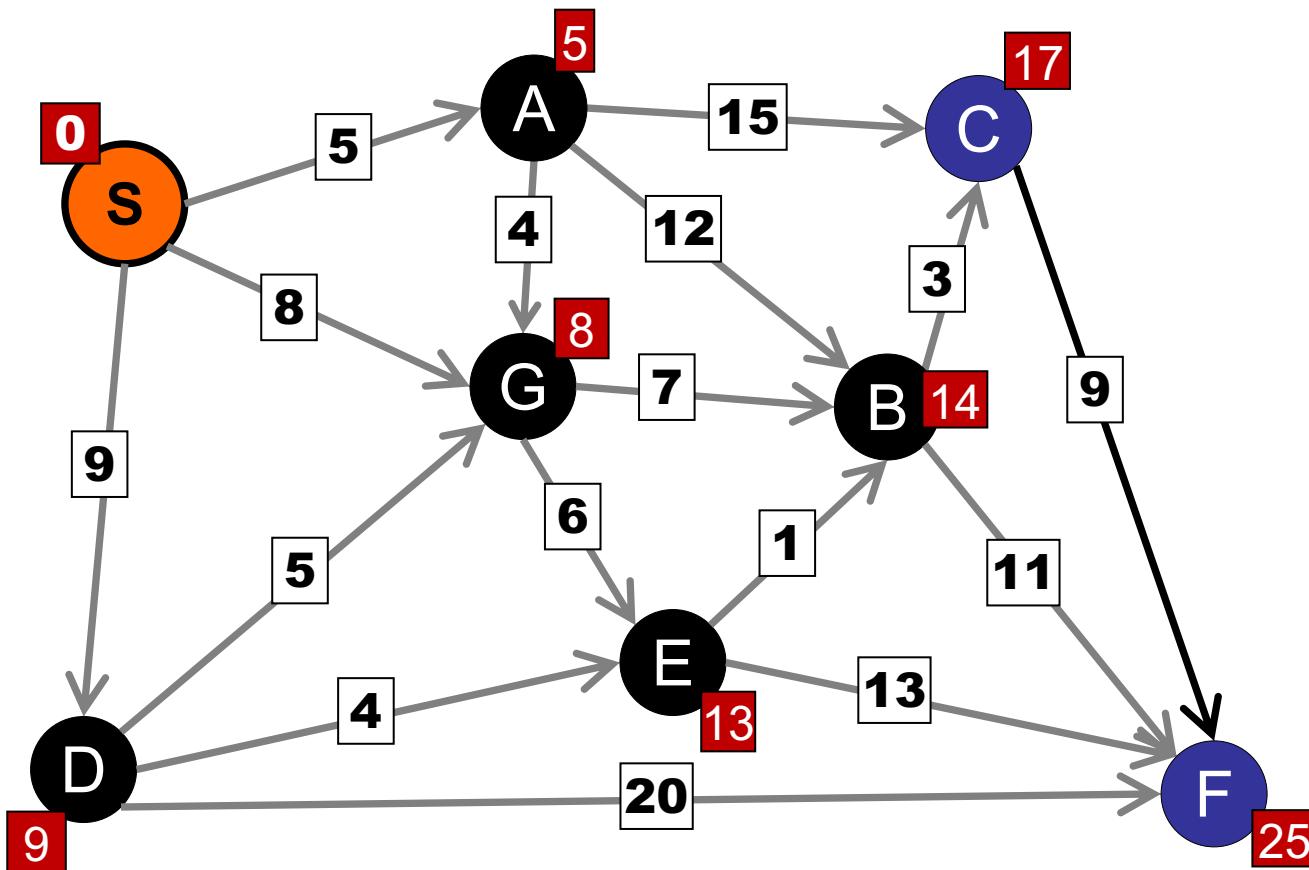
Extract B



Vertex	Dist.
B	14
C	20
F	26

# Dijkstra's Algorithm

Extract C

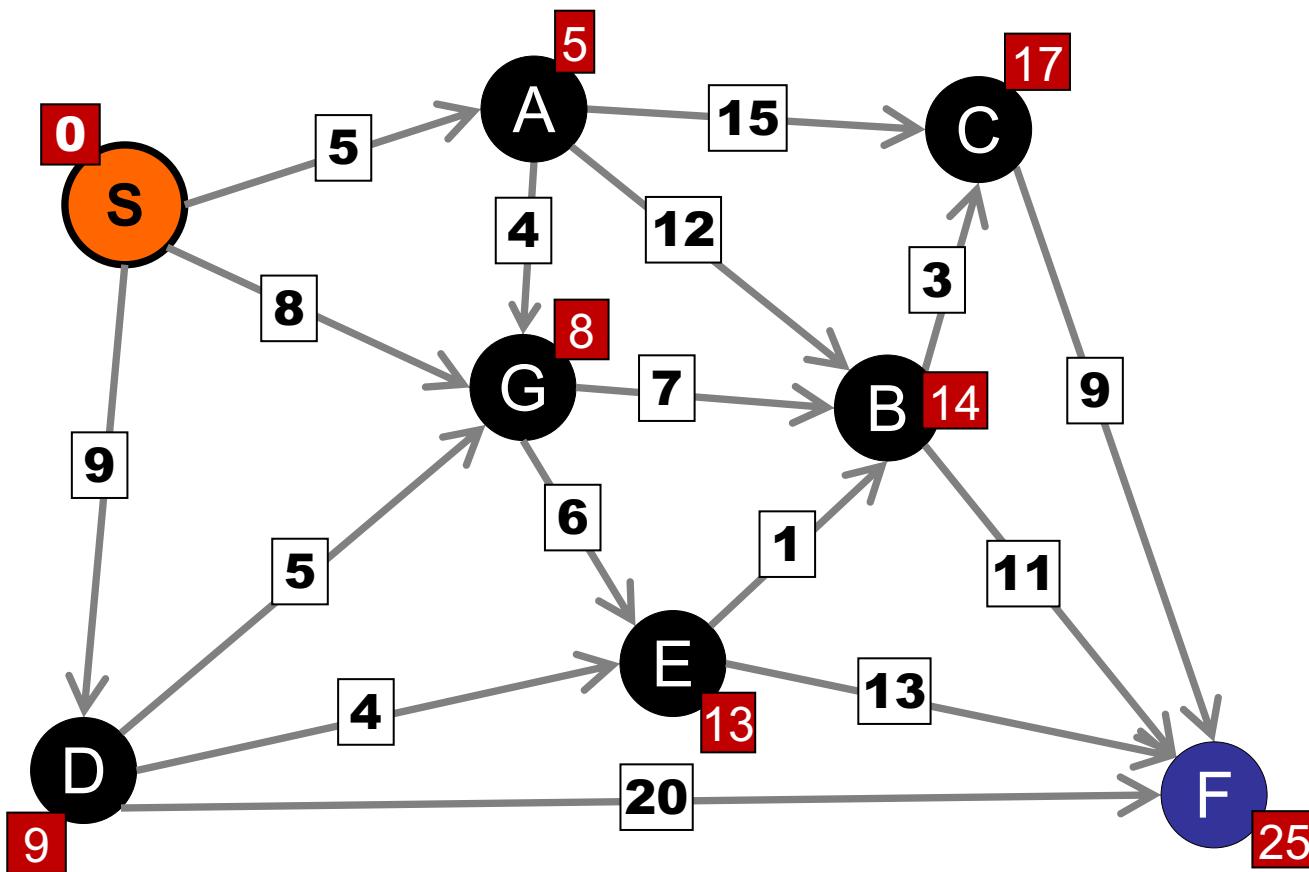


Vertex	Dist.
C	20
F	25

# Dijkstra's Algorithm

Extract F

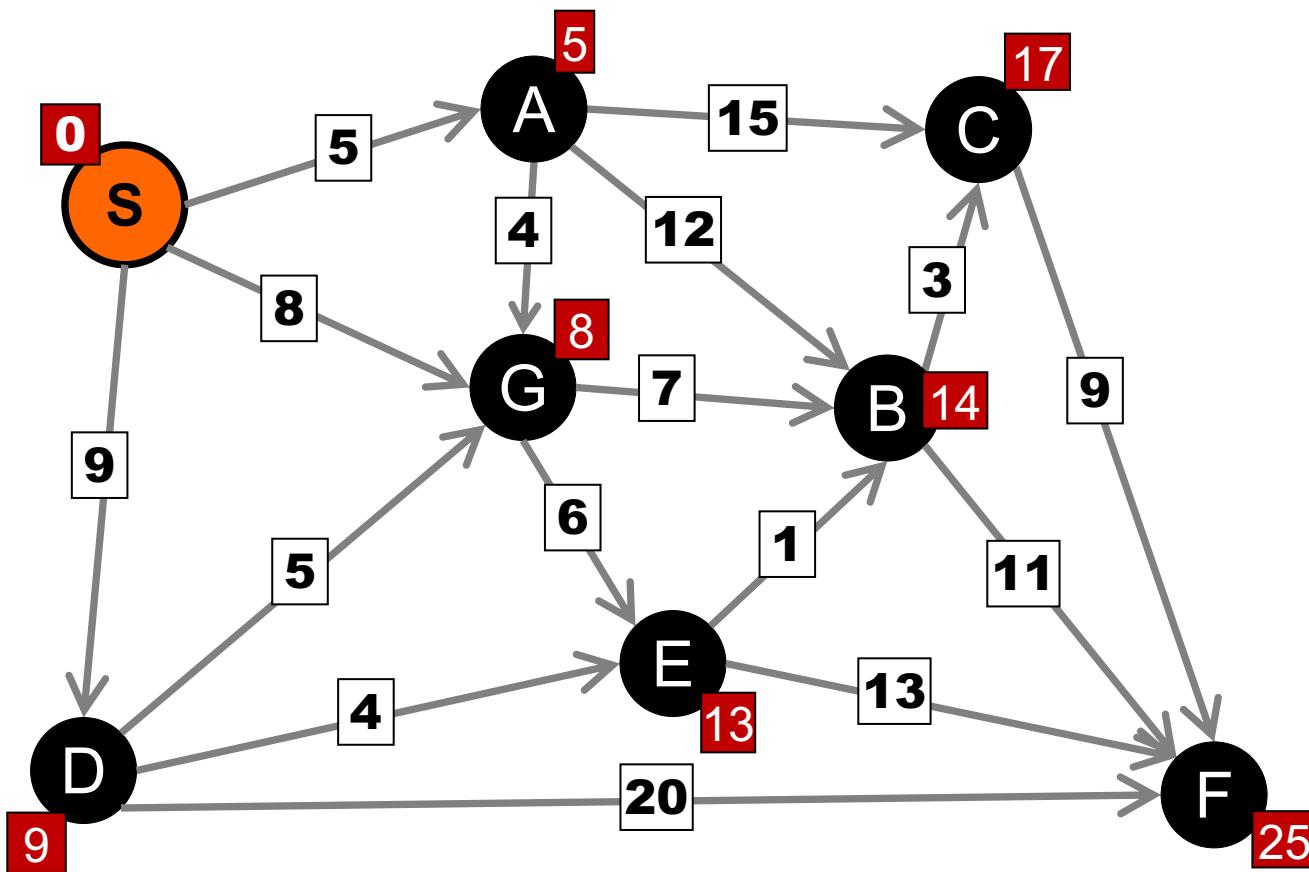
Vertex	Dist.
F	25



# Dijkstra's Algorithm

Done!

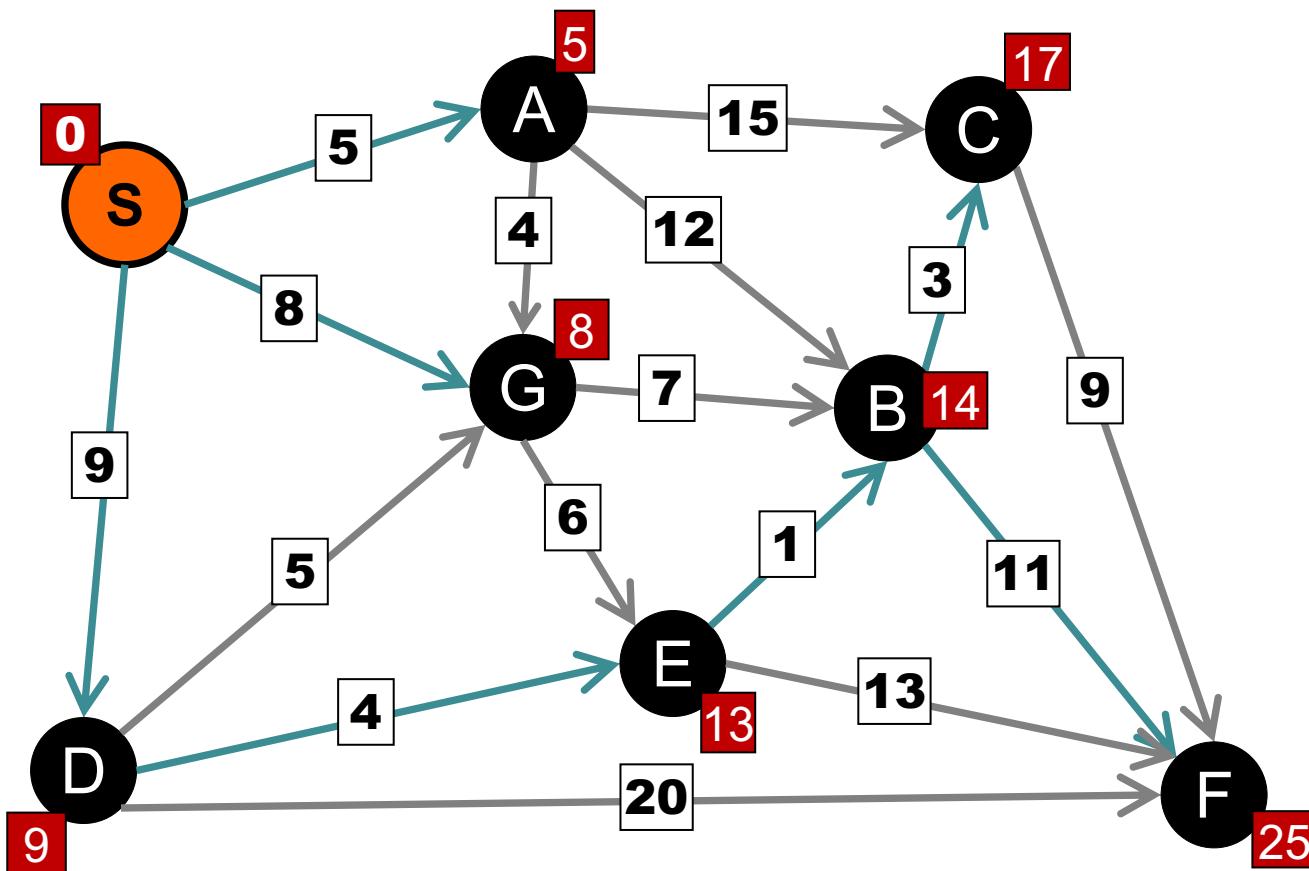
Vertex	Dist.



# Dijkstra's Algorithm

Done

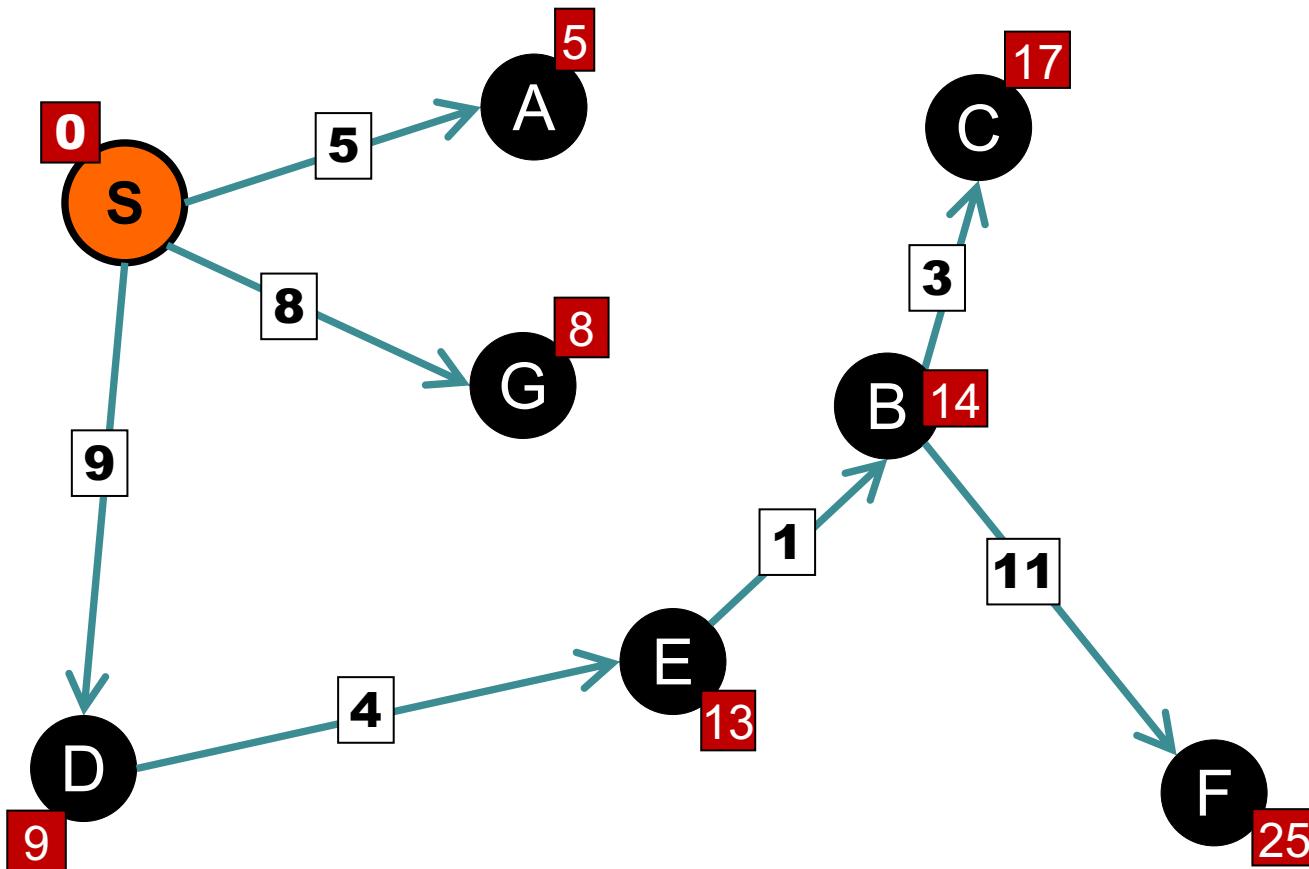
Vertex	Dist.



# Dijkstra's Algorithm

## Shortest Path Tree

Vertex	Dist.



# Abstract Data Type

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## Priority Queue

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void insert(Key k, Priority p)

*insert k with priority p*

Data extractMin()

*remove key with minimum priority*

void decreaseKey(Key k, Priority p)

*reduce the priority of key k to priority p*

boolean contains(Key k)

*does the priority queue contain key k?*

boolean isEmpty()

*is the priority queue empty?*

### Notes:

Assume data items are unique.

# Dijkstra's Algorithm

---

```
relax(Edge e) {  
    int v = e.from();  
    int w = e.to();  
    double weight = e.weight();  
    if (distTo[w] > distTo[v] + weight) {  
        distTo[w] = distTo[v] + weight;  
        parent[w] = v;  
        pq.decreaseKey(w, distTo[w]);  
    }  
}
```

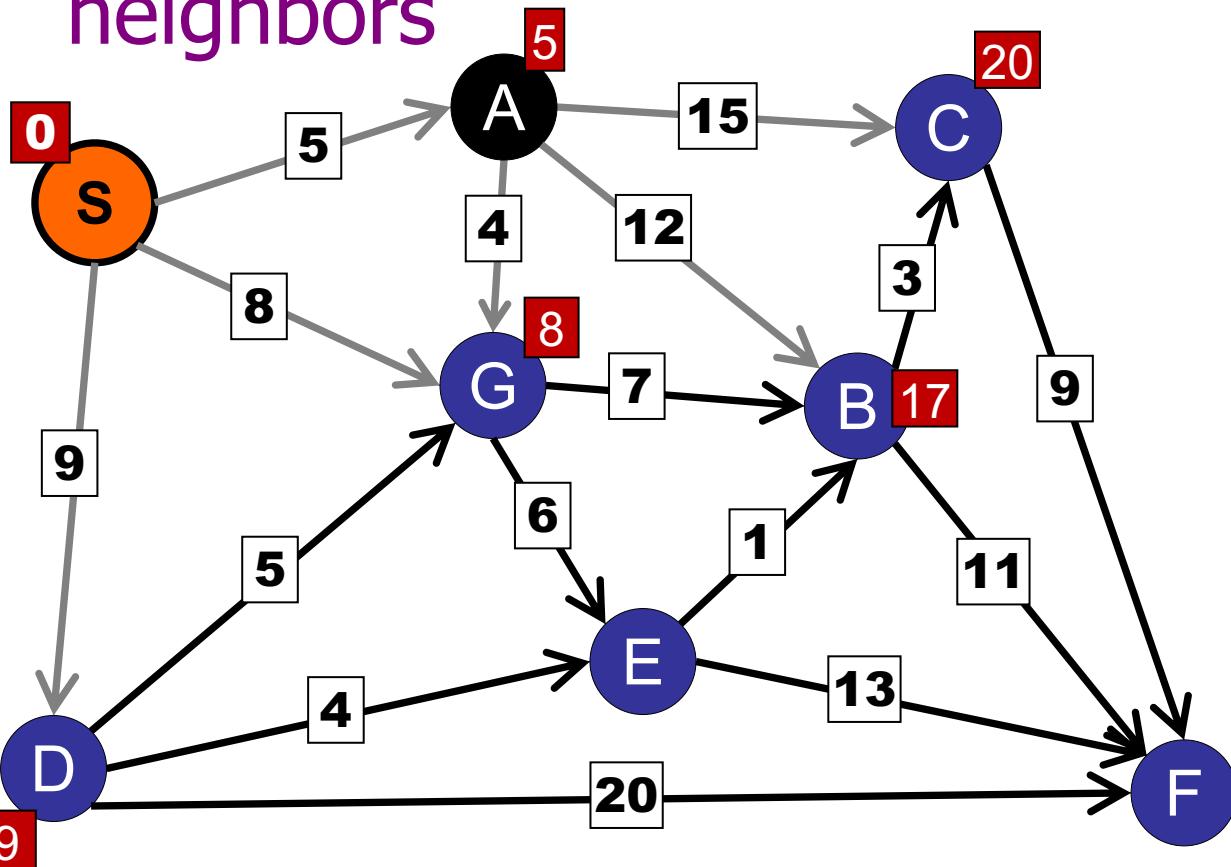
# Dijkstra's Algorithm

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        parent[w] = v;  
        pq.decreaseKey(w, distTo[w]);  
    }  
}
```

# Dijkstra's Algorithm

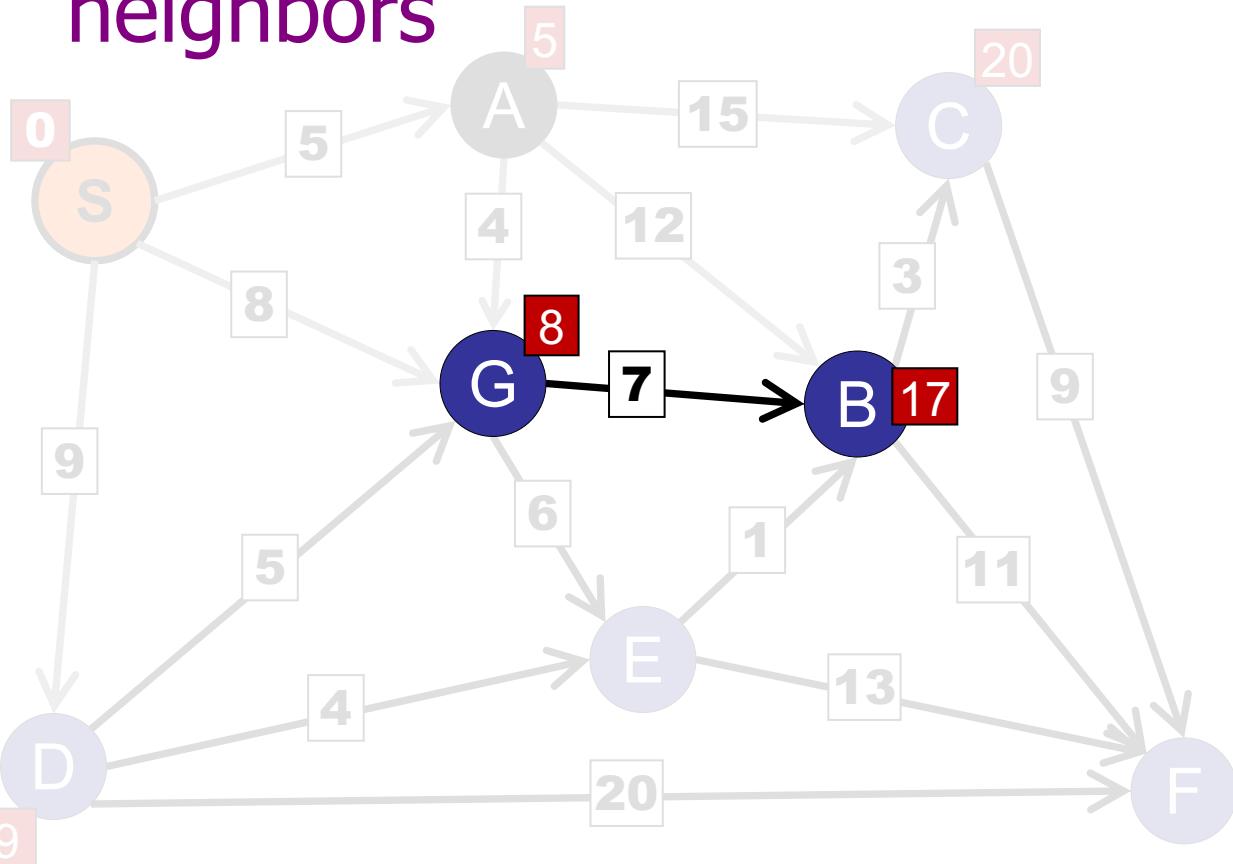
Remove G and relax/update neighbors



Vertex	Dist.
G	8
D	9
B	17
C	20

# Dijkstra's Algorithm

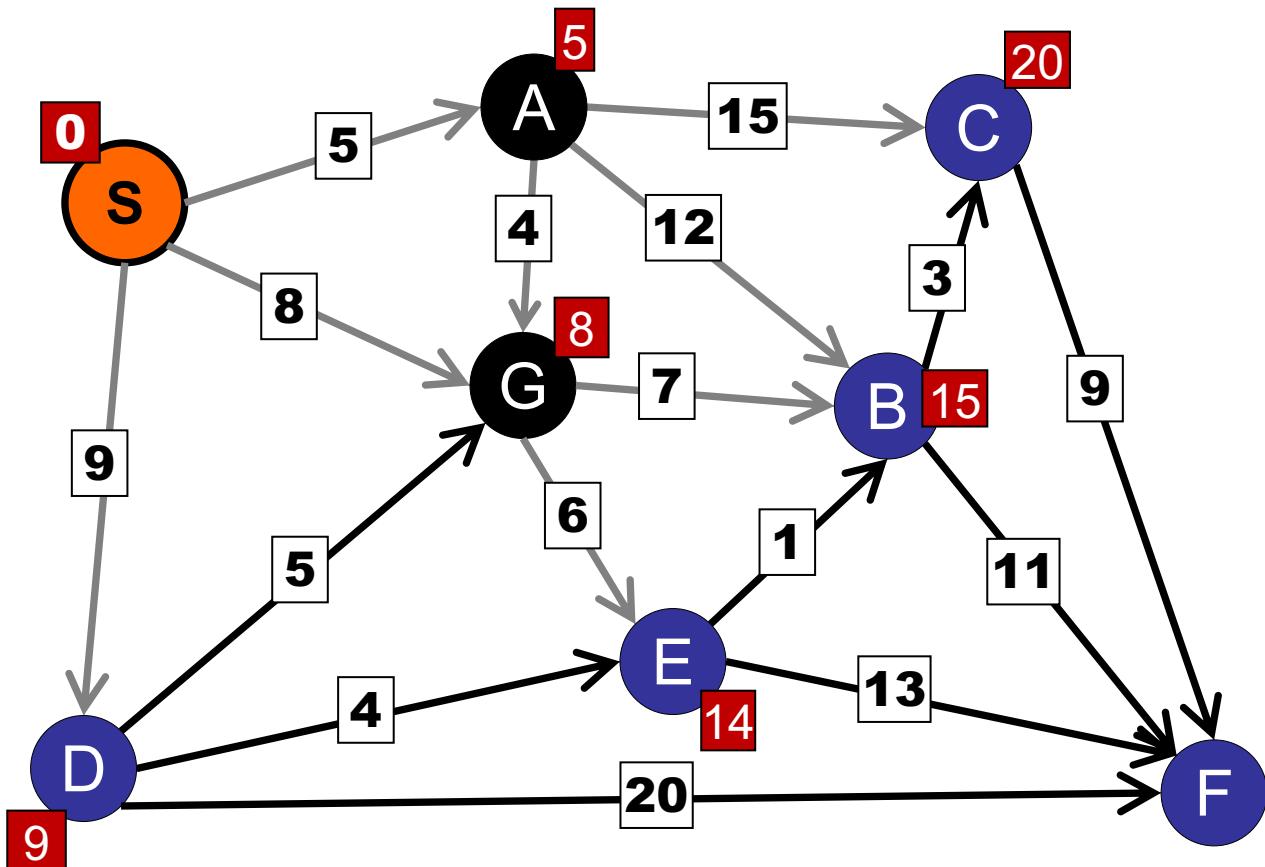
Remove G and relax/update neighbors



Vertex	Dist.
G	8
D	9
B	17
C	20

# Dijkstra's Algorithm

Remove G and relax.



Vertex	Dist.
D	9
E	14
B	15
C	20

# Dijkstra's Algorithm

---

```
relax(Edge e) {  
    int v = e.from();  
    int w = e.to();  
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    }  
}
```

# Dijkstra's Algorithm

---

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    if (distTo[w] > distTo[v] + weight) {  
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        pq.decreaseKey(w, distTo[w]);  
    }  
}
```

# Dijkstra's Algorithm

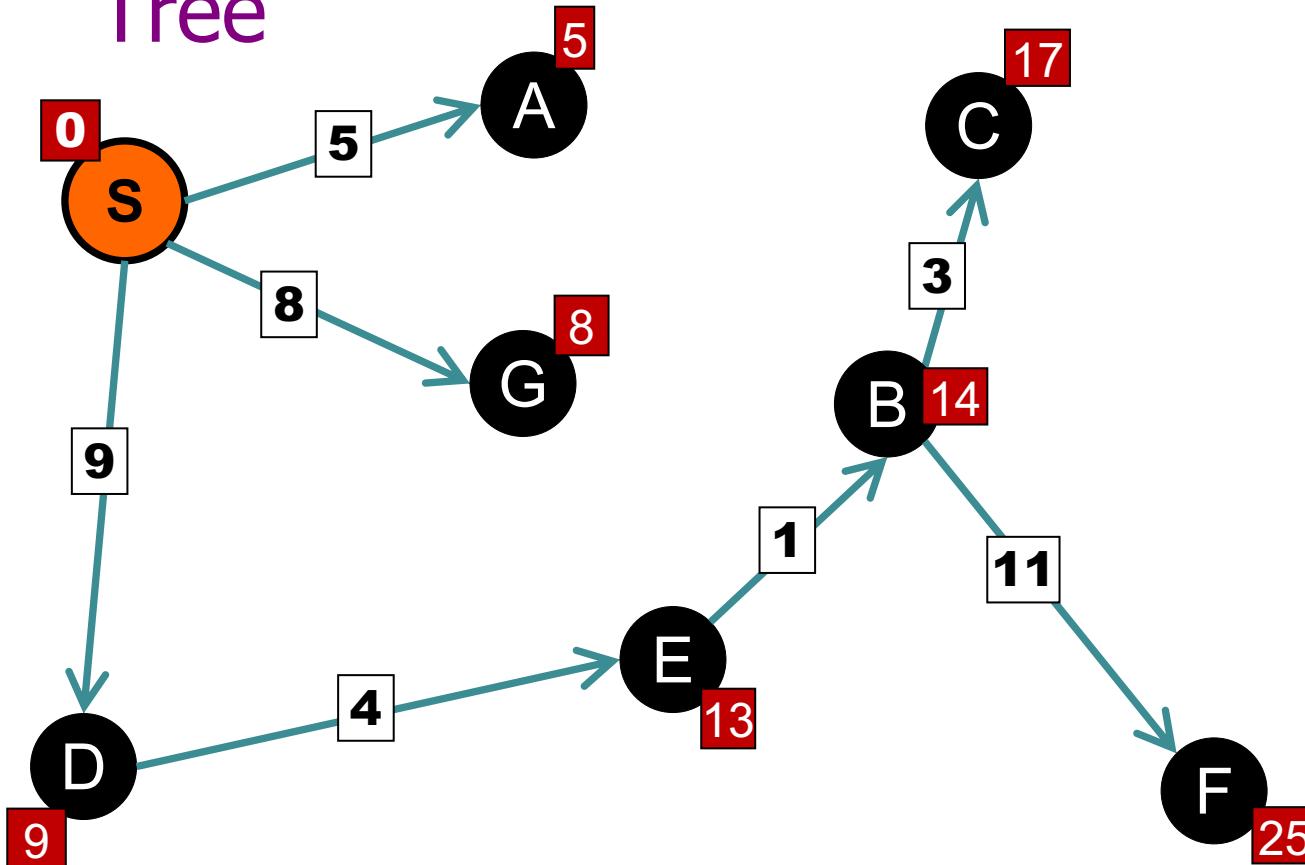
---

## Analysis:

- `deleteMin`:  $|V|$  times each
  - Each node is added to the priority queue **once**.
- `relax / decreaseKey`:  $|E|$  times
  - Each edge is relaxed once.
- Priority queue operations:  $O(\log V)$
- Total:  $O((V+E)\log V) = O(E \log V)$

# Dijkstra's Algorithm

Following the parents: Yields the Shortest Path Tree



# Dijkstra's Algorithm

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Why does it work?

# Dijkstra's Algorithm

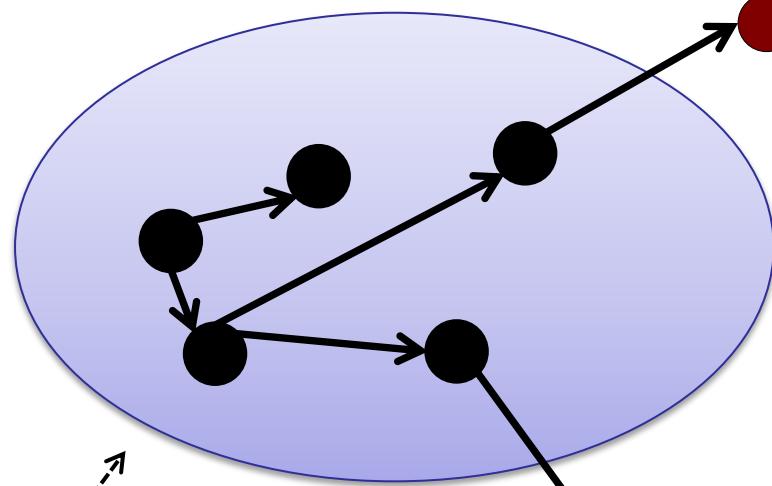
---

Proof by induction:

- Every “finished” (dequeued) vertex has a correct estimate.
  - Namely, shortest path is found for that vertex
- Initially: only “finished” vertex is start.

# Dijkstra's Algorithm

Every edge crossing the boundary **has been relaxed**.



**finished vertices:**  
distance is accurate.

**fringe vertices:** **top in priority queue**  
neighbor of a finished vertex.

**fringe vertices:**  
neighbor of a  
finished vertex.

**other vertices:**  
no known  
estimate

# Dijkstra's Algorithm

---

Proof by induction:

- Every “finished” vertex has correct estimate.
- Initially: only “finished” vertex is start.

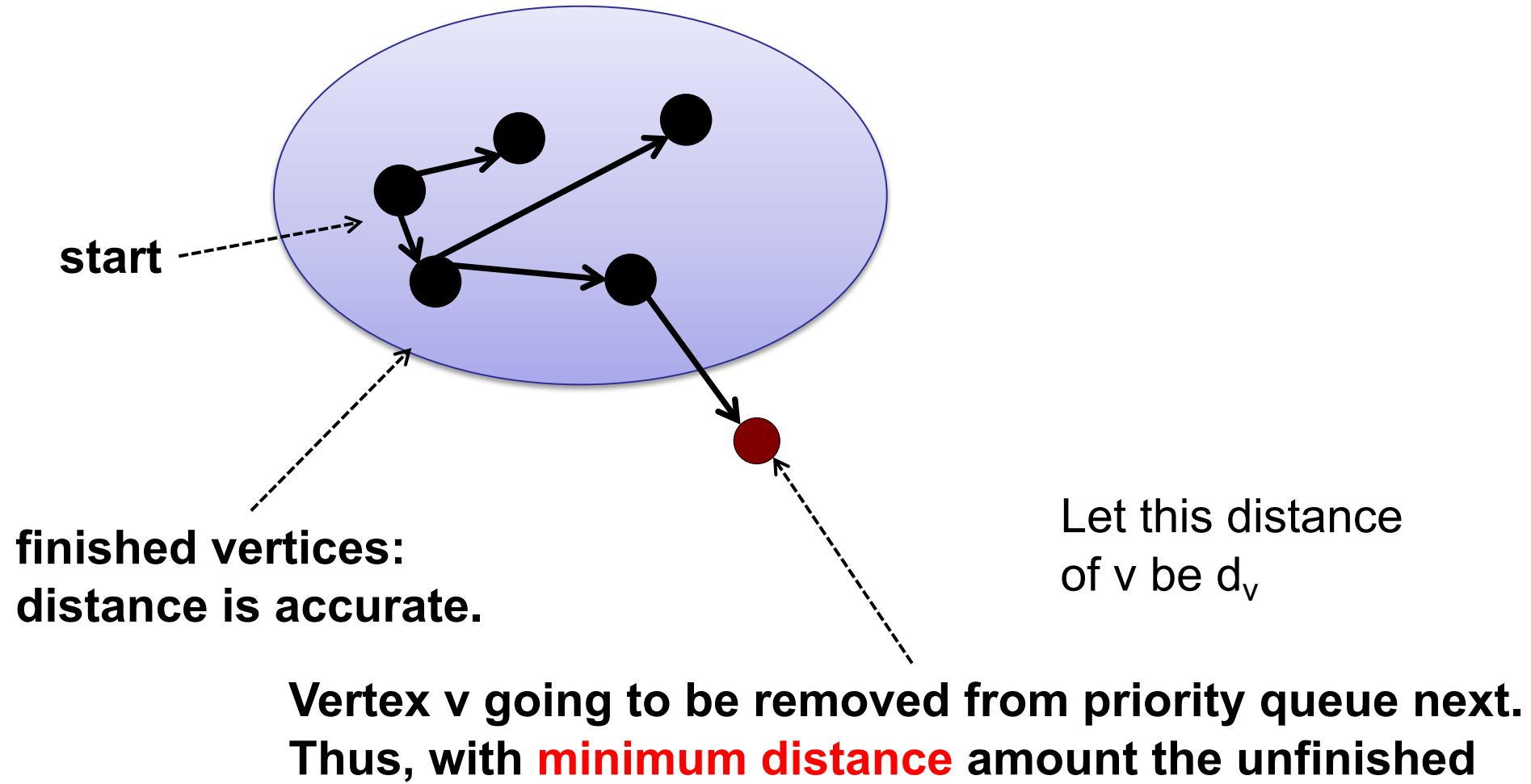
# Dijkstra's Algorithm

---

Proof by induction:

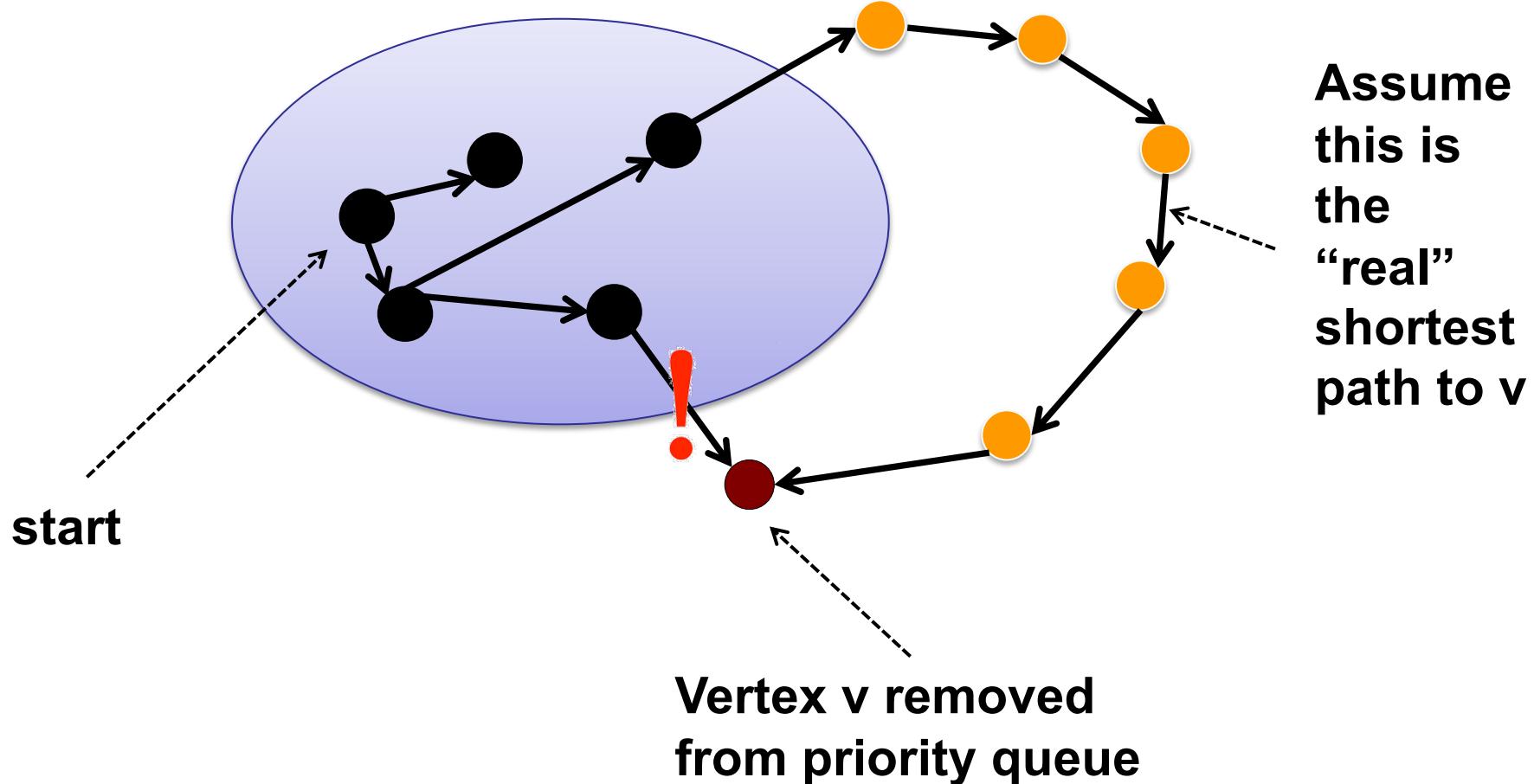
- Every “finished” vertex has correct estimate.
- Initially: only “finished” vertex is start.
- Inductive step:
  - Remove vertex from priority queue.
  - Relax its edges.
  - Add it to finished.
  - **Claim: it has a correct estimate.**

# Dijkstra's Algorithm



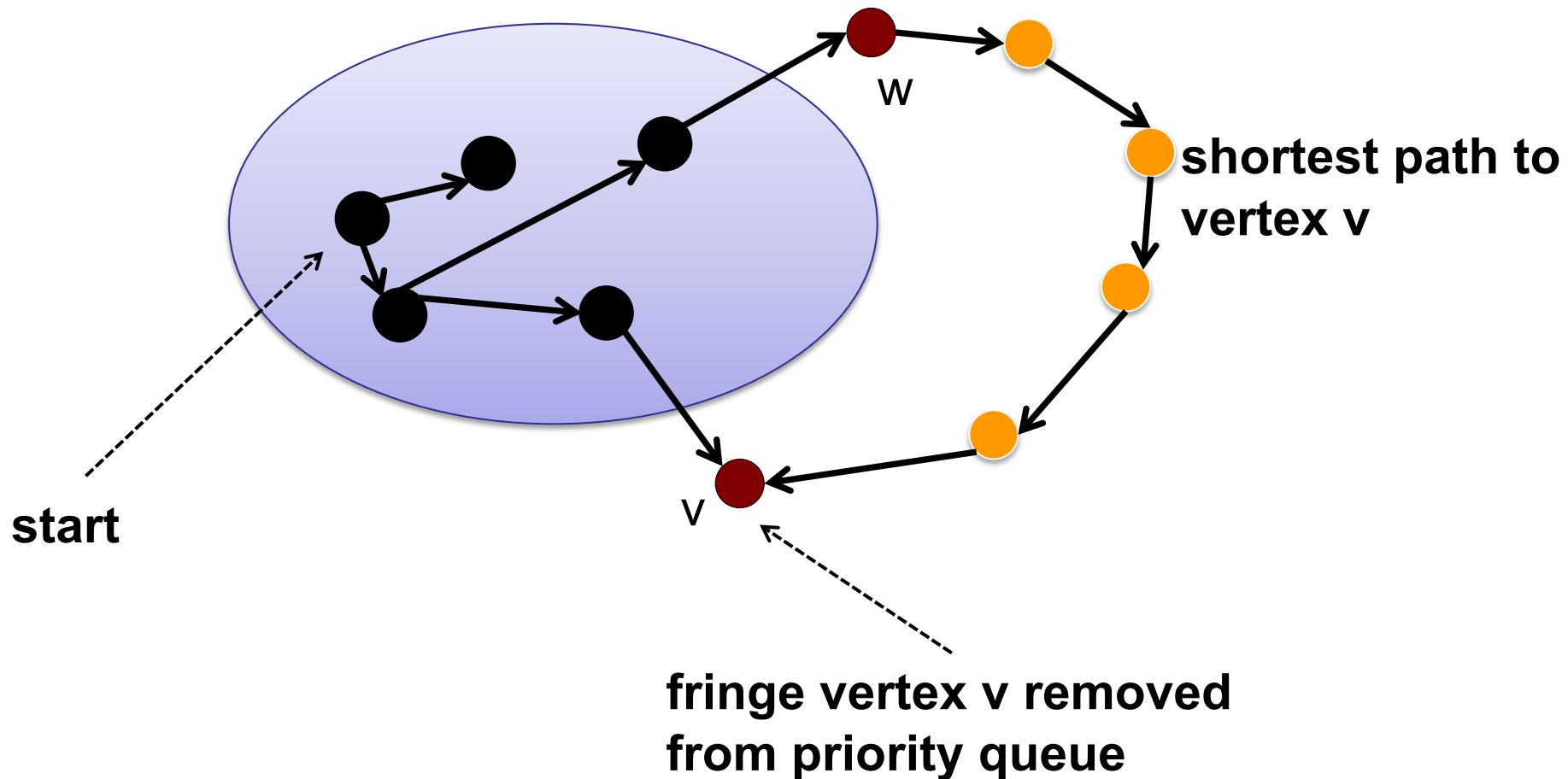
# Dijkstra's Algorithm

Assume NOT. The current estimate is not the shortest path. And the new path has dist.  $< d_v$



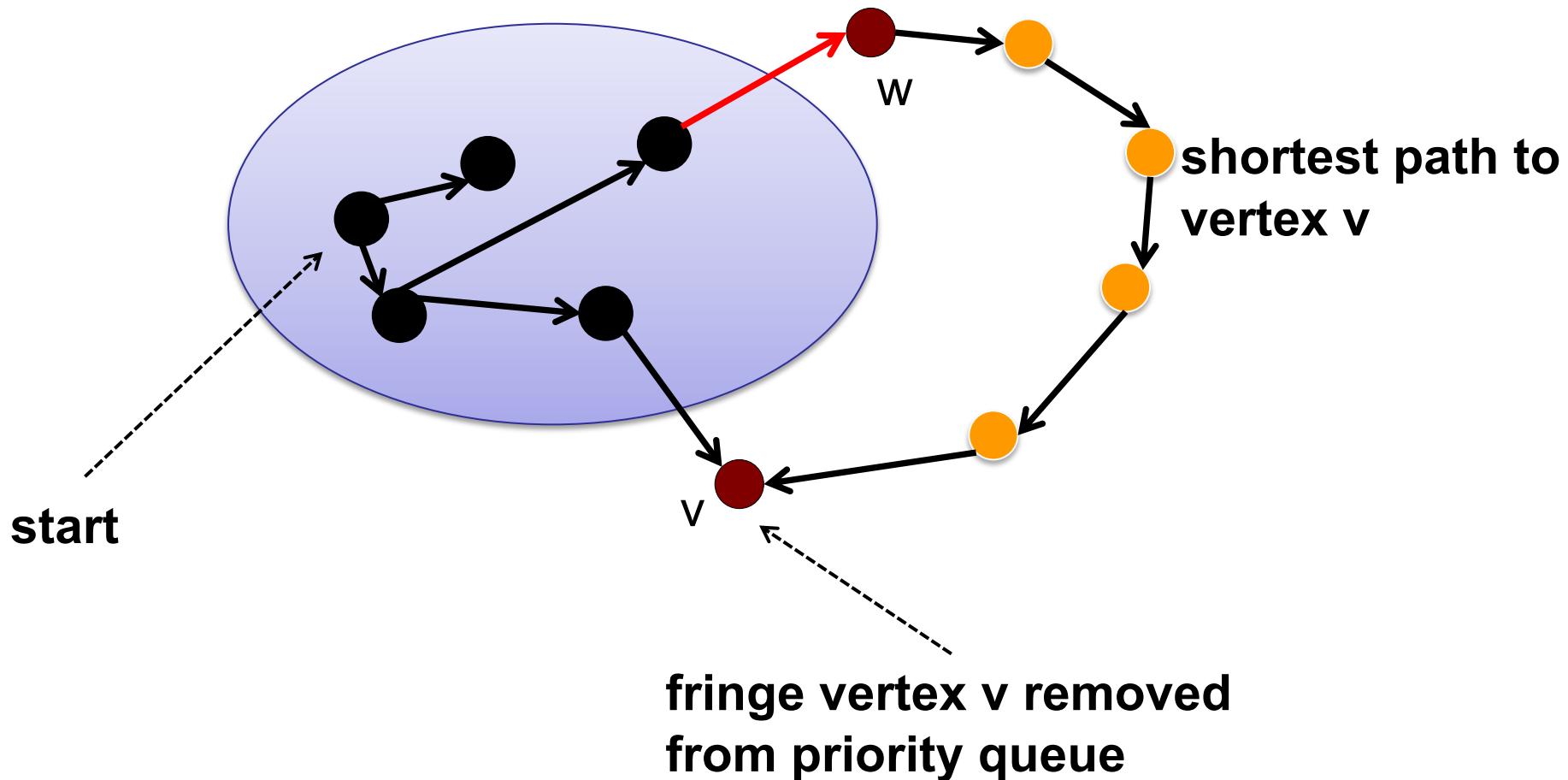
# Dijkstra's Algorithm

There must be a vertex  $w$  in the current PQ on this “real” path.  
And let this “real” path has distance  $r_v < d_v$



# Dijkstra's Algorithm

If  $P$  is shortest path to  $v$ , then prefix of  $P$  is shortest path to  $w$ .  
Then  $\text{distTo}[w]$  is accurate. And  $\text{distTo}[w] < r_v < d_v = \text{distTo}[v]$

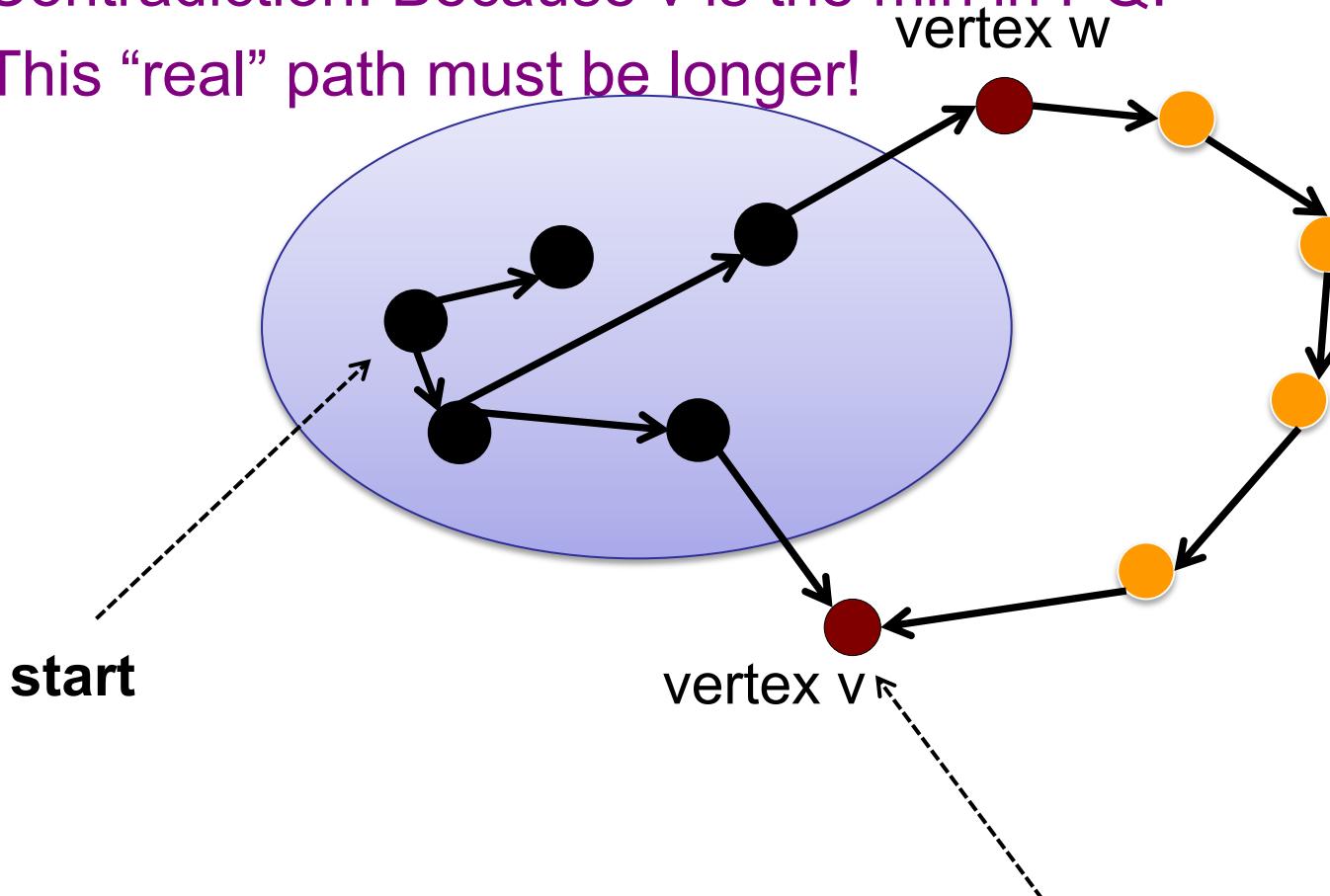


# Dijkstra's Algorithm

But  $\text{distTo}[w] \geq \text{distTo}[v]$  according to PQ!

Contradiction! Because v is the min in PQ!

This “real” path must be longer!



**Vertex v going to be removed from priority queue next.**  
**Thus, with minimum distance amount the unfinished**

# Dijkstra's Algorithm

---

Proof by induction:

- Every “finished” vertex has correct estimate.
- Initially: only “finished” vertex is start.
- Inductive step:
  - Remove vertex from priority queue.
  - Relax its edges.
  - Add it to finished.
  - **Claim: it has a correct estimate.**

# Dijkstra's Algorithm

---

```
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    int v = e.from();  
    int w = e.to();  
    double weight = e.weight();  
    if (distTo[w] > distTo[v] + weight) {  
        distTo[w] = distTo[v] + weight;  
        parent[w] = v;  
        pq.decreaseKey(w, distTo[w]);  
    }  
}
```

# Dijkstra's Algorithm

---

## Analysis:

- `insert / deleteMin`:  $|V|$  times each
  - Each node is added to the priority queue **once**.
- `decreaseKey`:  $|E|$  times
  - Each edge is relaxed once.
- Priority queue operations:  $O(\log V)$
- Total:  $O((V+E)\log V) = O(E \log V)$

# Dijkstra's Algorithm

---

## Source-to-Destination:

- What if you stop the first time you dequeue the destination?
  
- Recall:
  - a vertex is “finished” when it is dequeued
  - if the destination is finished, then stop

# Dijkstra Summary

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Basic idea:

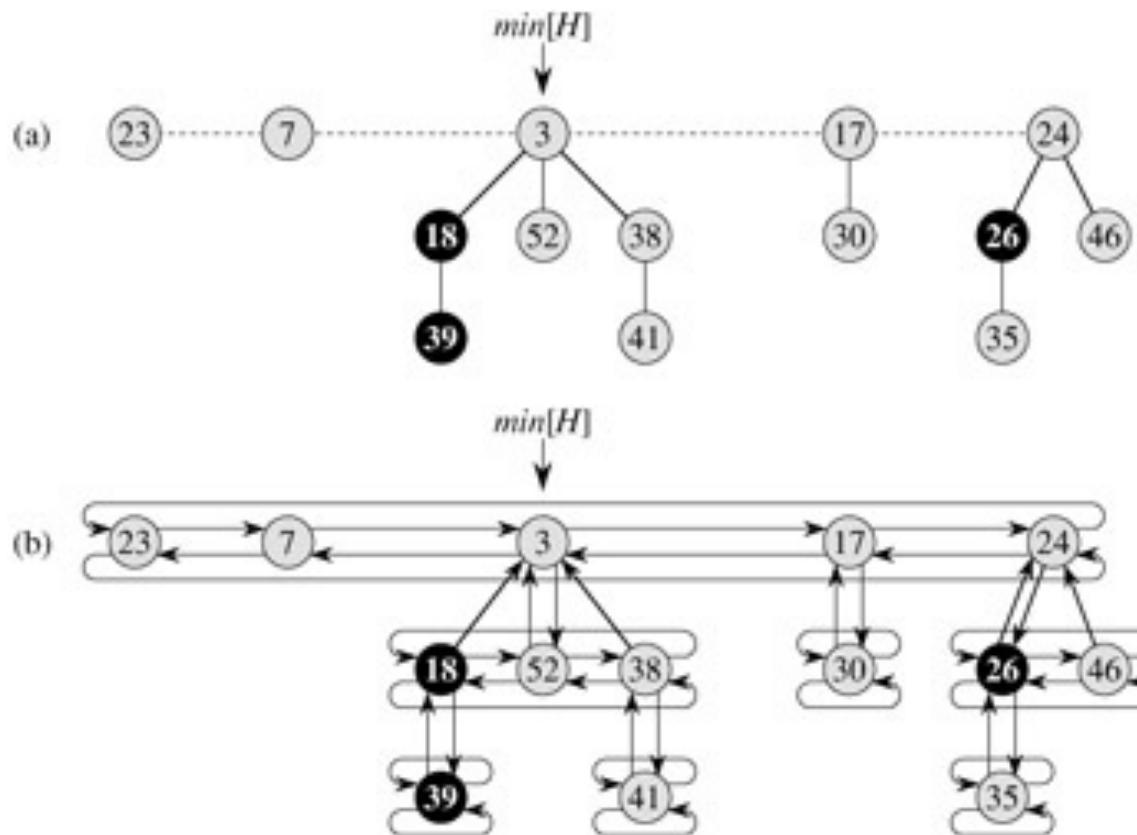
- Maintain distance estimates.
- Repeat:
  - Find unfinished vertex with smallest estimate.
  - Relax all outgoing edges.
  - Mark vertex finished.
- $O(E \log V)$  time

# Dijkstra's Performance

PQ Implementation	insert	deleteMin	decreaseKey	Total
Array	1	$V$	1	$O(V^2)$
AVL Tree	$\log V$	$\log V$	$\log V$	$O(E \log V)$
d-way Heap	$d\log_d V$	$d\log_d V$	$\log_d V$	$O(E \log_{E/V} V)$
Fibonacci Heap	1	$\log V$	1	$O(E + V \log V)$

# Fibonacci Heap

- Not in this course



# Dijkstra Summary

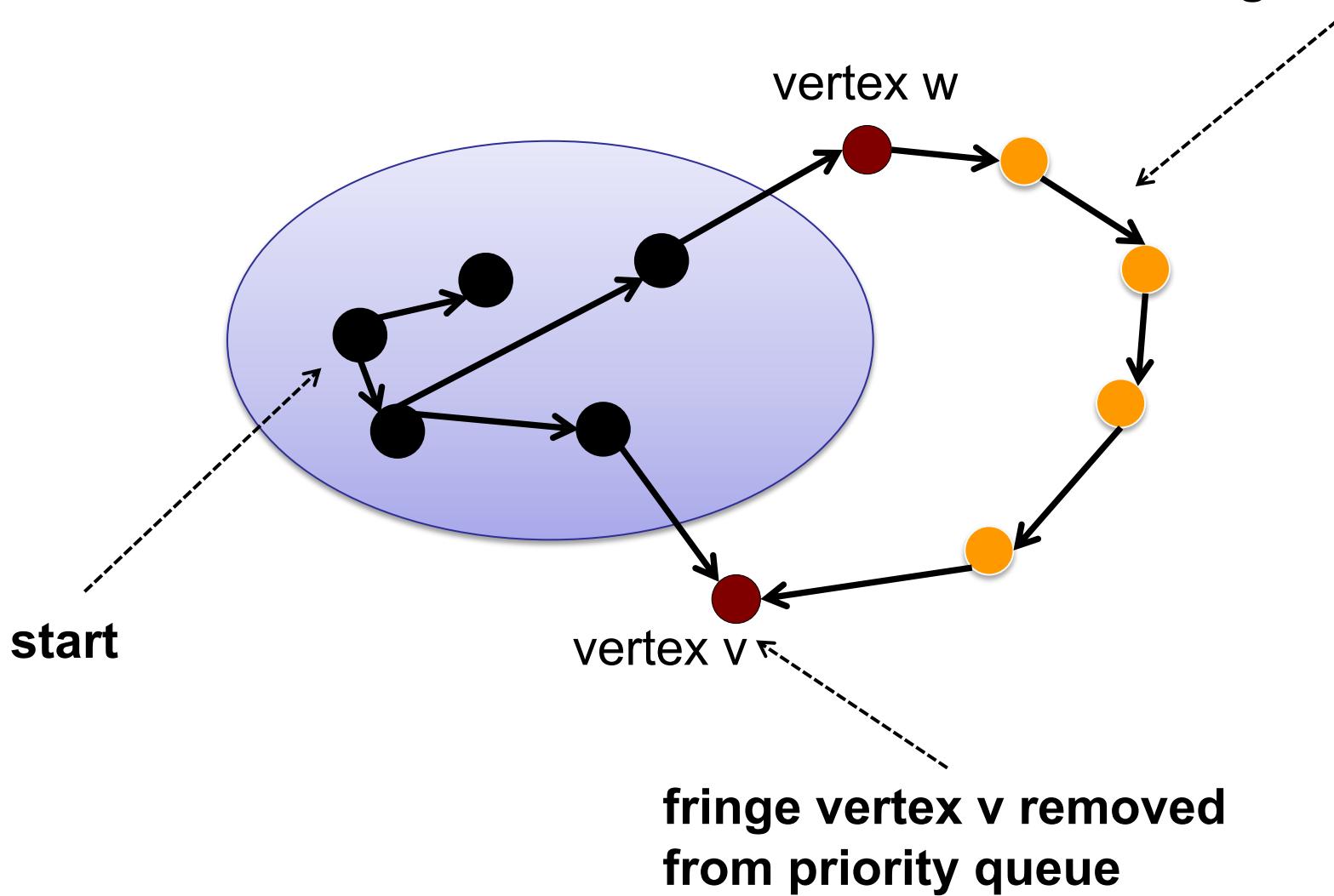
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Edges with negative weights?

# Dijkstra's Algorithm

What goes wrong with negative weights?

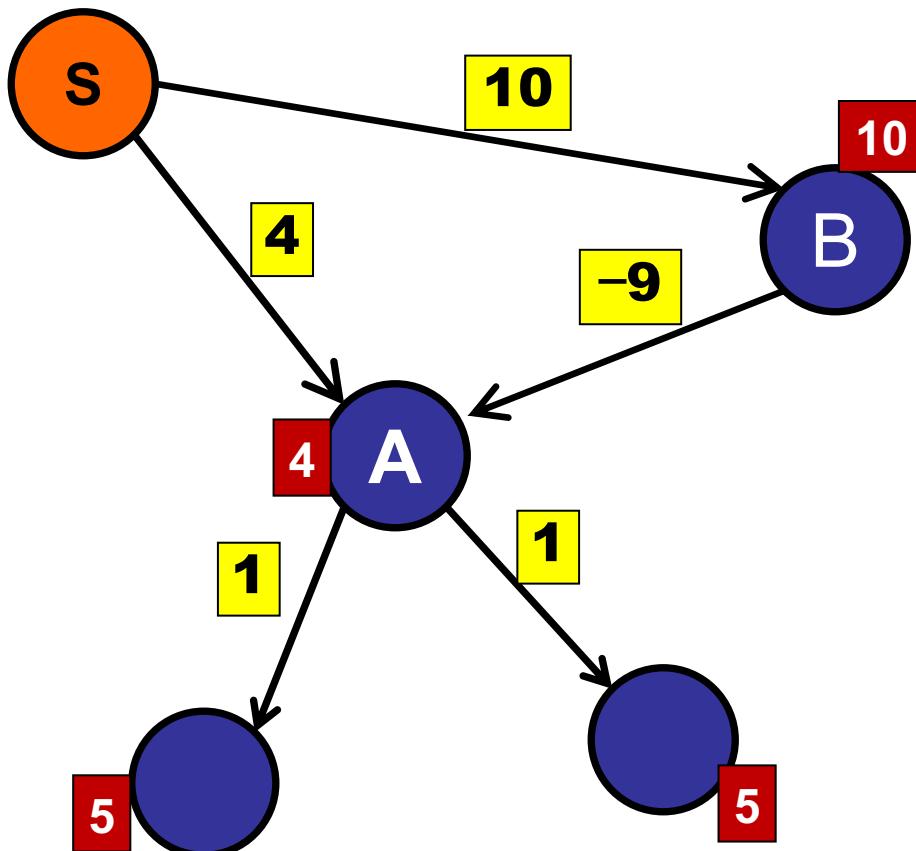
**shortest path to  
fringe vertex v**



**fringe vertex v removed  
from priority queue**

# Dijkstra's Algorithm

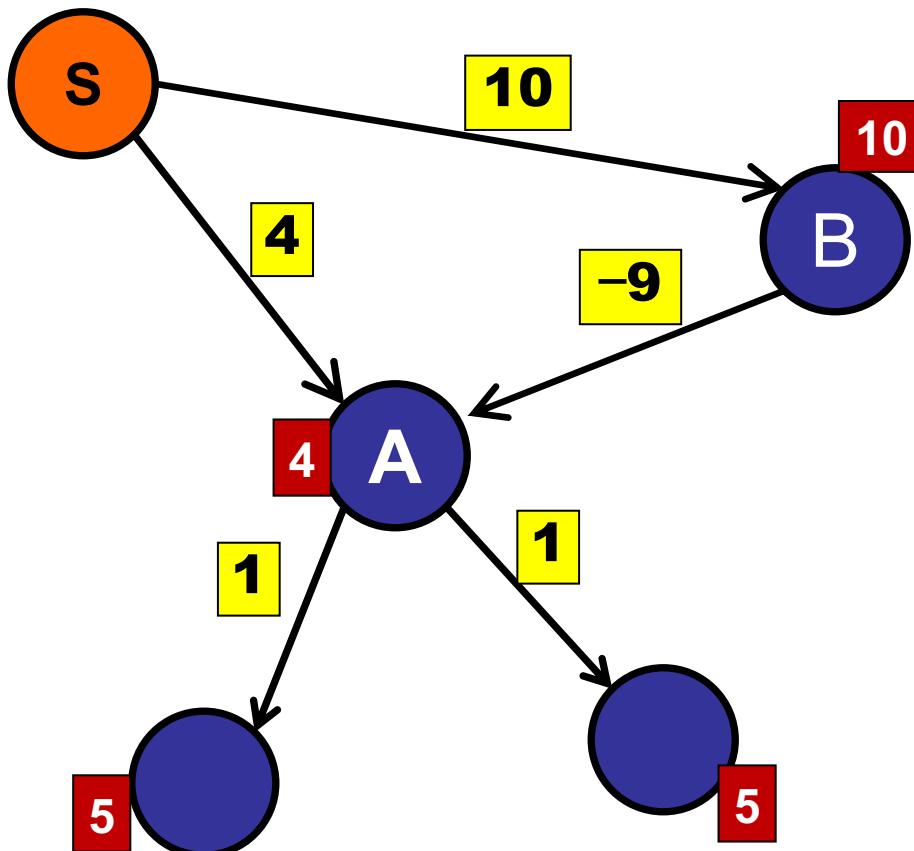
Edges with negative weights?



Step 1: Remove A.  
Relax A.  
Mark A done.

# Dijkstra's Algorithm

Edges with negative weights?



Step 1: Remove A.  
Relax A.  
Mark A done.

...

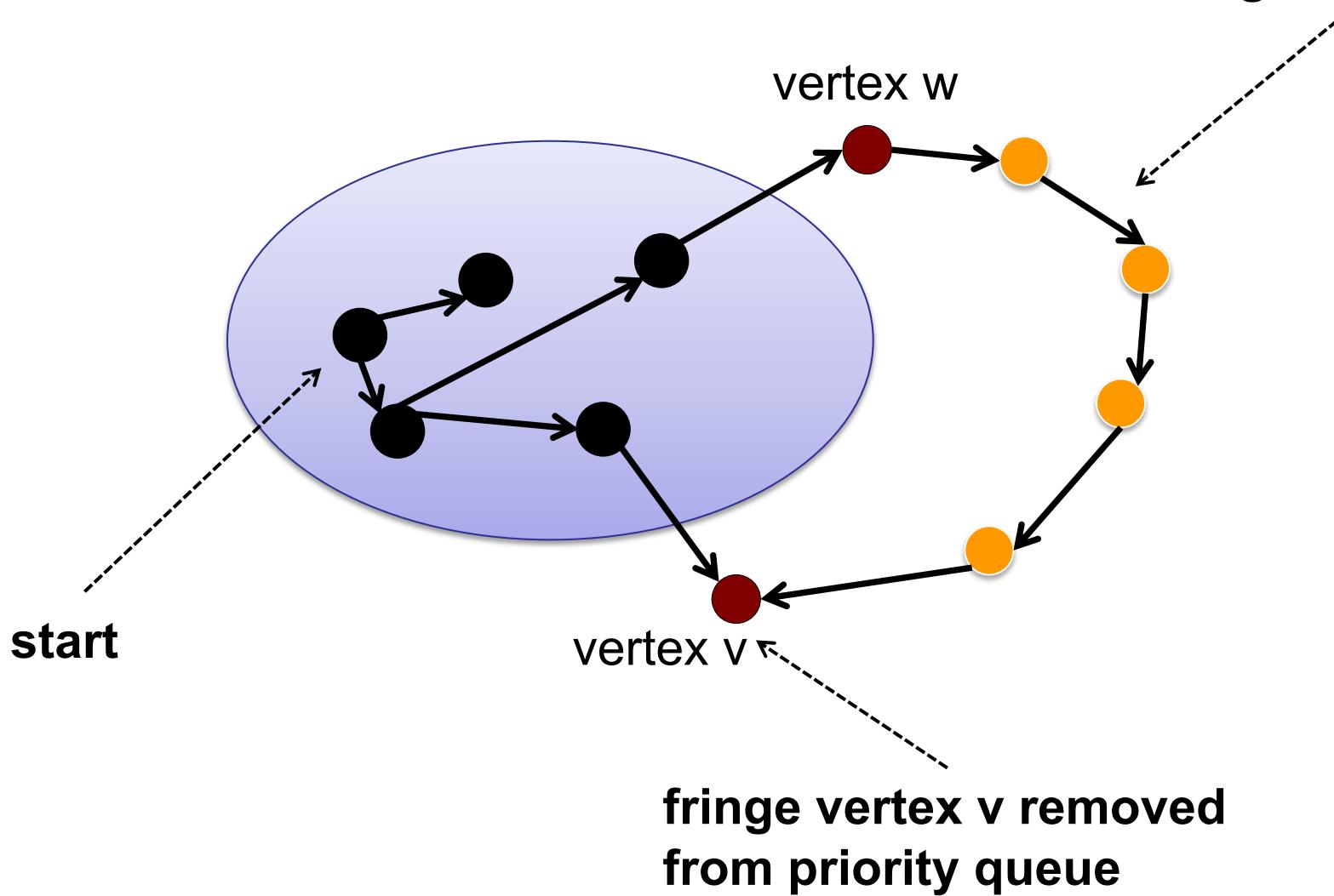
Step 4: Remove B.  
Relax B.  
Mark B done.

Oops: We need to update A.

# Dijkstra's Algorithm

What goes wrong with negative weights?

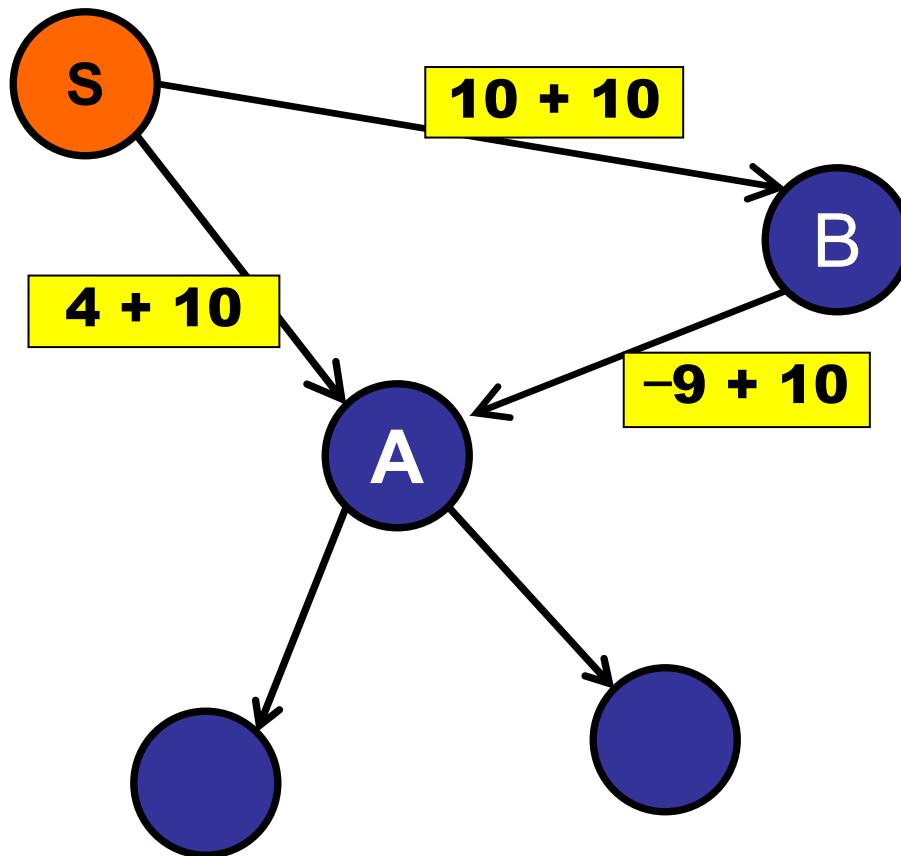
**shortest path to  
fringe vertex v**



# Dijkstra's Algorithm

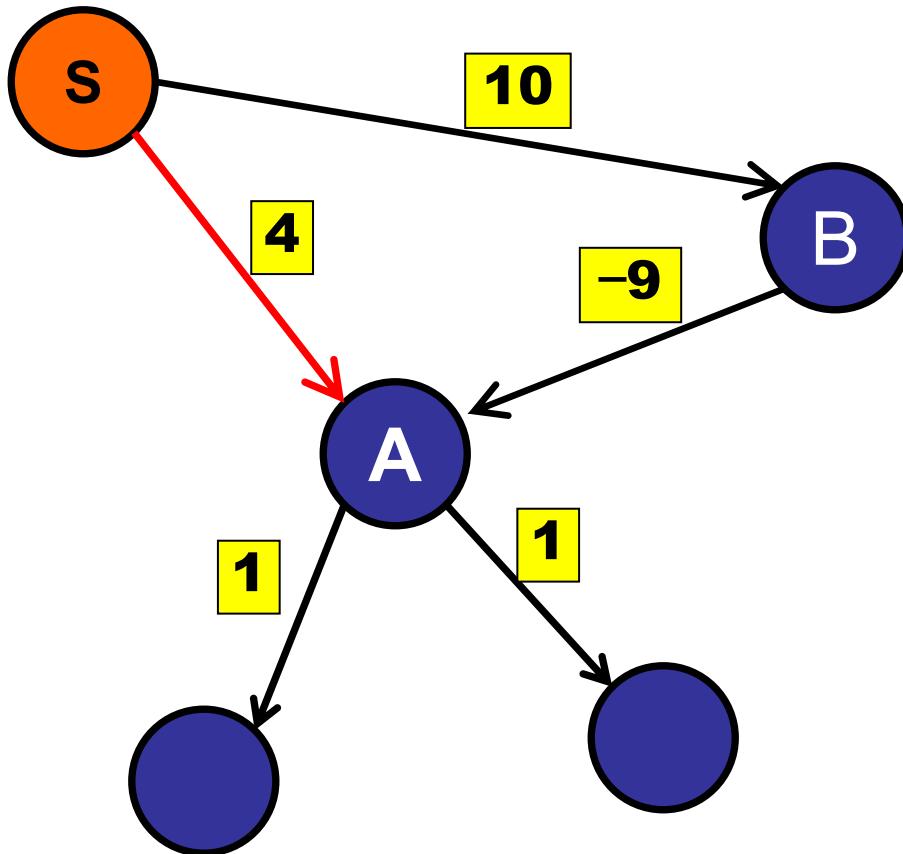
Can we reweight?

e.g.: weight += 10



# Dijkstra's Algorithm

Can we reweight?

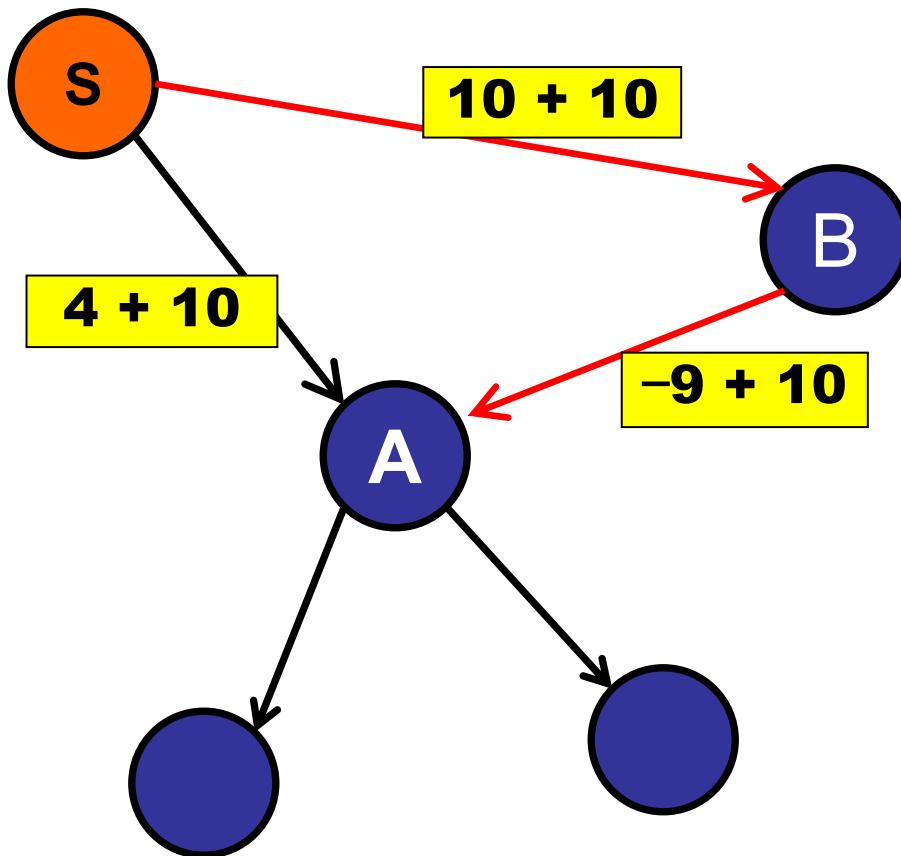


Path S-B-A: 1

Path S-A: 4

# Dijkstra's Algorithm

Can we reweight?



Path S-B-A: 21

Path S-A: 14

# Dijkstra Summary

---

Basic idea:

- Maintain distance estimates.
- Repeat:
  - Find unfinished vertex with smallest estimate.
  - Relax all outgoing edges.
  - Mark vertex finished.
- $O(E \log V)$  time (with AVL tree Priority Queue).
- No negative weight edges!

# Dijkstra Comparison

---

Same algorithm:

- Maintain a set of explored vertices.
  - Add vertices to the explored set by following edges that go from a vertex in the explored set to a vertex outside the explored set.
- 
- BFS: Take edge from vertex that was discovered **least** recently.
  - DFS: Take edge from vertex that was discovered **most** recently.
  - Dijkstra's: Take edge from vertex that is **closest** to source.

# Dijkstra Comparison

Same algorithm:

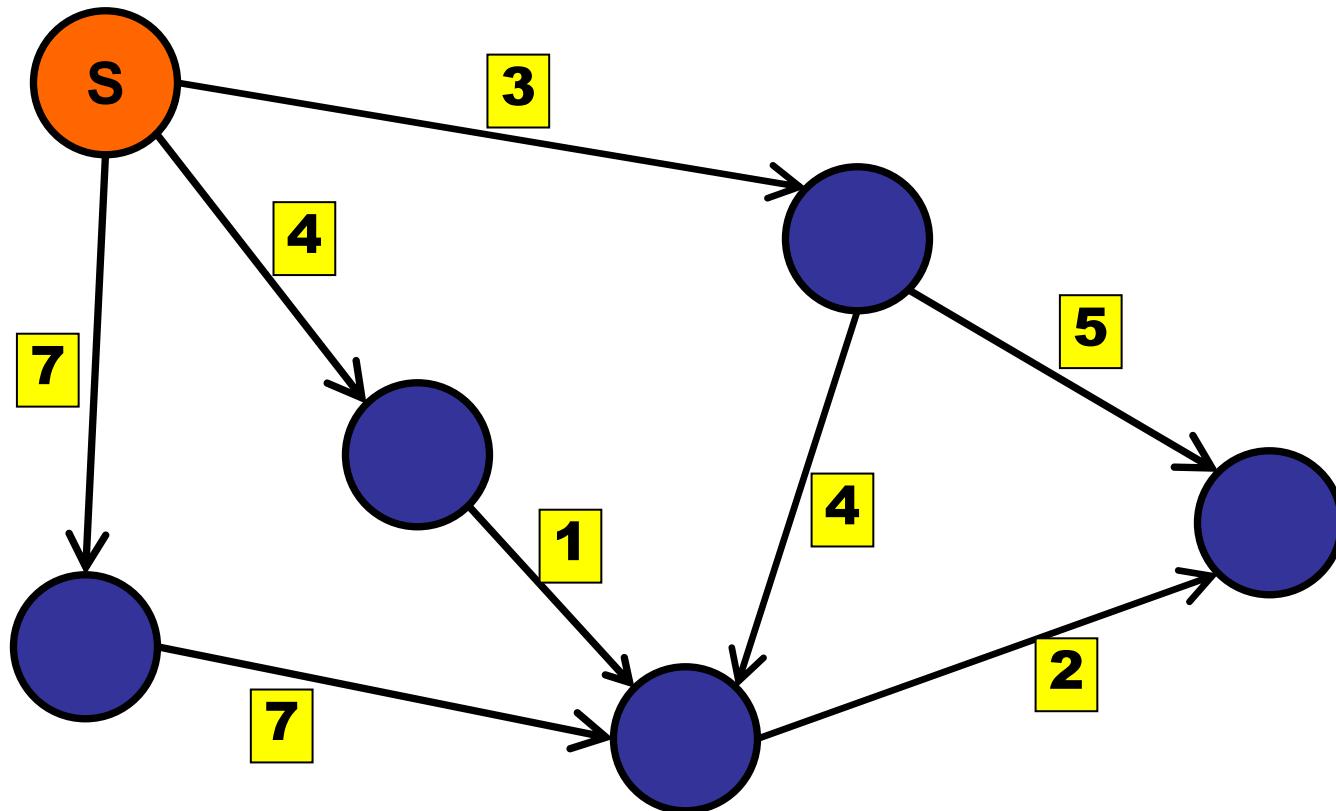
- Maintain a set of explored vertices.
  - Add vertices to the explored set by following edges that go from a vertex in the explored set to a vertex outside the explored set.
- 
- BFS: Use queue.
  - DFS: Use stack.
  - Dijkstra's: Use priority queue.



# Longest Paths

---

Any ideas?

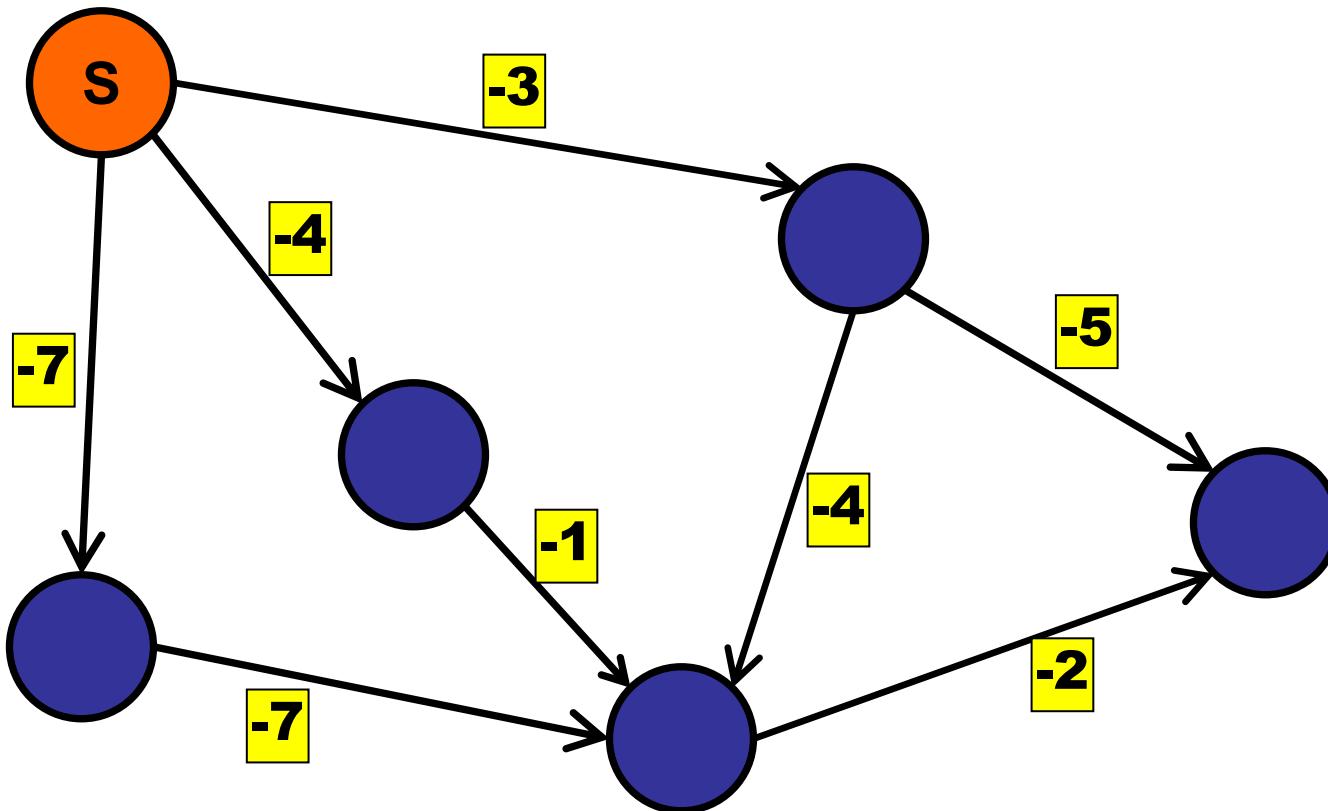


# Longest Paths

---

Negate the edges?

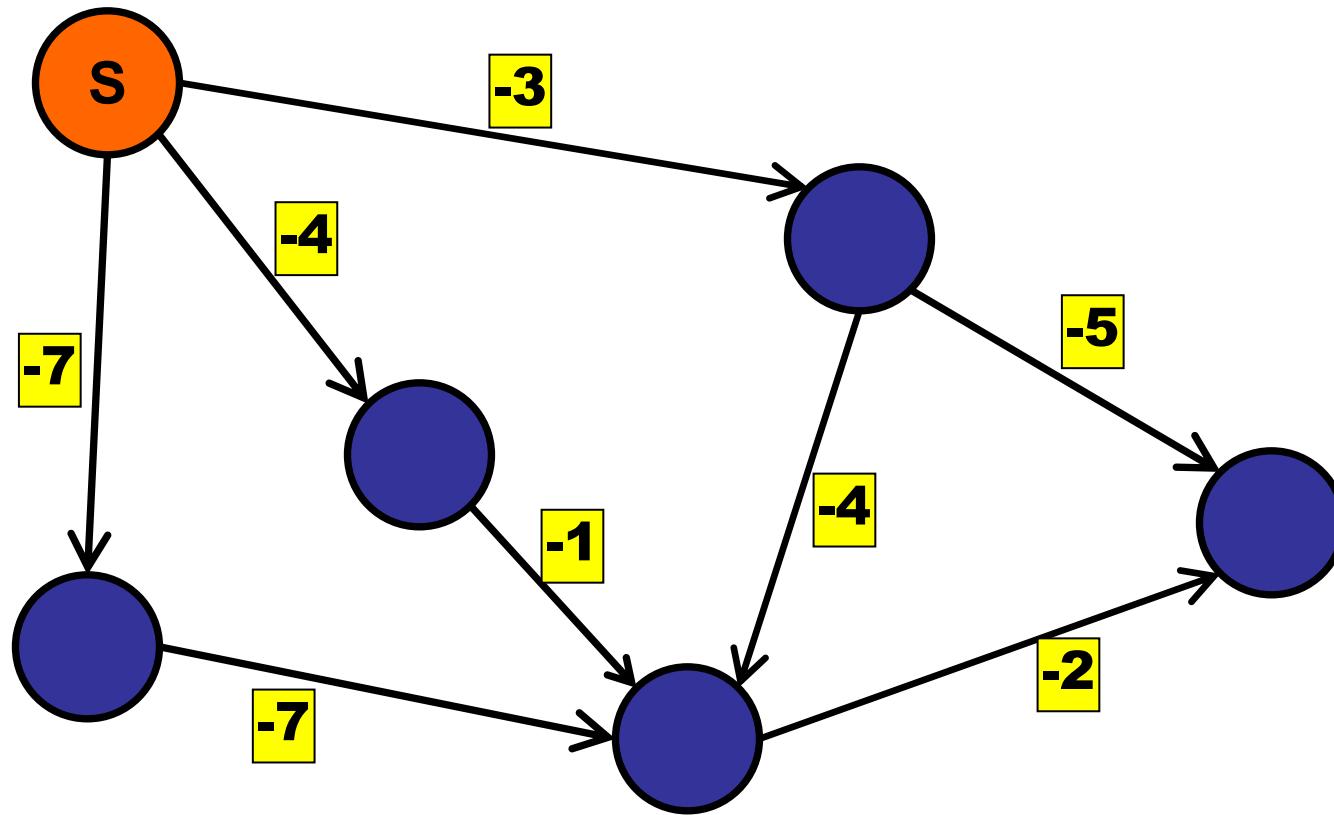
Only if your graph does not have a cycle!



# Longest Paths

Acyclic Graph:

shortest path in negated=longest path in regular



# Longest Path

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## Directed Acyclic Graph:

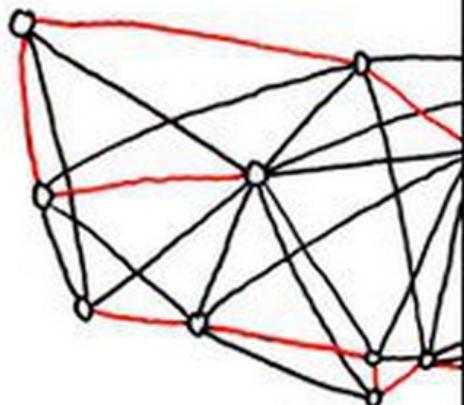
- Solvable efficiently using topological sort

## General (cyclic) Graphs:

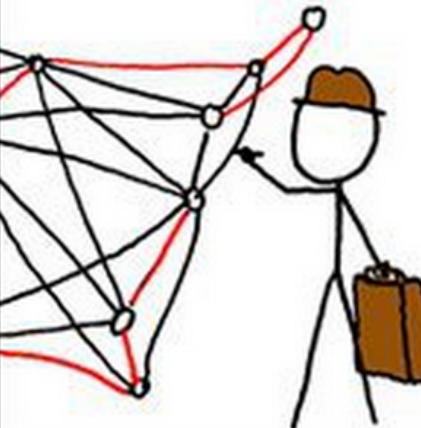
- NP-Hard
- Reduction from Hamiltonian Path:
  - If you could find the longest simple path, then you could decide if there is a path that visits every vertex.
  - Any polynomial time algorithm for longest path thus implies a polynomial time algorithm for HAMPATH.

# Also called the Travelling Salesmans Problem

BRUTE-FORCE  
SOLUTION:  
 $O(n!)$



DYNAMIC  
PROGRAMMING  
ALGORITHMS:  
 $O(n^2 2^n)$



SELLING ON EBAY:  
 $O(1)$

STILL WORKING  
ON YOUR ROUTE?



# MY HOBBY: EMBEDDING NP-COMPLETE PROBLEMS IN RESTAURANT ORDERS

## CHOTCHKIES RESTAURANT

### ~ APPETIZERS ~

MIXED FRUIT	2.15
FRENCH FRIES	2.75
SIDE SALAD	3.35
HOT WINGS	3.55
MOZZARELLA STICKS	4.20
SAMPLER PLATE	5.80

### ~ SANDWICHES ~

BARBECUE 6.55

WE'D LIKE EXACTLY \$15.05  
WORTH OF APPETIZERS, PLEASE.

... EXACTLY? UHH ...

HERE, THESE PAPERS ON THE KNAPSACK  
PROBLEM MIGHT HELP YOU OUT.

LISTEN, I HAVE SIX OTHER  
TABLES TO GET TO -

- AS FAST AS POSSIBLE, OF COURSE. WANT  
SOMETHING ON TRAVELING SALESMAN?



# Roadmap

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## Part I: Shortest Paths

## Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

# Example: DNA Alignment

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Input: two DNA strings:

- AGGAACCGTA
- AGAATCCGAA

How similar are they?

- Metric: edit distance
  - How many operations to transform one DNA string into another?

# Example: DNA Alignment

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Input: two DNA strings:

- AG~~G~~AACCGT~~A~~ ← delete G, delete T
- AGAATCCGA ← add T

Three operations:

- Delete a character
- Add a character
- Transform a character

# Example: DNA Alignment

Input: two DNA strings:

- AG~~G~~AACCGT~~A~~ ← delete G, delete T
- AGAA~~T~~CCGA ← add T

Three operations:

- Delete a character                    cost = d
- Add a character                    cost = a
- Transform a character                    cost = t

OR: minimum *cost* to transform A to B?

# Example: DNA Alignment

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Model question as a directed graph:

- For each character  $i$ , character  $j$ :
  - Create a node in the graph  $N(i,j)$
  - $N(i,j)$  represents adapting position  $i$  of the old string to match position  $j$  of the new string.
- For node  $N(i,j)$ , three outgoing edges:
  - insert character  $j+1$  from new string after position  $i$
  - delete character  $i+1$  from old string
  - transform character  $i+1$  to character  $j+1$

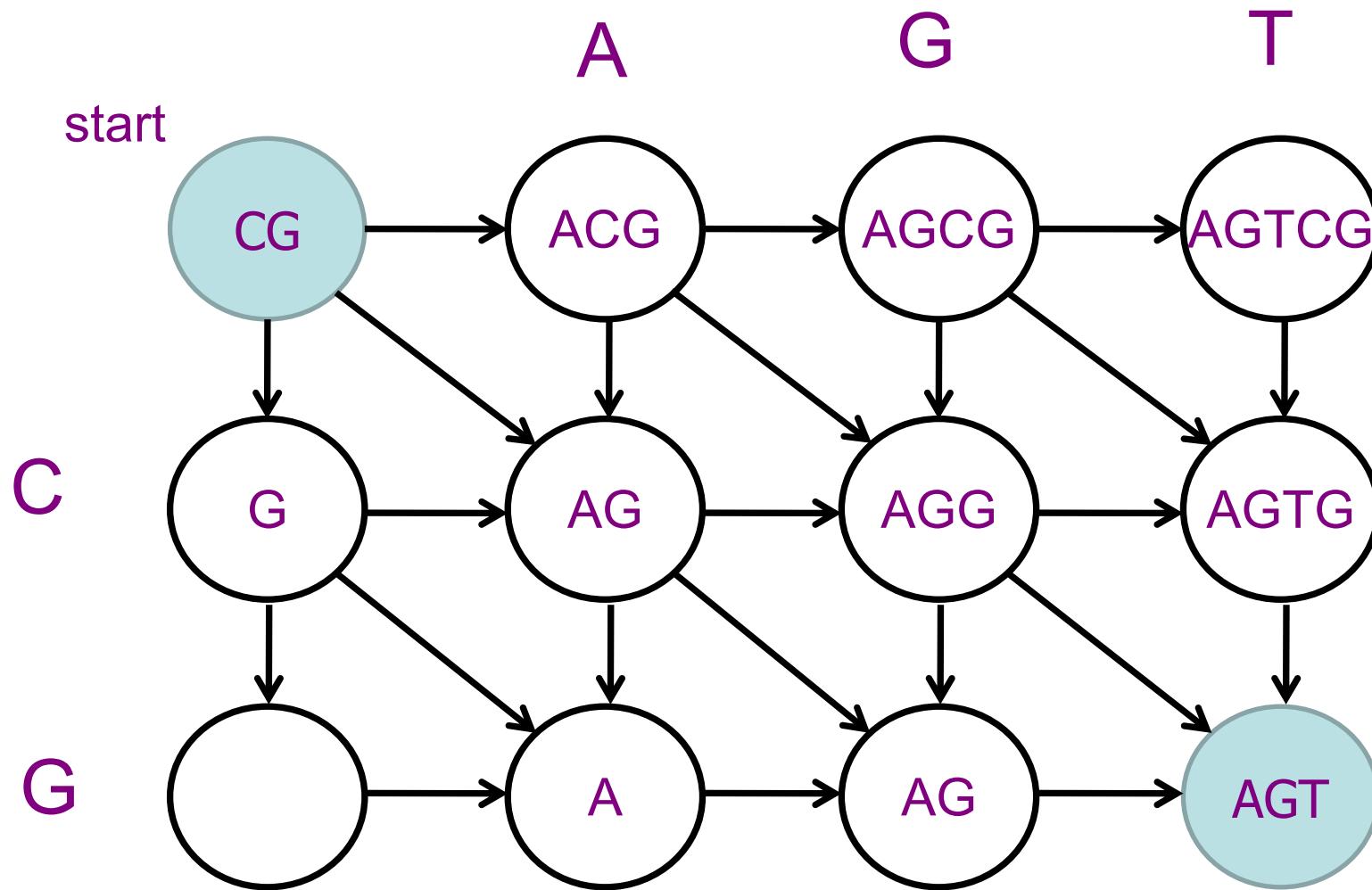
# Example: DNA Alignment

Vertical:  
delete  
character

Transform: CG to AGT

Horizontal:  
add  
character

Diagonal:  
transform  
character



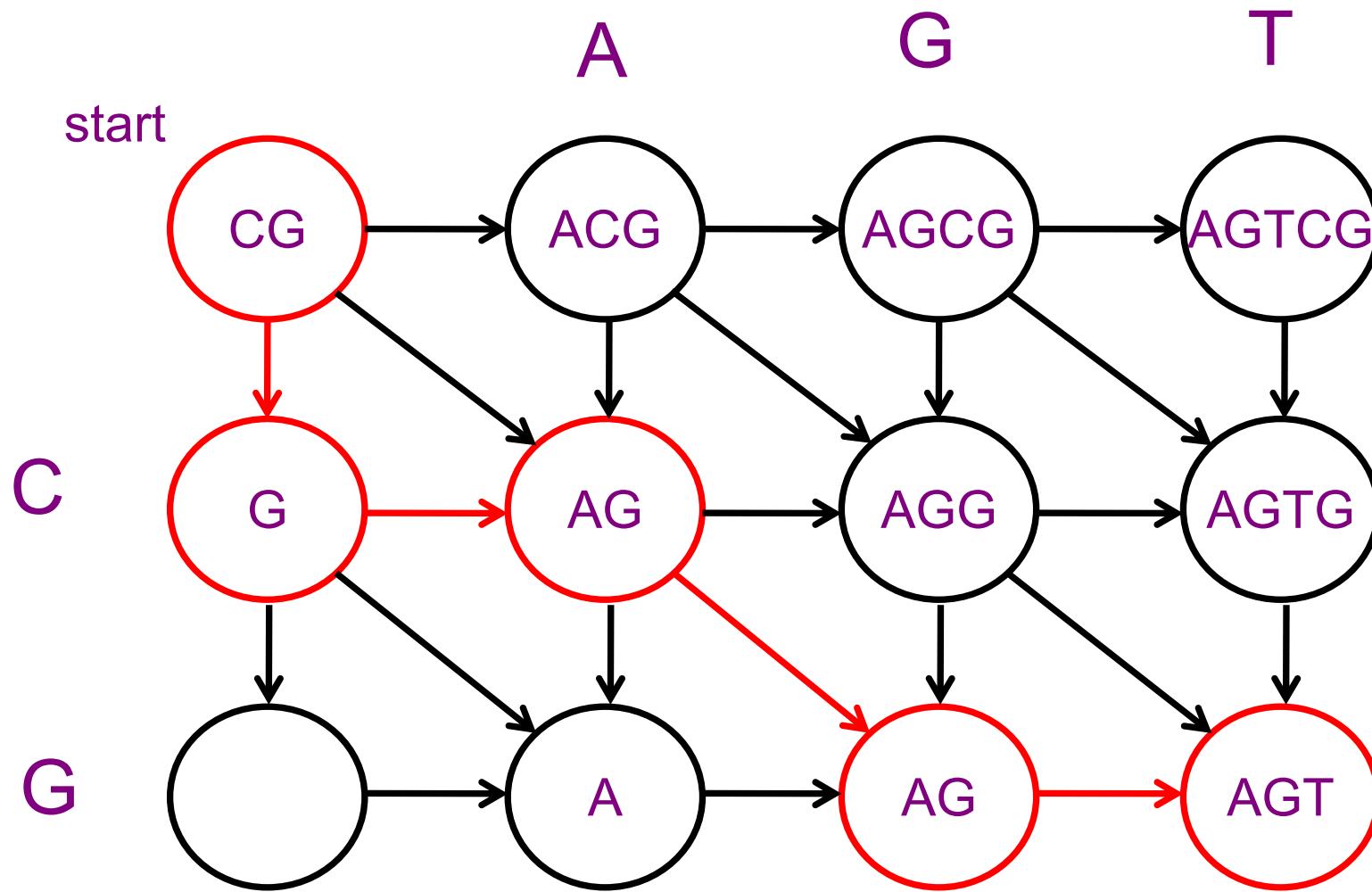
# CG to AGT

Vertical:  
delete  
character

Delete C, Add A, Leave G, Add T:

Horizontal:  
add  
character

Diagonal:  
transform  
character



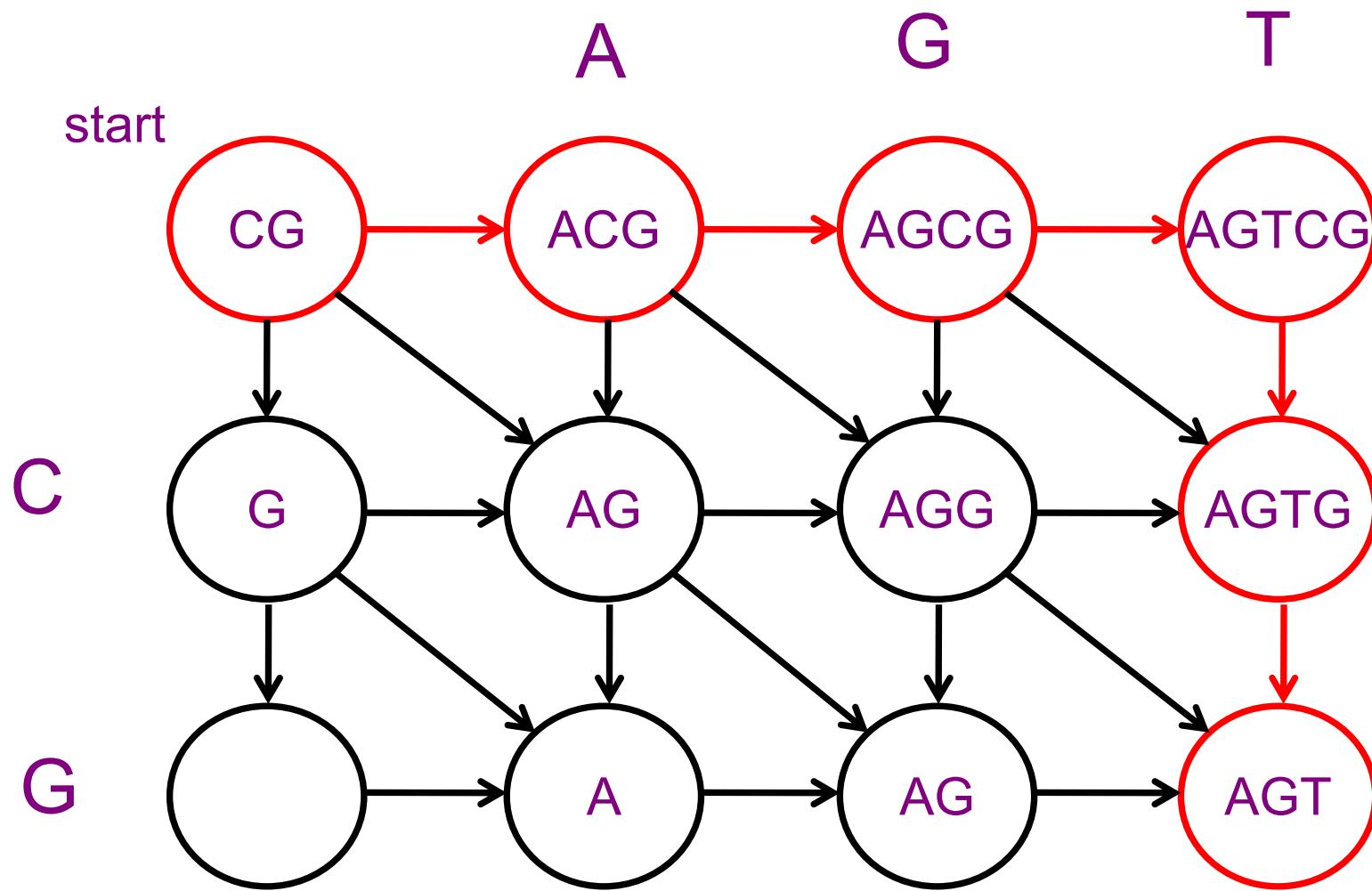
# CG to AGT

Add A, Add G, Add T, Delete C, Delete G:

Vertical:  
delete  
character

Horizontal:  
add  
character

Diagonal:  
transform  
character



# Example: DNA Alignment

---

Model question as a directed graph:

- For node  $N(i,j)$ :
  - The first  $i$  letters of the old string have been replaced with the first  $j$  letters of the new string.
  - The shortest path to  $N(i,j)$  is the shortest set of changes to change the first  $i$  letters of the old string to the first  $j$  letters of the new string.

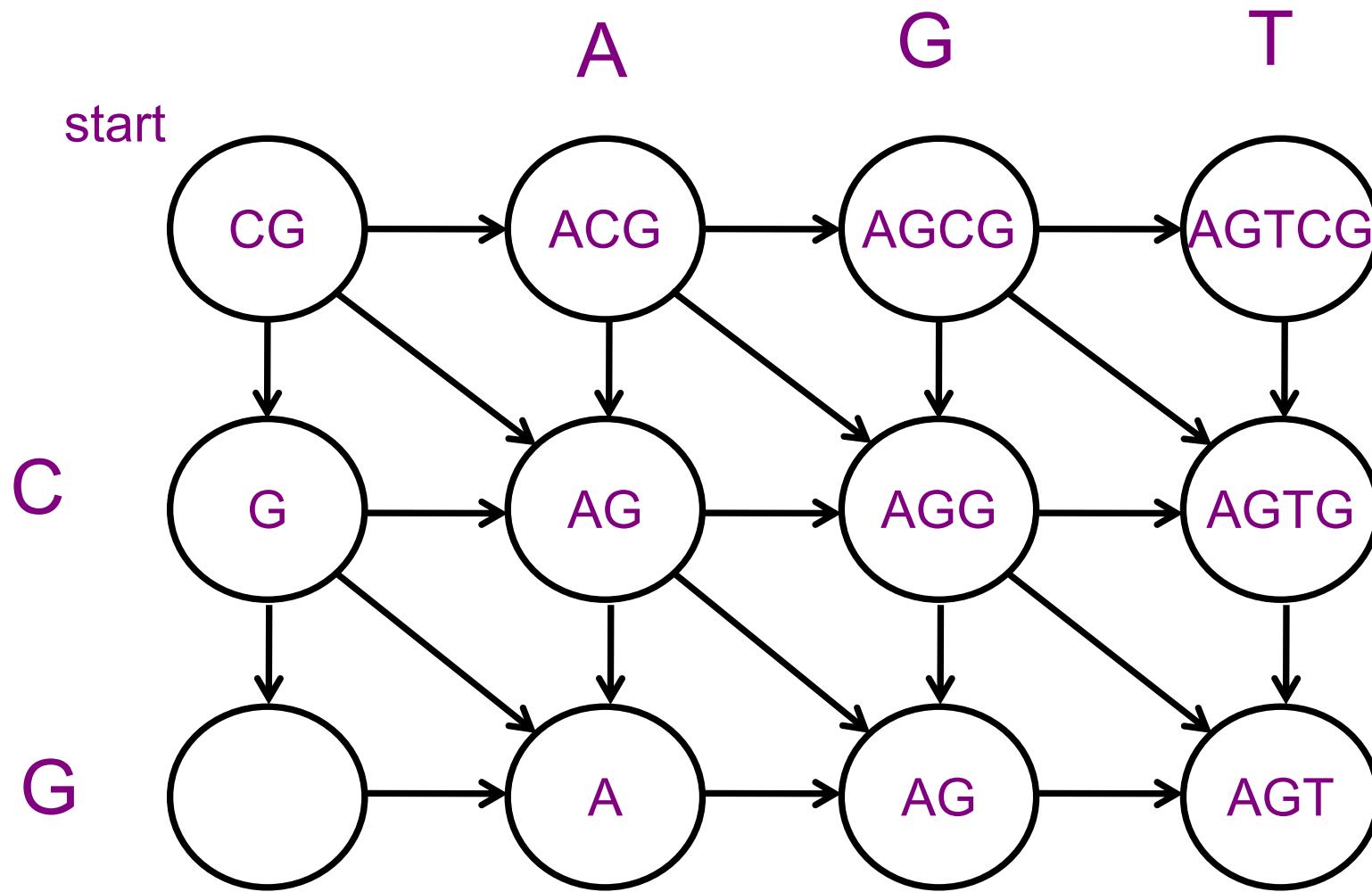
# Example: DNA Alignment

Vertical:  
delete  
character

Transform: CG to AGT

Horizontal:  
add  
character

Diagonal:  
transform  
character



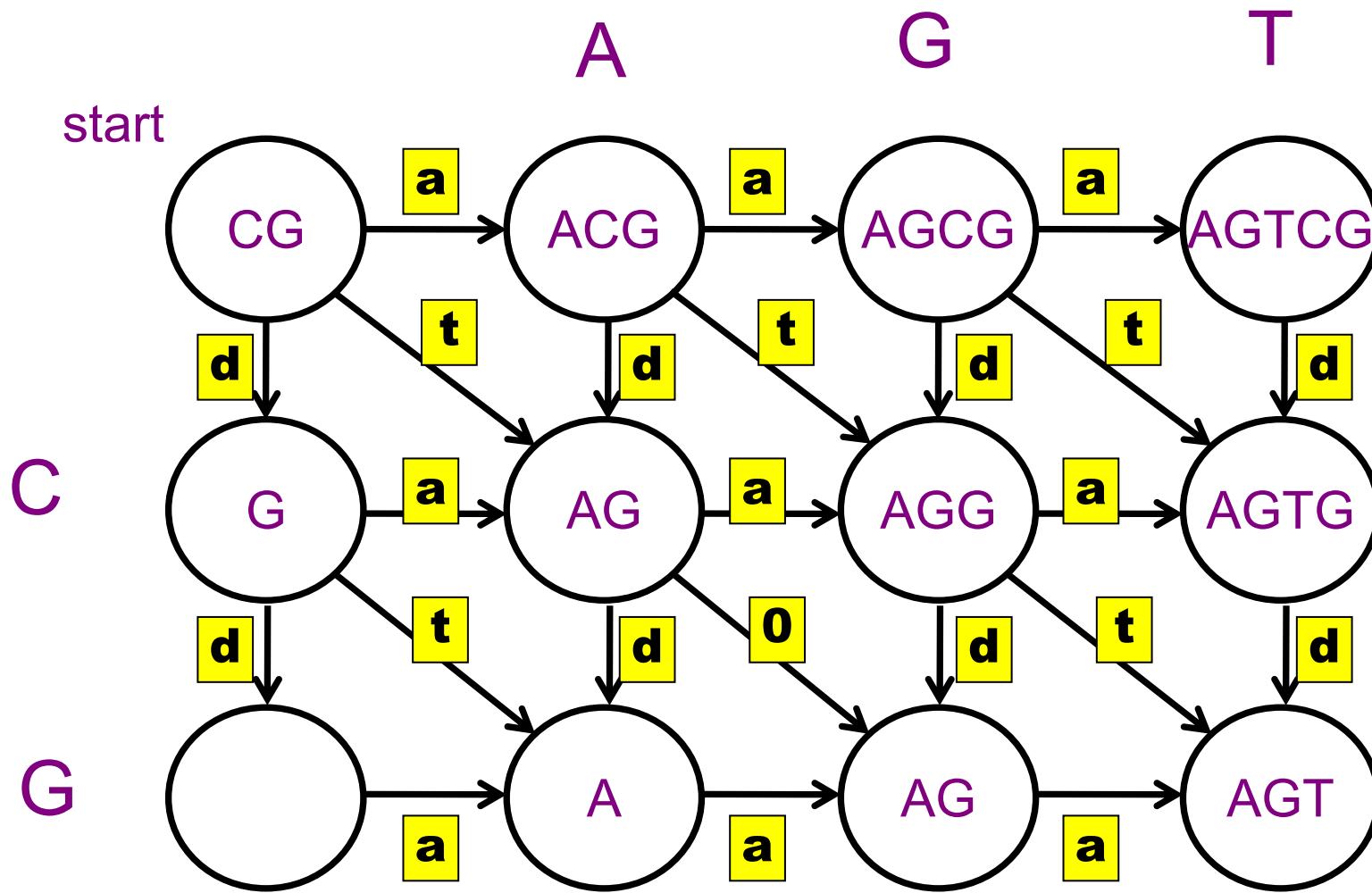
# CG to AGT

Vertical:  
delete  
character

Edge costs:

Horizontal:  
add  
character

Diagonal:  
transform  
character



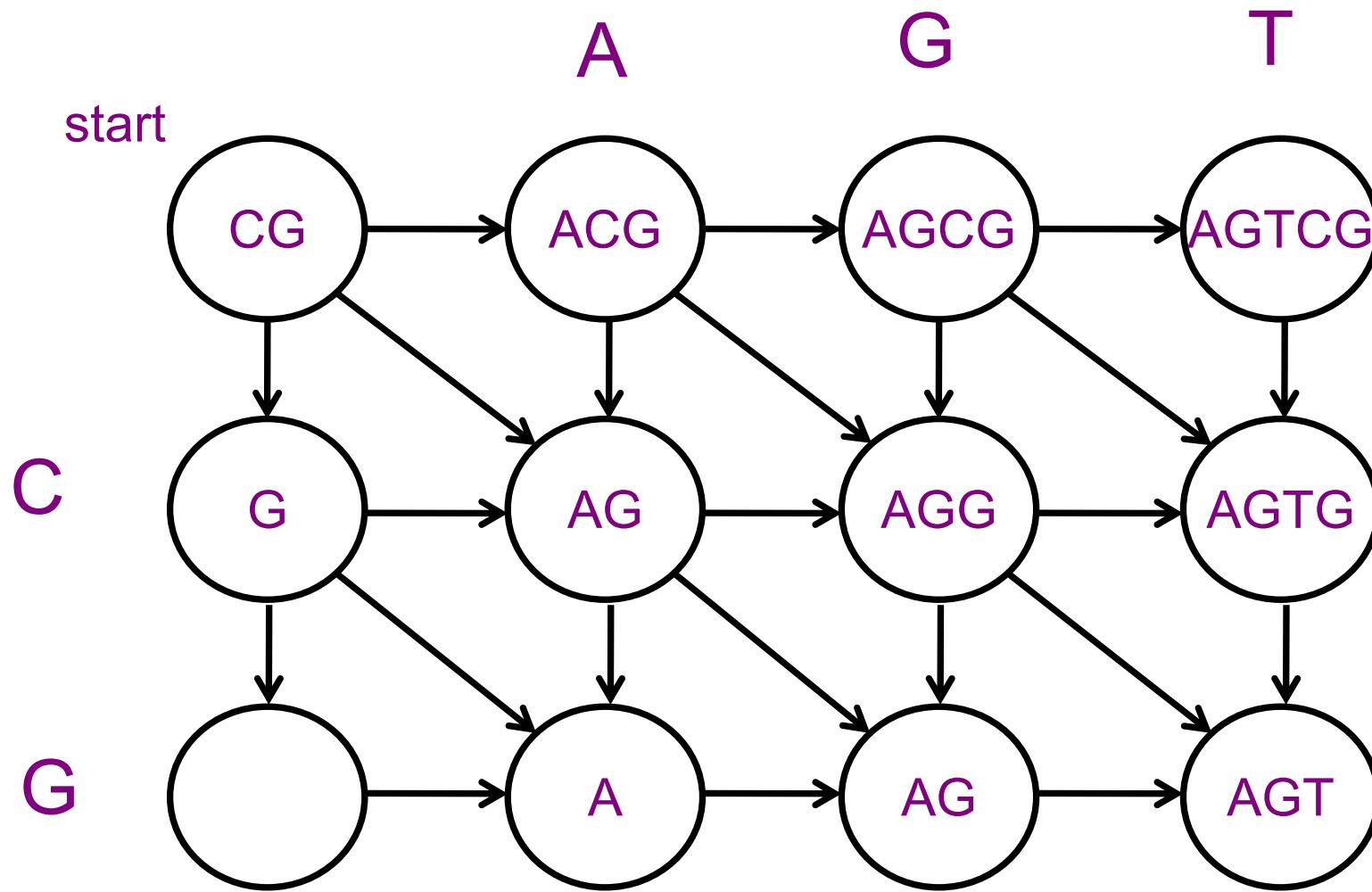
# Example: DNA Alignment

Vertical:  
delete  
character

Transform: CG to AGT

Horizontal:  
add  
character

Diagonal:  
transform  
character



# Roadmap

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## Part I: Shortest Paths

## Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

# Example: Scheduling

---

Input:

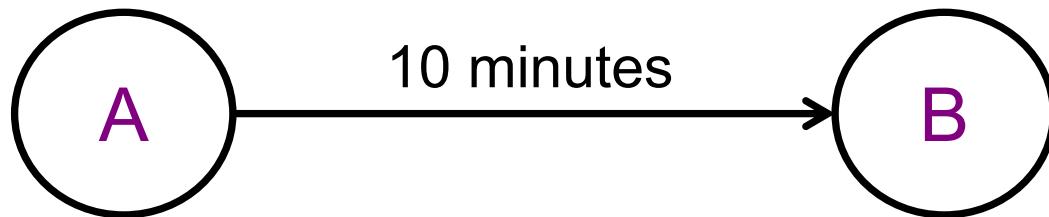
- Set of tasks: A, B, C, D, E, F
- Constraints:
  - A must be done at least 10 minutes before C
  - D must be done at most 20 minutes after E
  - B must be done after F

Output:

- Feasible?
- Schedule?

# Example: Scheduling

B must be executed **at most** 10 minutes **after A**

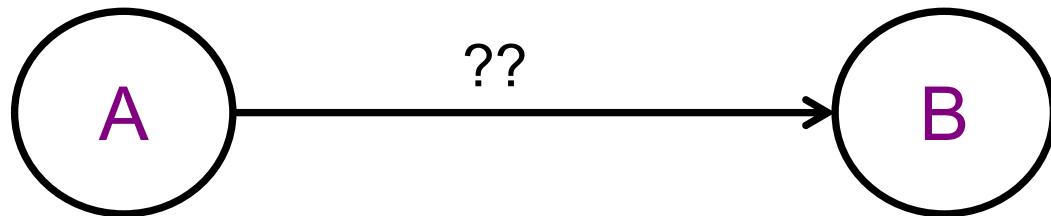


Shortest path = schedule time

triangle inequality: shortest path to B is at most 10 longer than shortest path to A

# Example: Scheduling

B must be executed **at least** 10 minutes **after A**

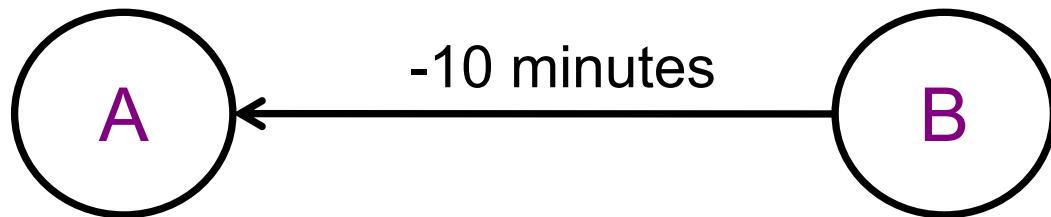


Shortest path = schedule time

triangle inequality: shortest path to B is at most 10 longer than shortest path to A

# Example: Scheduling

B must be executed **at least** 10 minutes **after** A

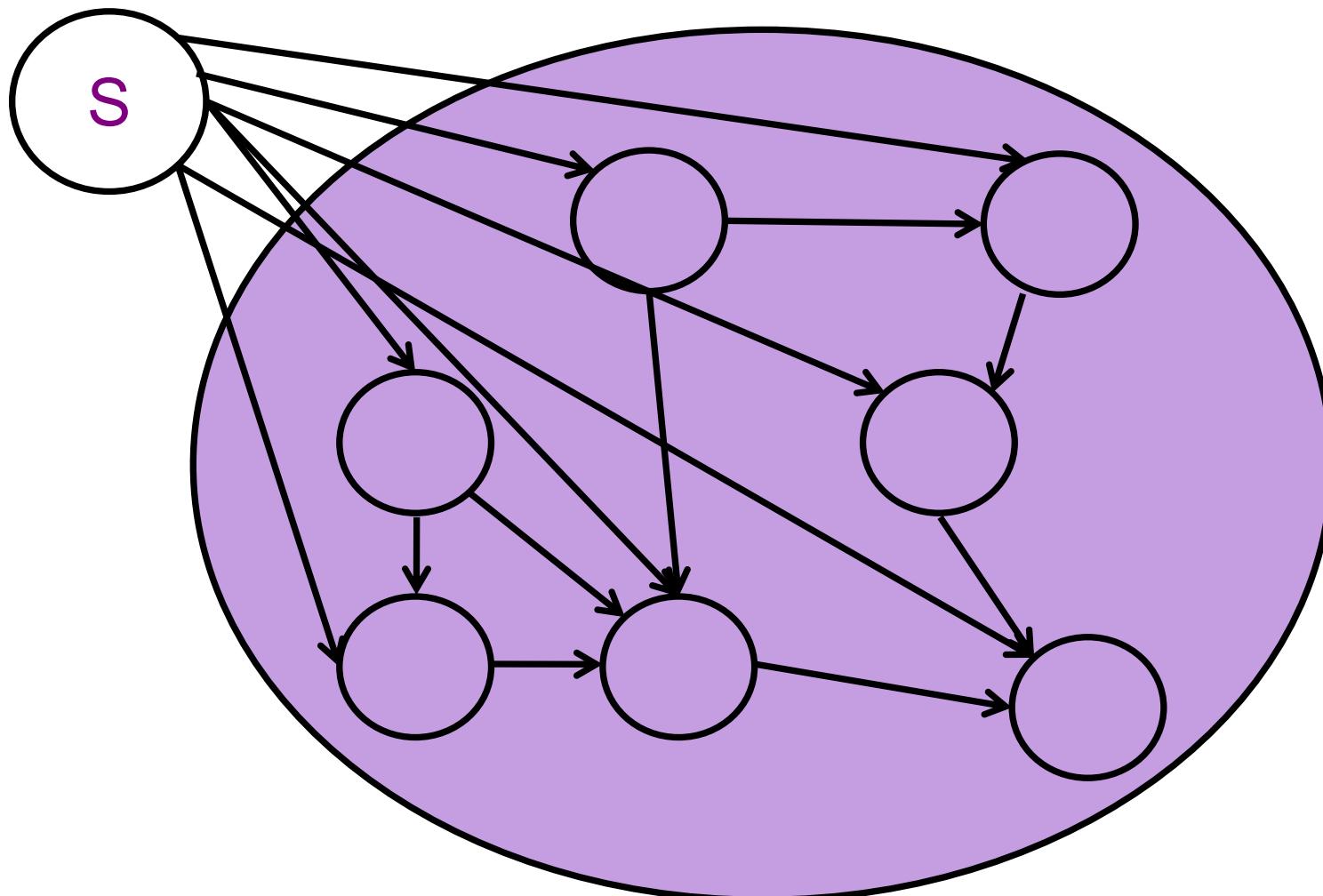


Shortest path = schedule time

triangle inequality: shortest path to B is at least 10 longer than shortest path to A

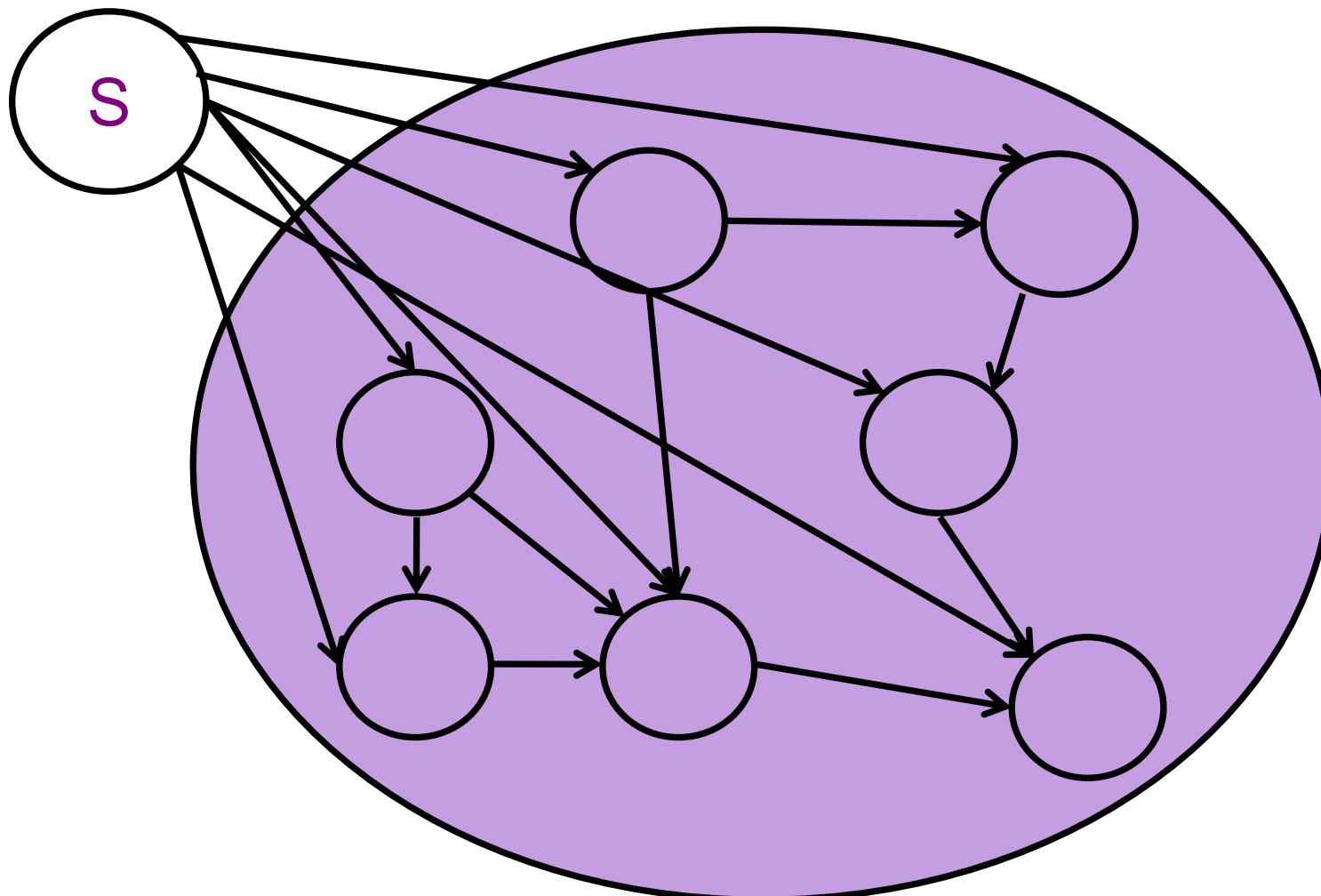
# Example: Scheduling

Add source  $S$  connected by 0 weight edges to all.



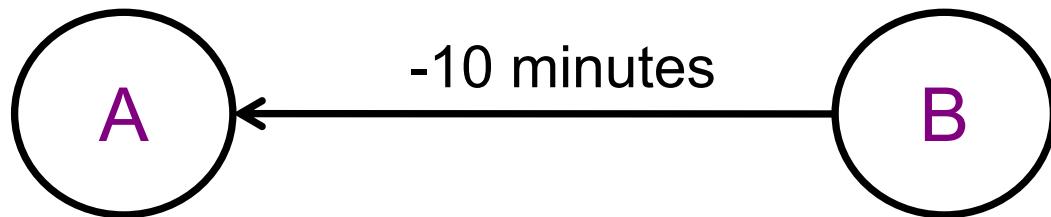
# Example: Scheduling

Solve shortest paths.



# Example: Scheduling

B must be executed **at least** 10 minutes **after** A



Negative edges: use Bellman-Ford!

Running time:  $O(nm)$

# Example: Scheduling

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## Input:

- Set of tasks: A, B, C, D, E, F
- Constraints:
  - A must be done at least 10 minutes before C
  - D must be done at most 20 minutes after E
  - B must be done after F

## Output:

- Shortest path guarantees constraints are met.
- Shortest path finishes all tasks in minimum time.

# Roadmap

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## Part I: Shortest Paths

## Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems