# CS2020 Data Structures and Algorithms

**Shortest Paths** 

### Roadmap

#### Part I: Shortest Paths

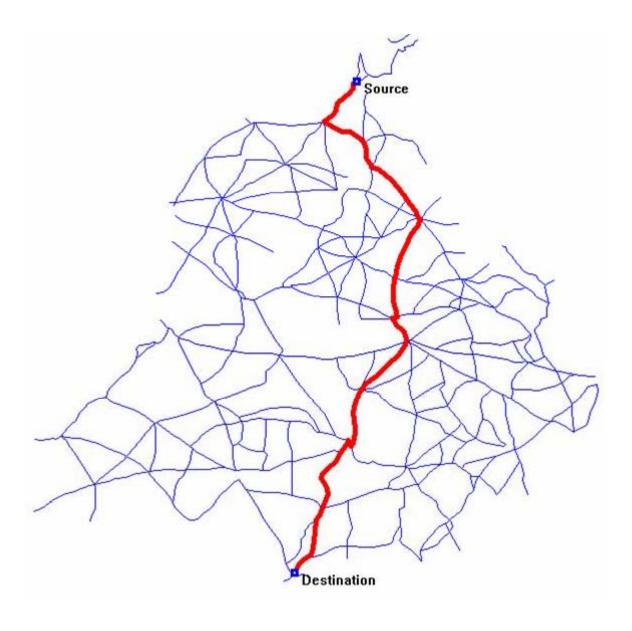
- Special Case: Tree
- Special Case: Non-negative weights (Dijkstra's)
- Special Case: Directed Acyclic Graphs

#### Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

### SHORTEST PATHS

(ON WEIGHTED GRAPHS)



### Shortest Path Problem

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

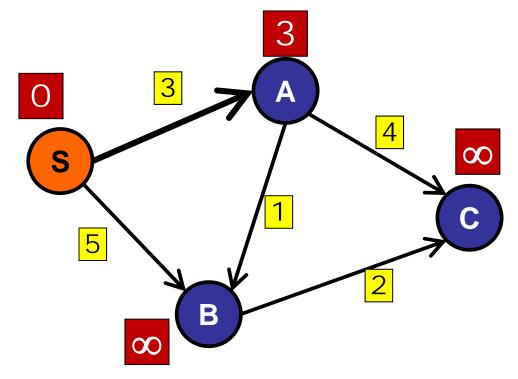
- Source-to-destination: one vertex to another
- Single source: one vertex to every other
- All pairs: between all pairs of vertices

#### Variants:

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

#### **Shortest Paths**

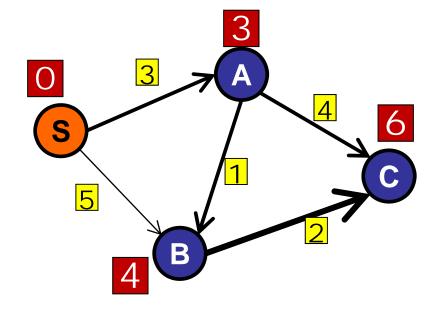
```
relax(int u, int v){
    if (dist[v] > dist[u] + weight(u,v))
        dist[v] = dist[u] + weight(u,v);
}
```



### Bellman-Ford

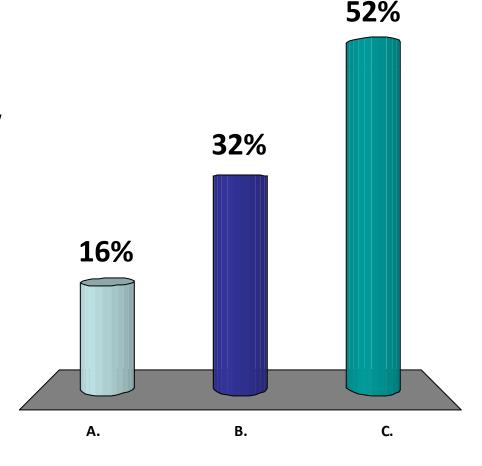
```
n = V.length;
for (i=0; i<n; i++)
    for (Edge e : graph)
        relax(e)</pre>
```





# What is the meaning of negative cycles in SSSP?

- A. Use Bellman-Ford in the last lecture
- B. We will talk about how to solve today
- C. The meaning of negative cycles will make the SSSP meaningless



### 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

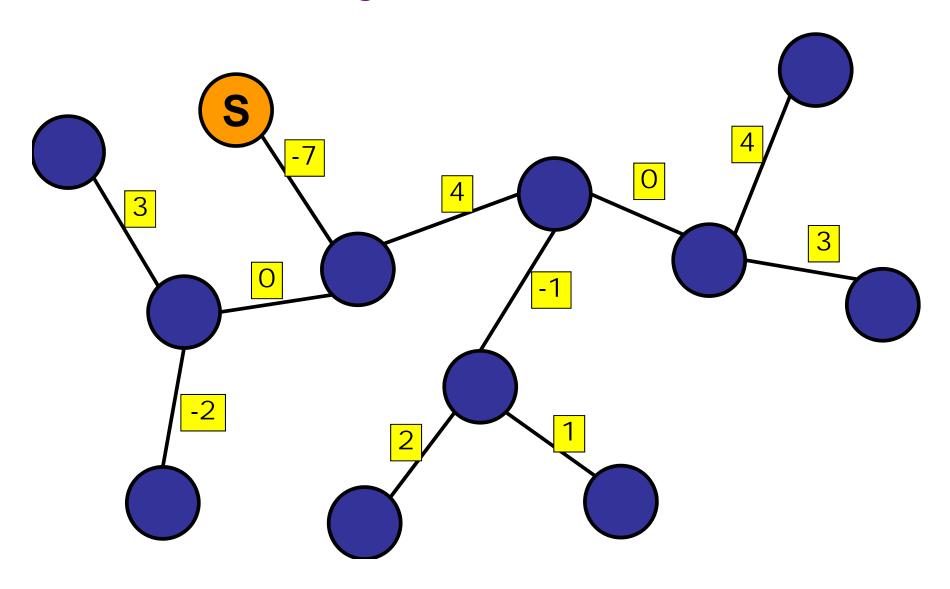
#### Key idea:

Relax the edges in the "right" order.

#### Only relax each edge once:

O(E) cost (for relaxation step).

Undirected, weighted



### Aside: Trees, Redefined

#### What is an (undirected) tree?

A graph with no cycles is an (undirected) tree.

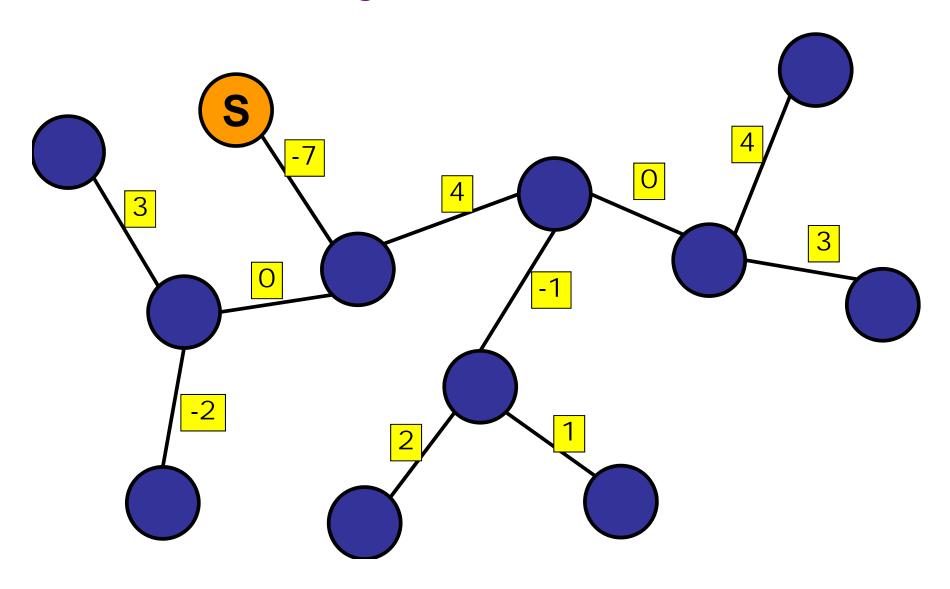
#### What is a *rooted* tree?

A tree with a special designated root note.

#### Our previous (recursive) definition of a tree:

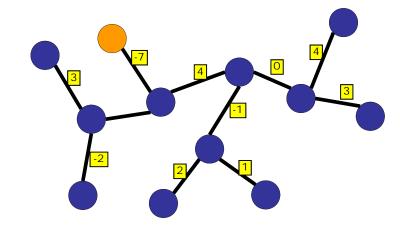
- A node with zero, one, or more sub-trees.
- I.e., a rooted tree.

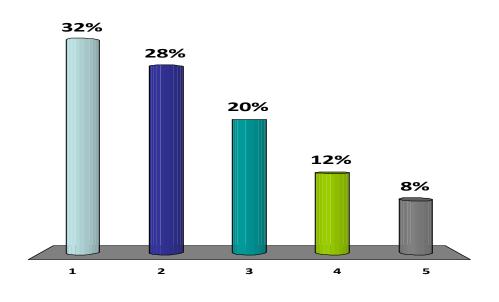
Undirected, weighted



# Which algorithm is best for checking if a graph is a tree?

- **✓**1. BFS
- ✓2. DFS
  - 3. Bellman Ford
  - 4. Topological Sort
  - 5. Dijkstra's Algorithm





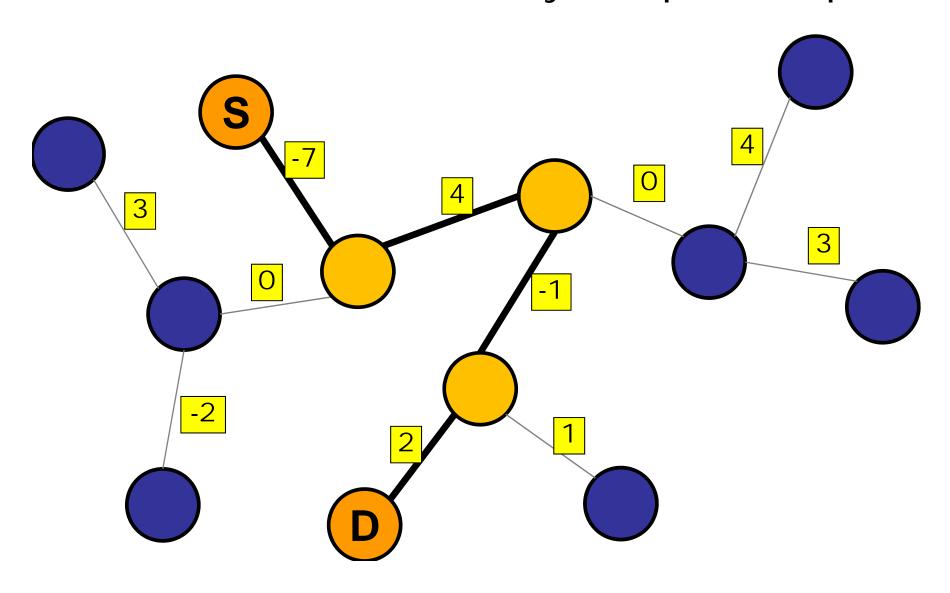
# Aside: Tree Checking

If it is connected...

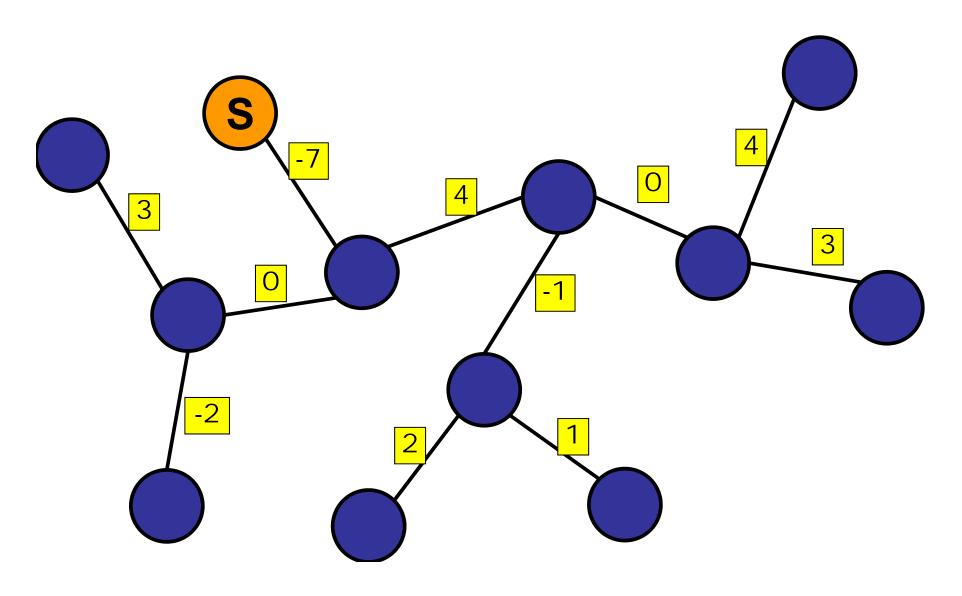
If it is disconnected...

If it is directed...

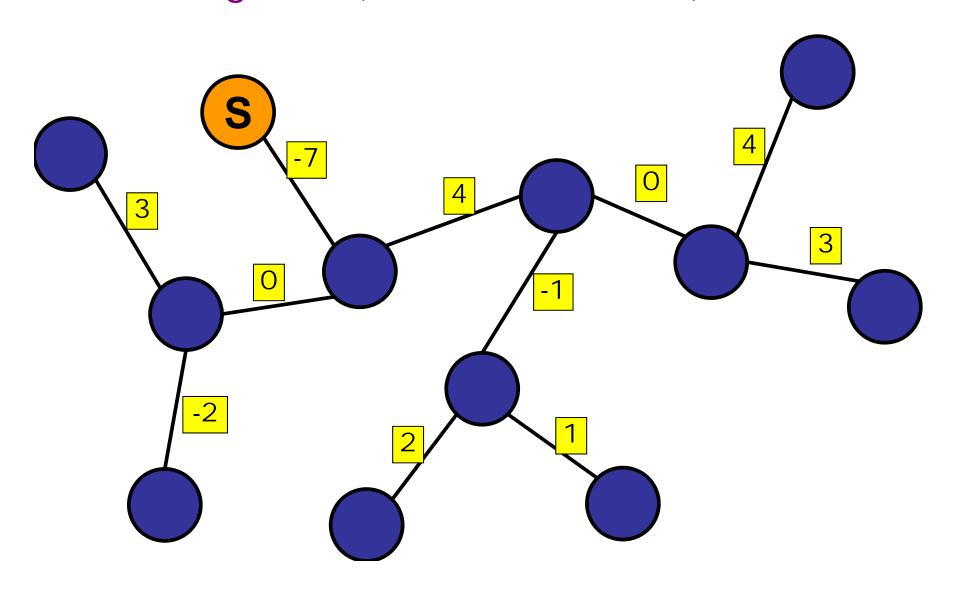
source-to-destination: only one possible path!

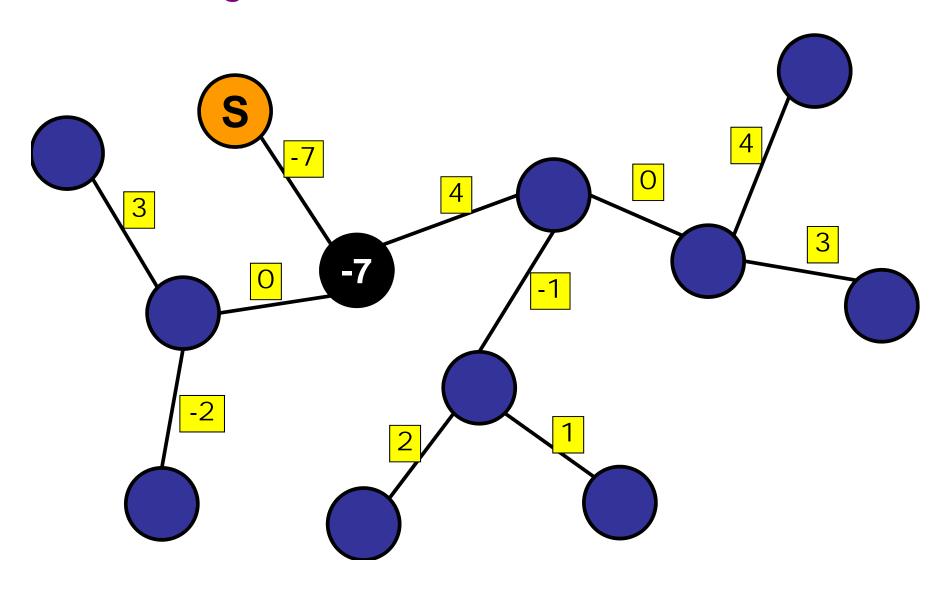


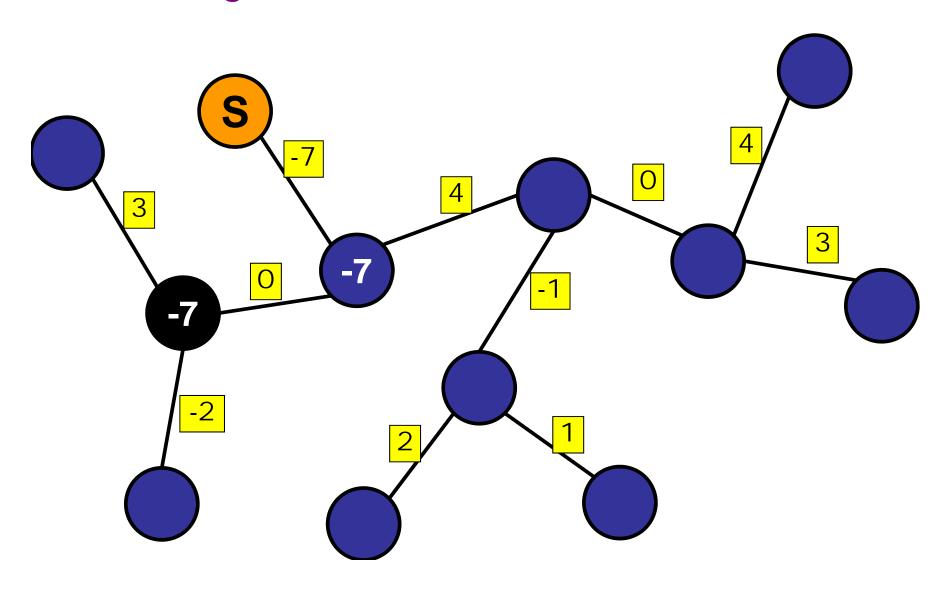
source-to-all: what order to relax?

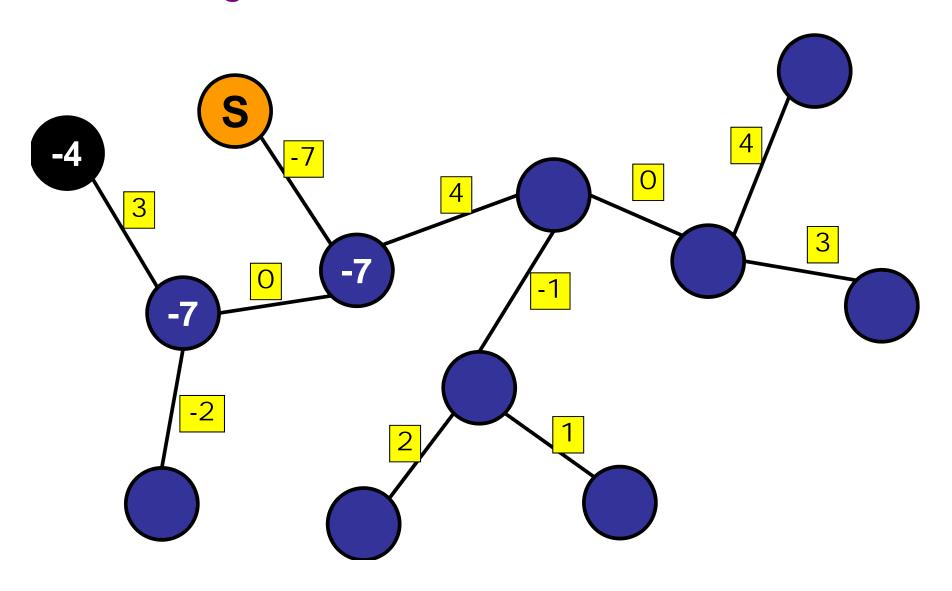


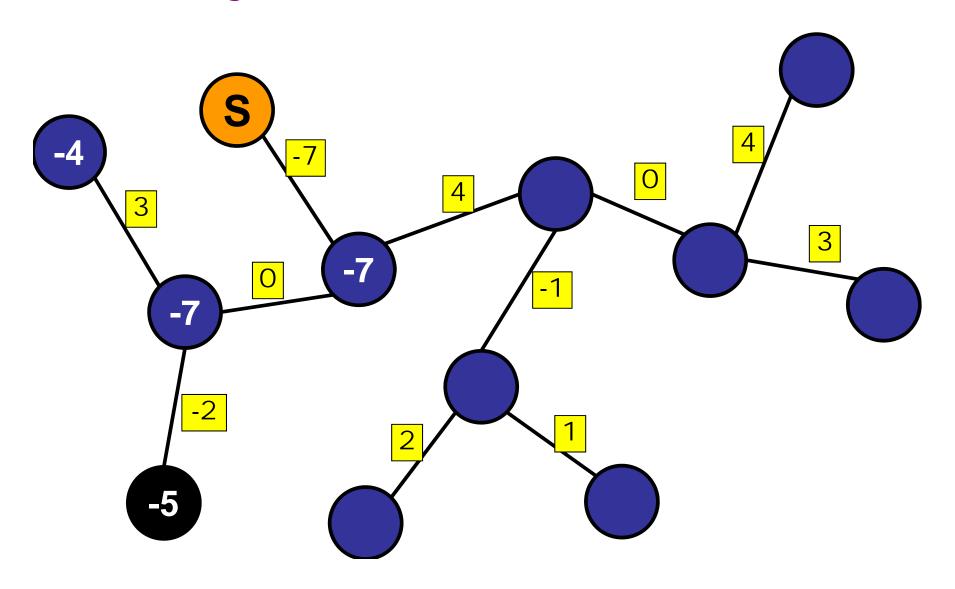
Relax edges in (BFS or DFS order).

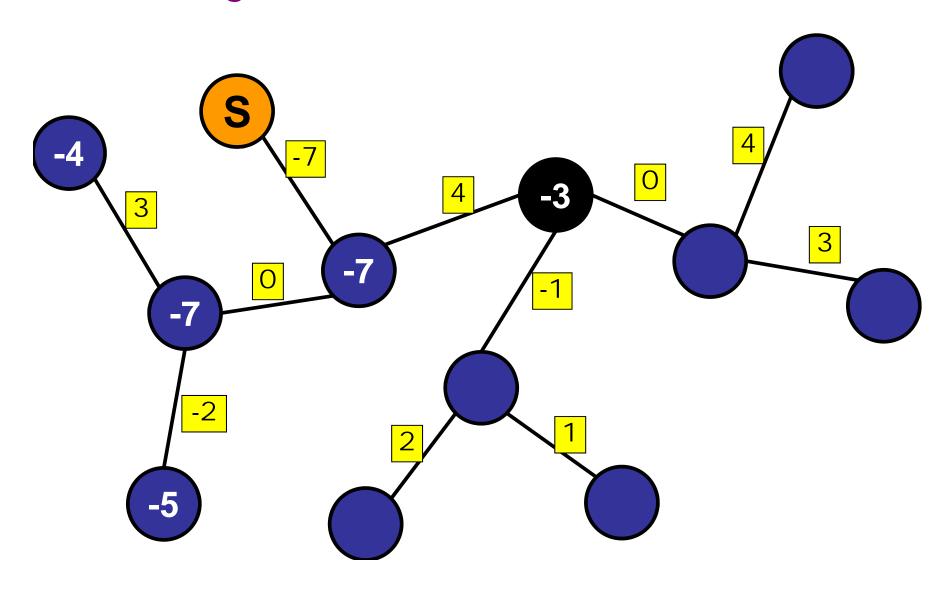


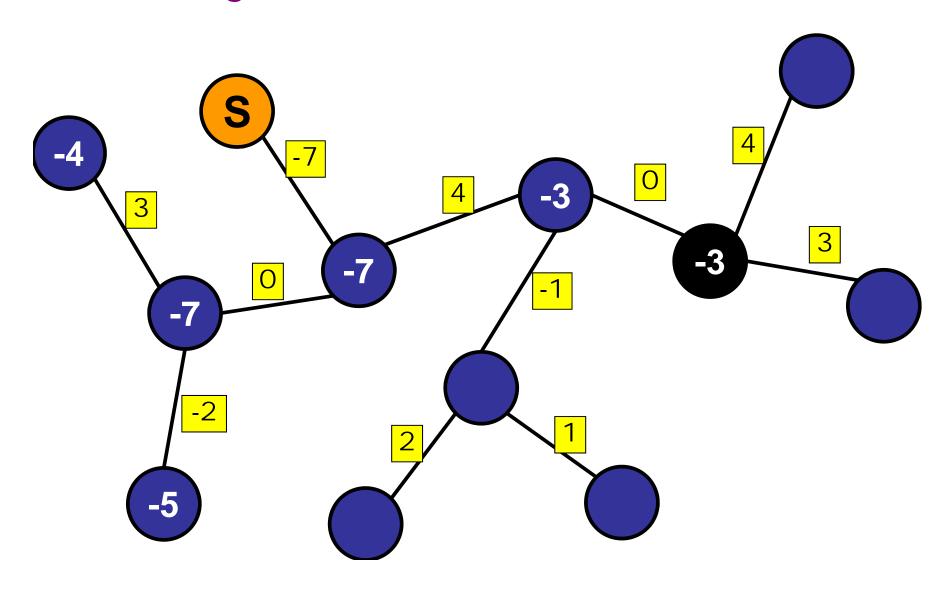


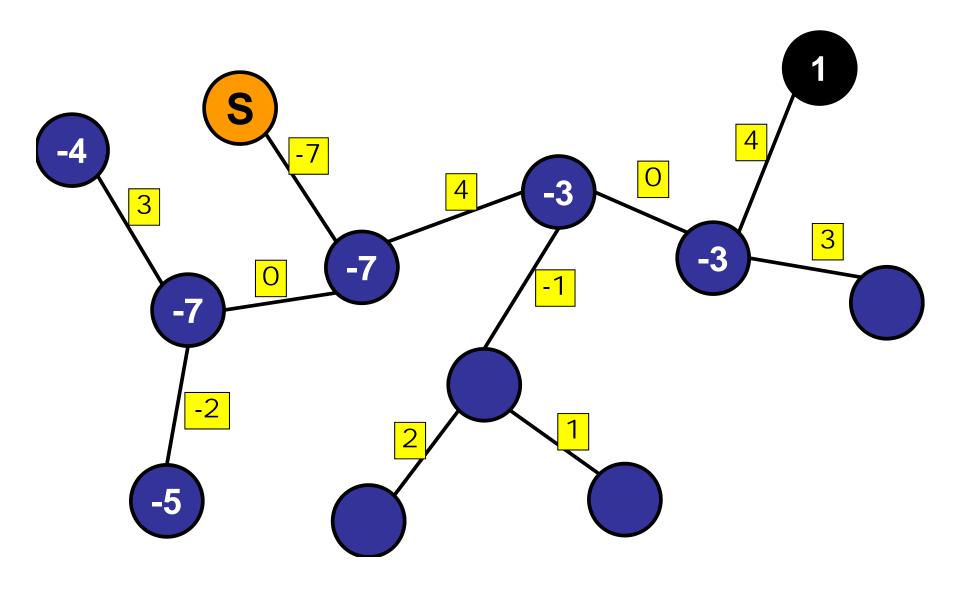


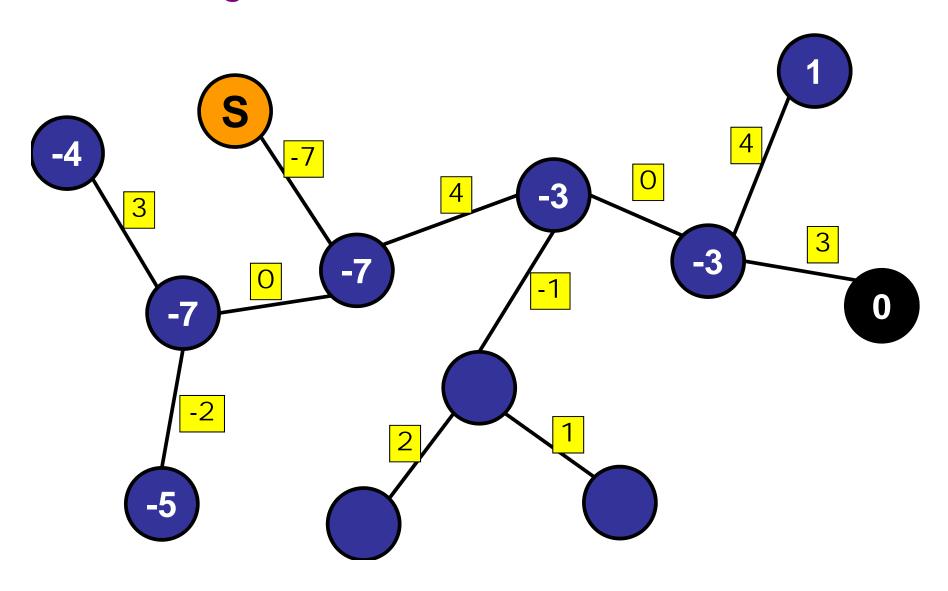


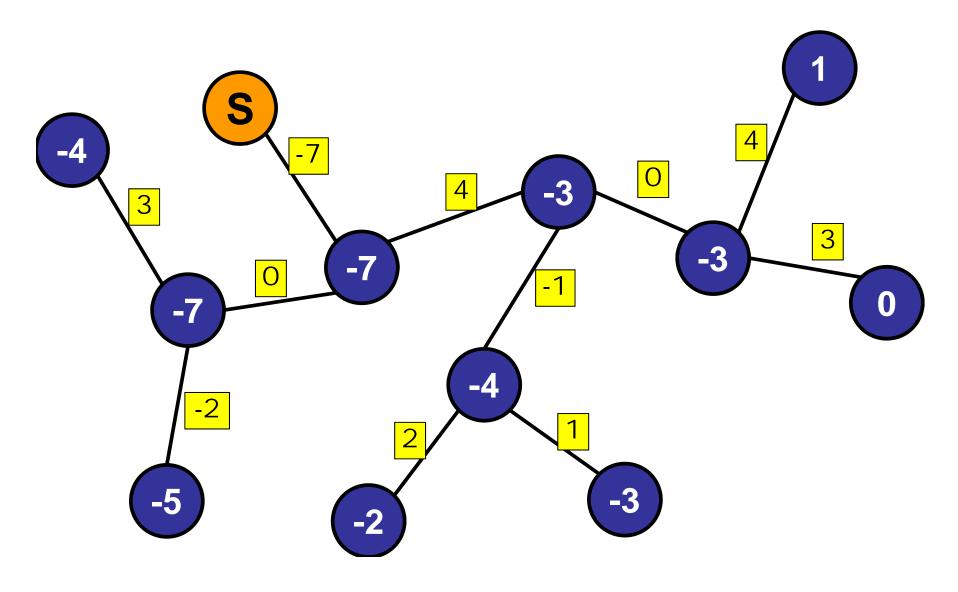












#### Basic idea:

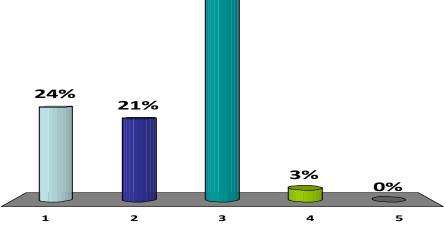
- Perform DFS or BFS
- Relax each edge the first time you see it.
- O(V) time.

#### **Assumptions:**

- Weighted edges
- Positive or negative weights
- Undirected tree

#### Why is the running time O(V)?

- 1. You only need to explore 1 outgoing edge for each vertex.
- 2. DFS/BFS run in O(V) time on a graph.
- ✓ 3. There are only O(V) edges in a tree.
  - 4. It is not O(V): you need to explore every edge!
  - 5. I'm confused.



#### Basic idea:

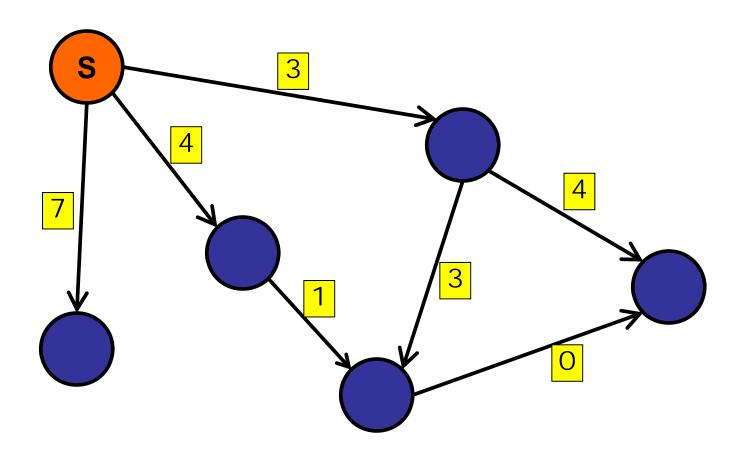
- Perform DFS or BFS
- Relax each edge the first time you see it.
- O(V) time.

#### **Assumptions:**

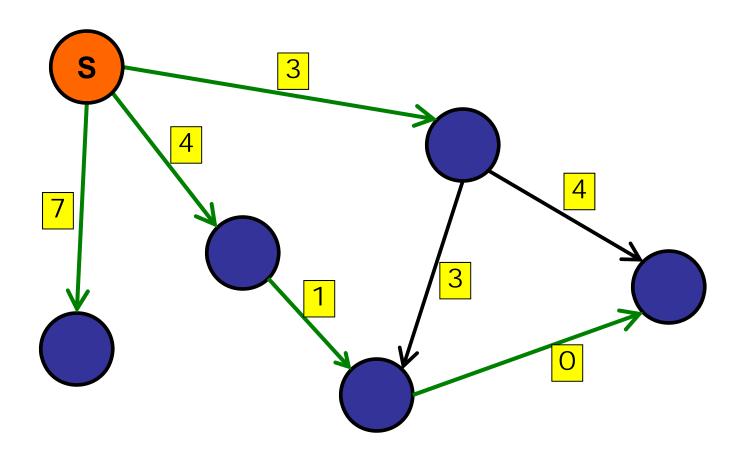
- Weighted edges
- Positive or negative weights
- Undirected tree

# General Graph

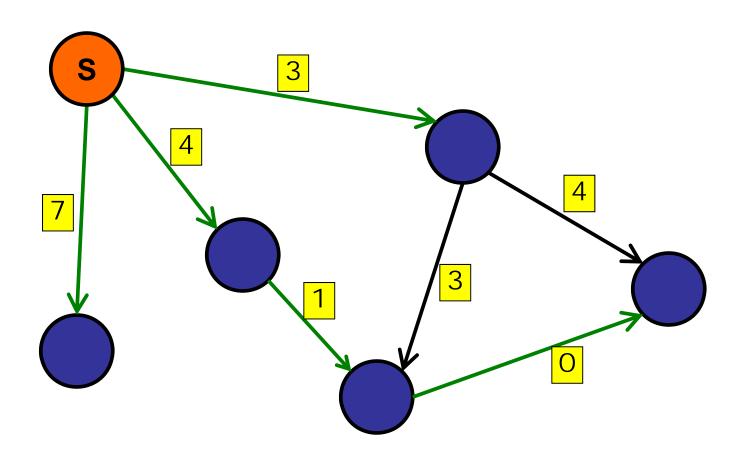
### Non-negative edges



For every node: add 1 shortest path to the tree.

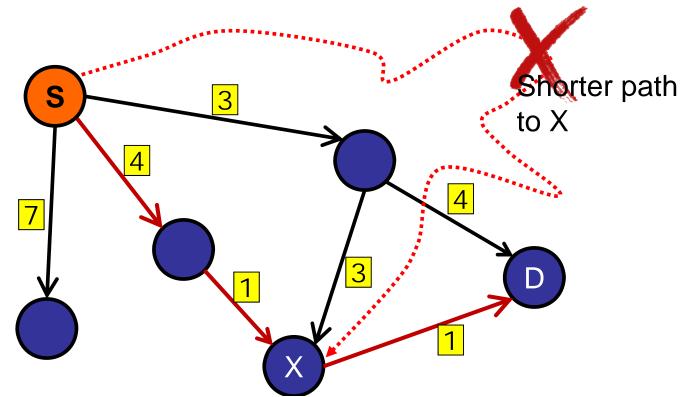


Why are there no cycles?



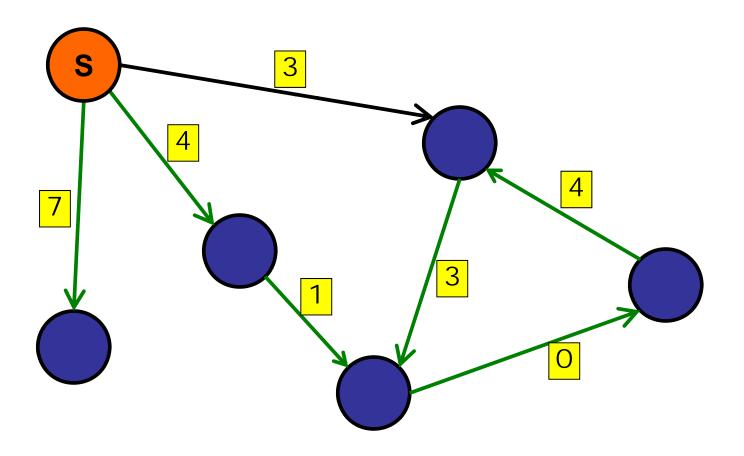
#### Key property:

If P is the shortest path from S to D, and if P goes through X, then P is also the shortest path from S to X (and from X to D).



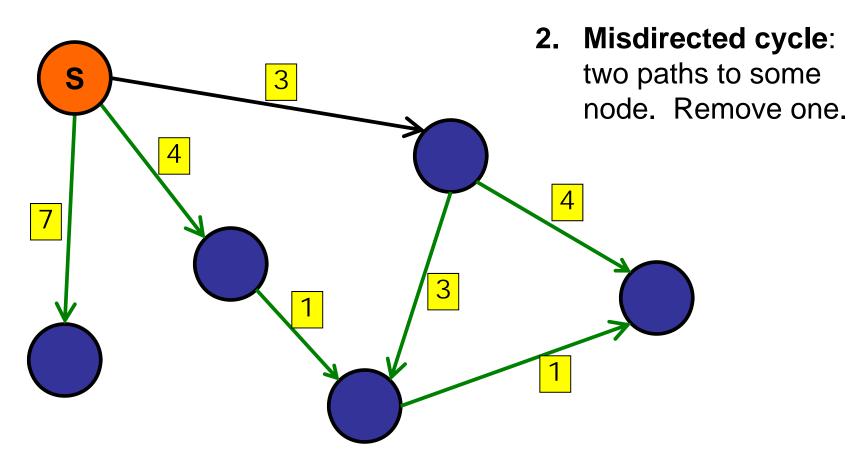
Why are there no cycles?

1. Directed cycle: remove one edge to get shorter paths.

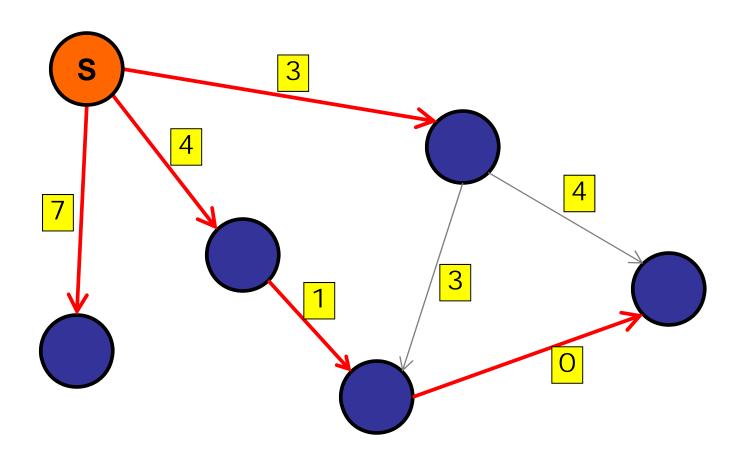


Why are there no cycles?

1. Directed cycle: remove one edge to get shorter paths.



No cycles in the shortest path tree.



### Today

#### Key idea:

Relax the edges in the "right" order.

### Only relax each edge once:

O(E) cost (for relaxation step).

### Edsger W. Dijkstra

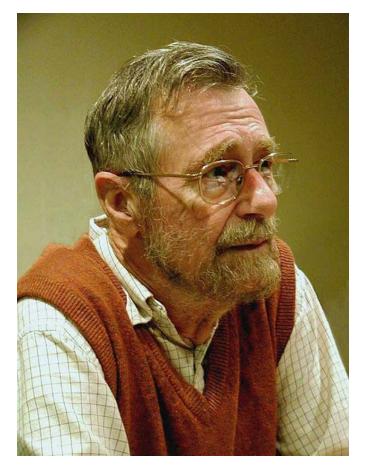
"Computer science is no more about computers than astronomy is about telescopes."

"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

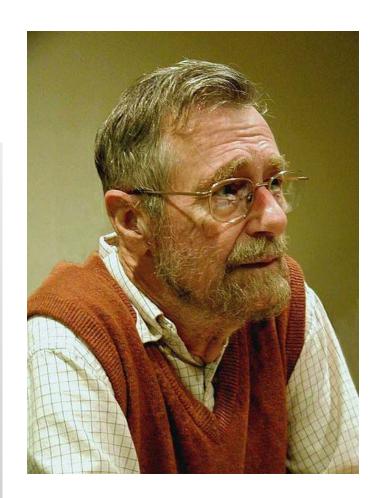
### Edsger W. Dijkstra

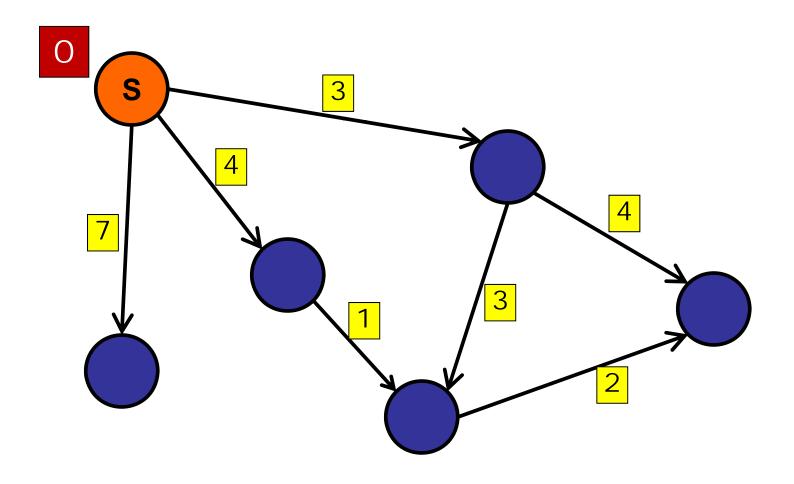
"It is practically impossible to teach good programming to students that have had a prior exposure to BASIC: as potential programmers they are mentally mutilated beyond hope of regeneration."

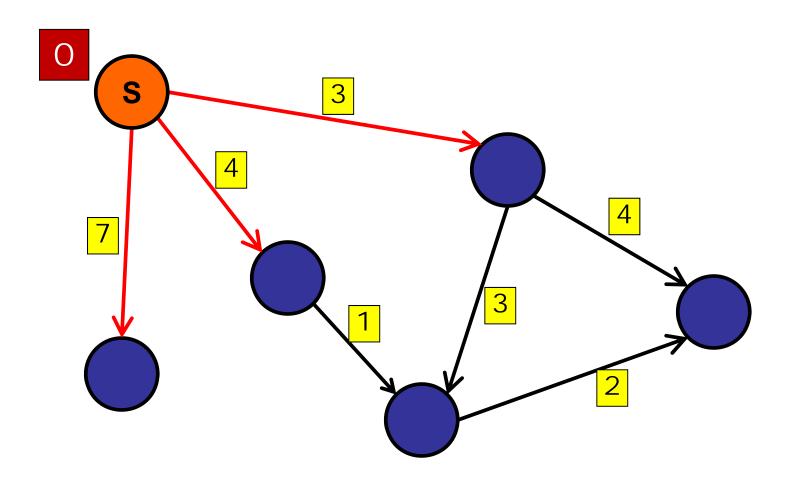
"The use of COBOL cripples the mind; its teaching should, therefore, be regarded as a criminal offense."

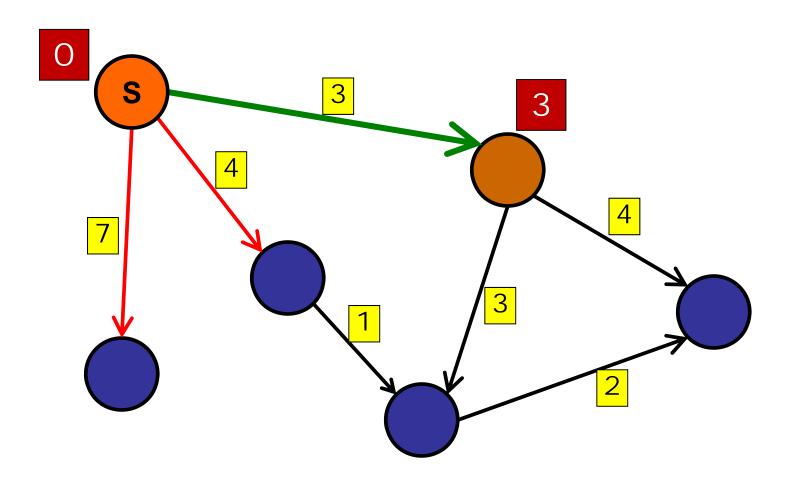
"APL is a mistake, carried through to perfection. It is the language of the future for the programming techniques of the past: it creates a new generation of coding bums."

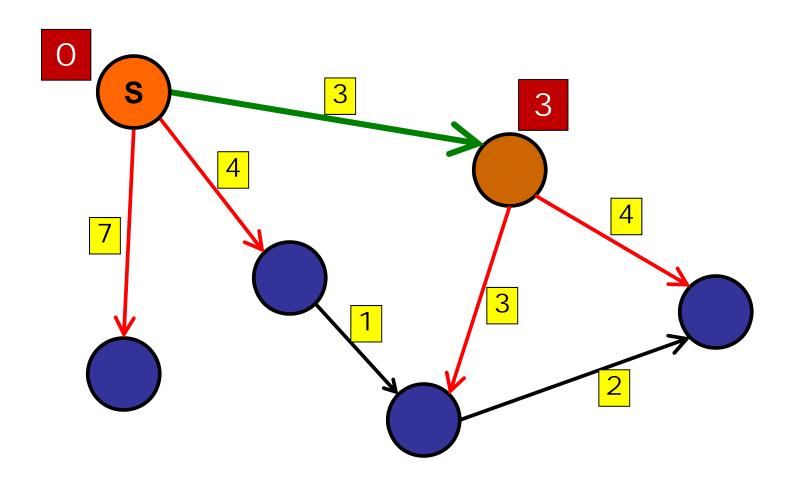
"Object-oriented programming is an exceptionally bad idea which could only have originated in California."



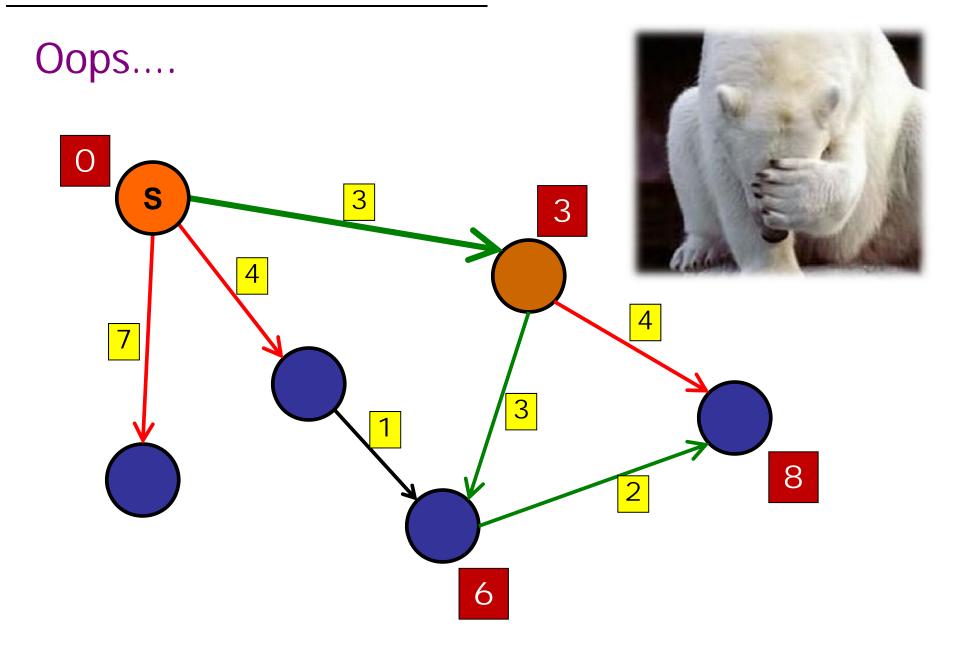








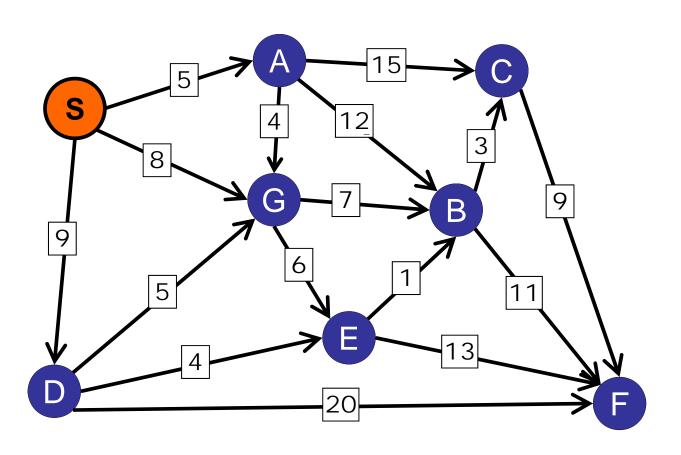
## Dijkstra's Algorithm (Failed Try)



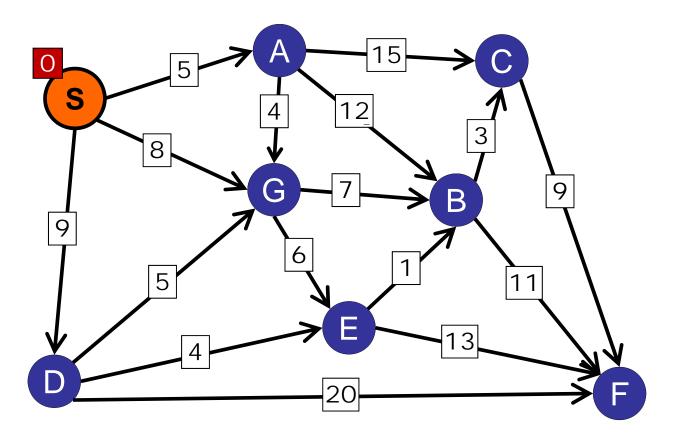
#### Basic idea:

- Maintain distance <u>estimate</u> for every node.
- Begin with empty shortest-path-tree.
- Repeat:
  - Consider vertex with minimum estimate.
  - Add vertex to shortest-path-tree.
  - Relax all outgoing edges.

### **Shortest Paths**

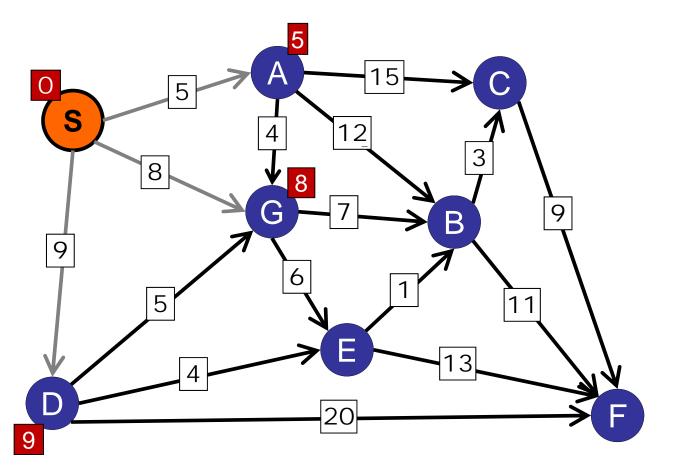


### Step 1: Add source



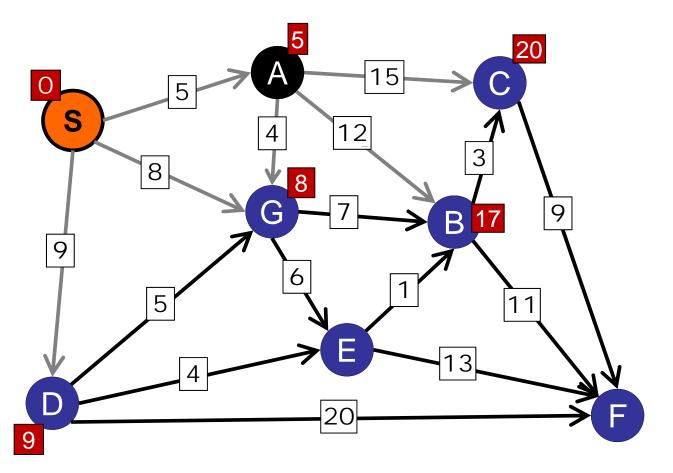
Vertex	Dist.
S	0

Step 2: Remove S and relax.



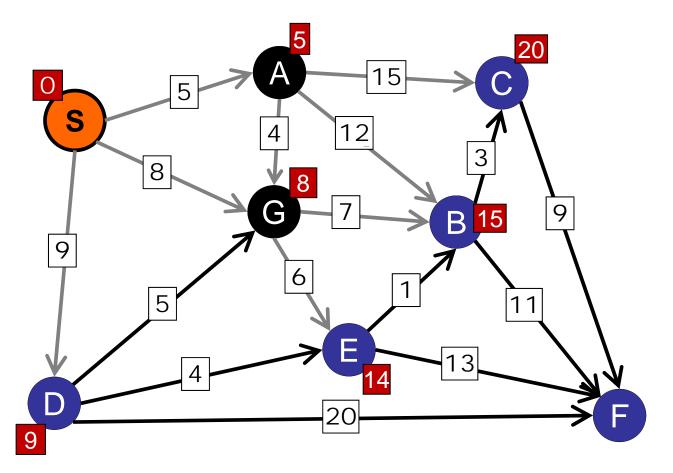
Vertex	Dist.
Α	5
G	8
D	9

Step 3: Remove A and relax.



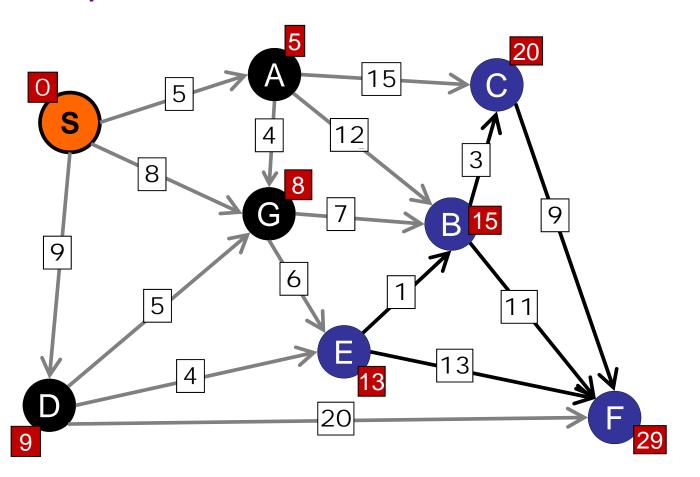
Vertex	Dist.
G	8
D	9
В	17
С	20

Step 4: Remove G and relax.



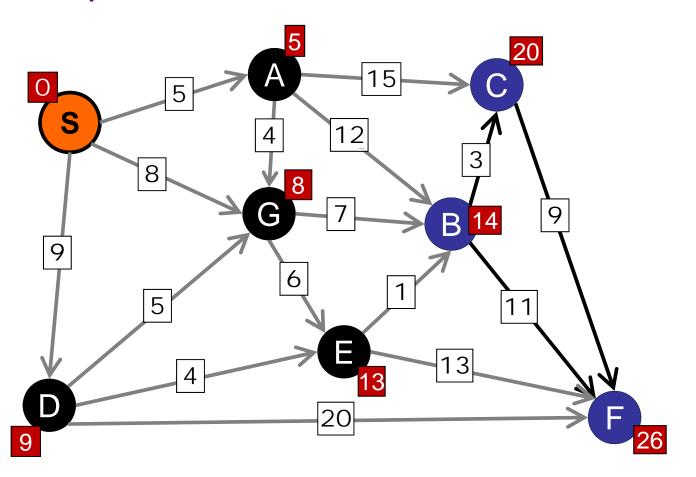
Vertex	Dist.
D	9
E	14
В	15
С	20

Step 5: Remove D and relax.



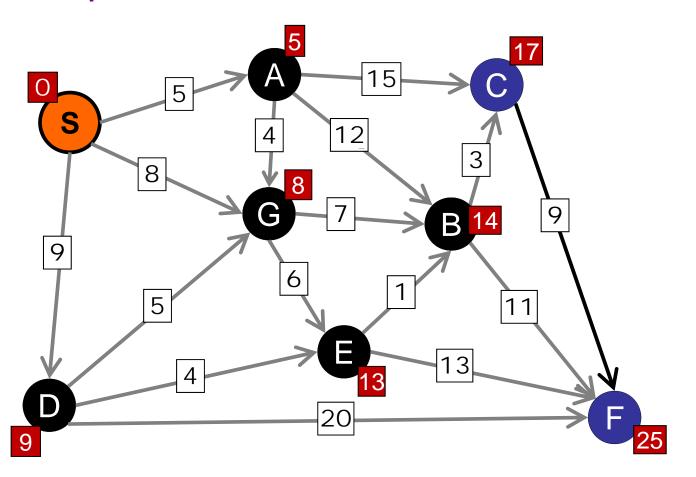
Vertex	Dist.
E	13
В	15
С	20
F	29

Step 5: Remove E and relax.



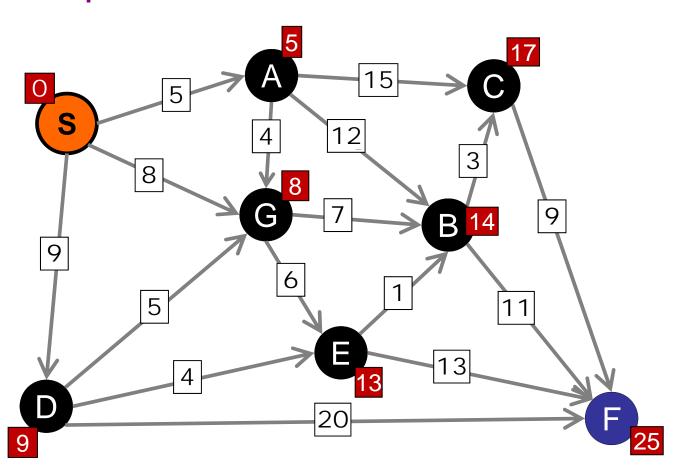
Vertex	Dist.
В	14
С	20
F	26

Step 5: Remove B and relax.



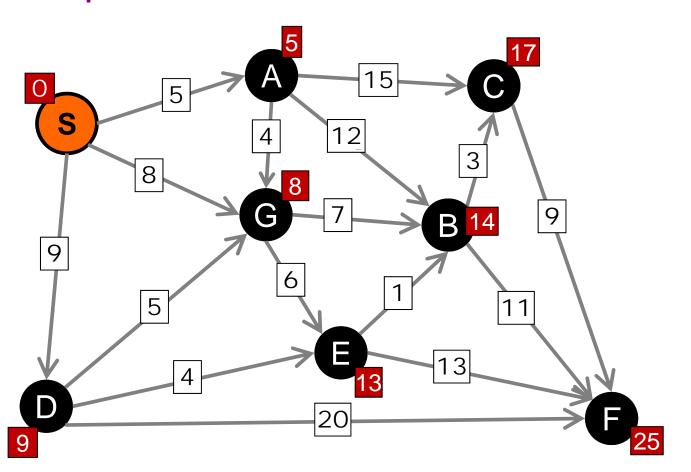
Vertex	Dist.
С	20
F	25

Step 5: Remove C and relax.



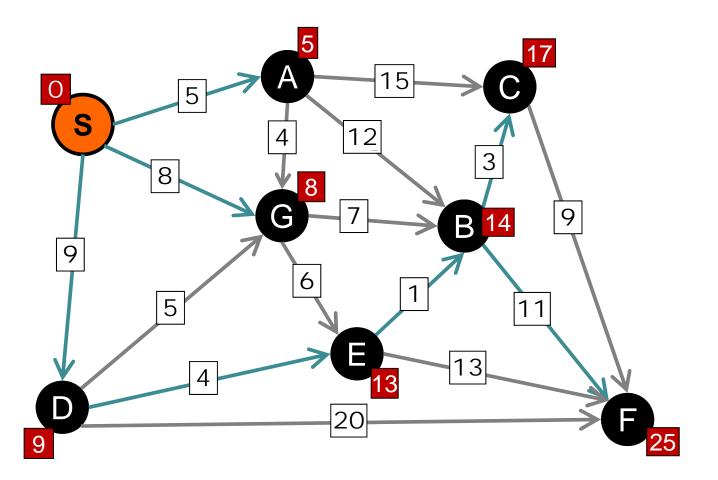
Vertex	Dist.
F	25

Step 5: Remove F and relax.



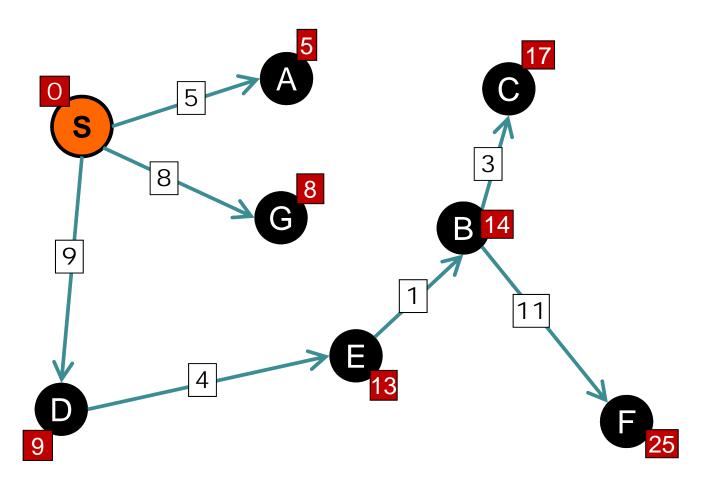


#### Done





#### **Shortest Path Tree**

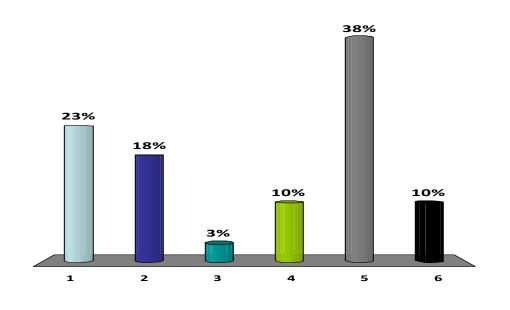




# What data structure to store vertices/distances?

- 1. Array
- 2. Linked list
- 3. Stack
- 4. Queue
- ✓5. AVL Tree
  - 6. Huh?

Vertex	Dist.
В	14
С	20
F	26



### Abstract Data Type

### Priority Queue

#### interface IPriorityQueue<Key, Priority>

```
void insert(Key k, Priority p)
                                         insert k with
                                         priority p
  Data extractMin()
                                         remove key with
                                         minimum priority
        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?
        isEmpty()
boolean
                                         is the priority queue
                                         empty?
```

#### Notes:

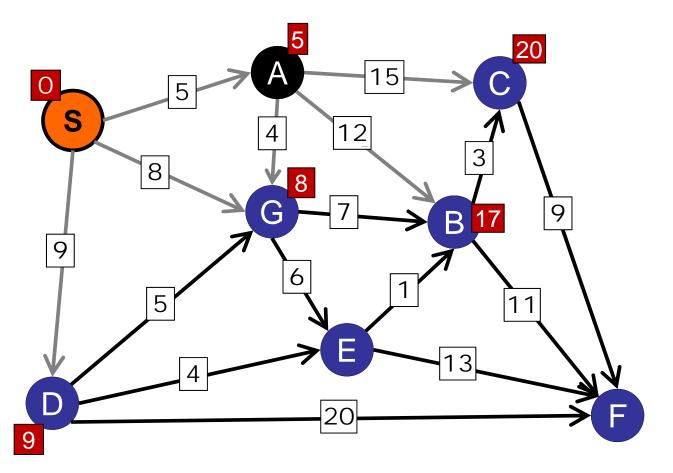
Assume data items are unique.

```
public Dijkstra{
     private Graph G;
     private IPriorityQueue pq = new PriQueue();
     private double[] distTo;
     searchPath(int start) {
           pq.insert(start, 0.0);
           distTo = new double[G.size()];
           Arrays.fill(distTo, INFTY);
           distTo[start] = 0;
           while (!pq.isEmpty()) {
                 int w = pq.deleteMin();
                 for (Edge e : G[w].nbrList)
                      relax(e);
```

```
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;
          if (pq.contains(w))
               pq.decreaseKey(w, distTo[w]);
          else
               pq.insert(w, distTo[w]);
```

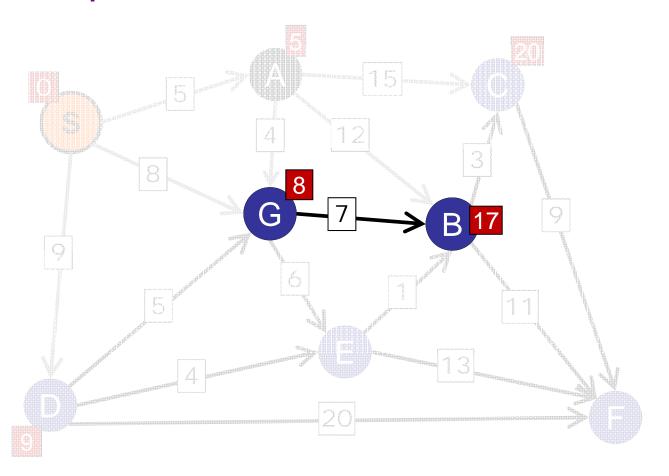
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          if (pq.contains(w))
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          else
               pq.insert(w, distTo[w]);
```

Step 3: Remove A and relax.



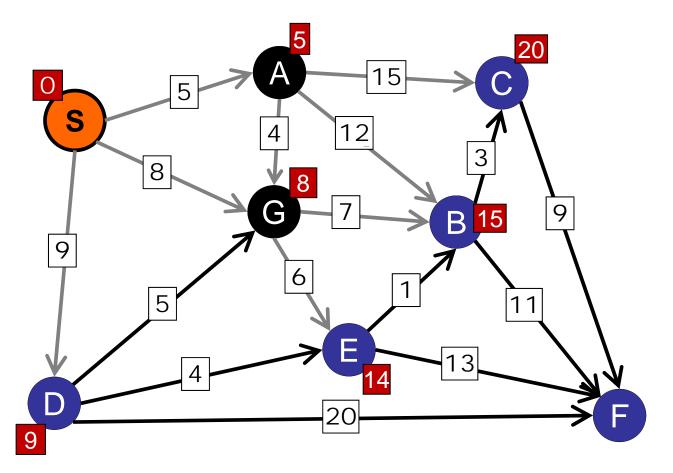
Vertex	Dist.
G	8
D	9
В	17
С	20

Step 3: Remove A and relax.



Vertex	Dist.
G	8
D	9
В	17
С	20

Step 4: Remove G and relax.



Vertex	Dist.
D	9
E	14
В	15
С	20

```
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;
          if (pq.contains(w))
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```

### Abstract Data Type

### Priority Queue

#### interface IPriorityQueue<Key, Priority>

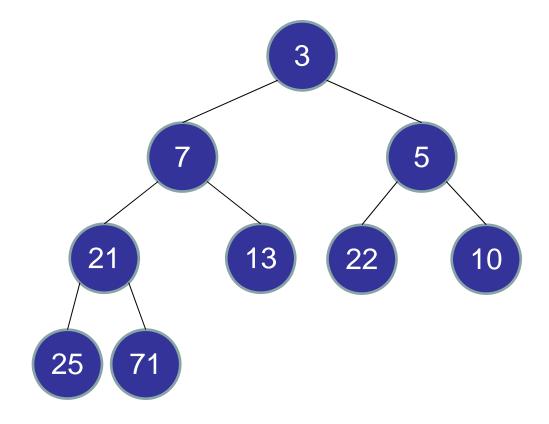
```
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boolean
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```

#### Notes:

Assume data items are unique.

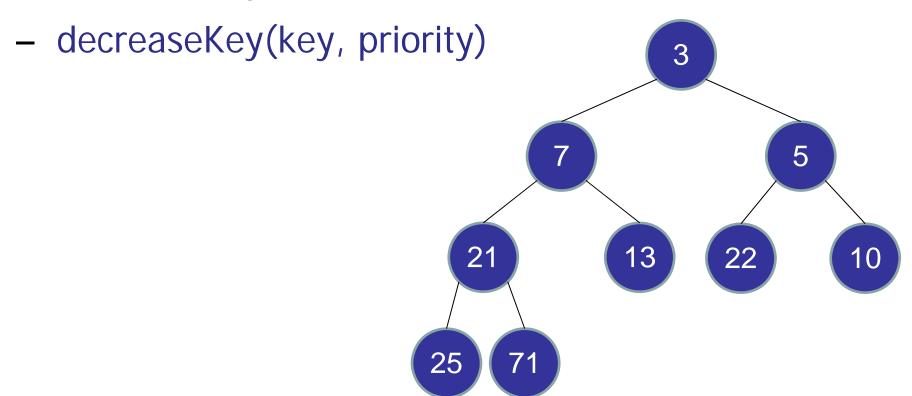
### Binary Heap

- Complete binary tree
- deleteMin: O(log n)
  - remove root
  - swap leaf to root
  - bubble down
- insert: O(log n)
  - add new leaf
  - bubble up



### Binary Heap

- How do we find a key? (Hint: not a search tree!)
- contains(key)



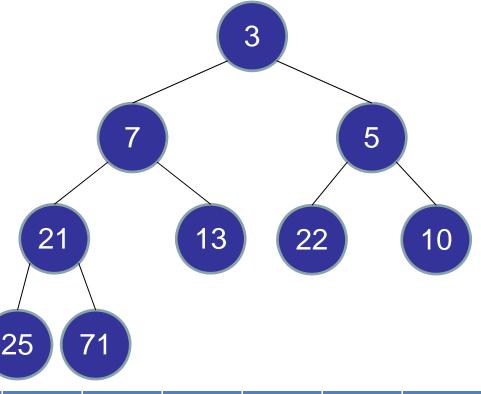
### Binary Heap

– decreaseKey(key, priority): O(logn)

#### – Hash Table:

 Map keys to locations in the binary tree.

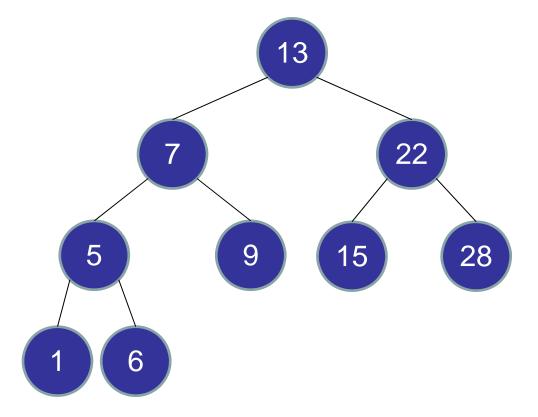
 Update hash table whenever the binary tree changes.



0	1	2	3	4	5	6	7	8	[9]	10	11
	3	7	6	21	13	22	10	25	71		

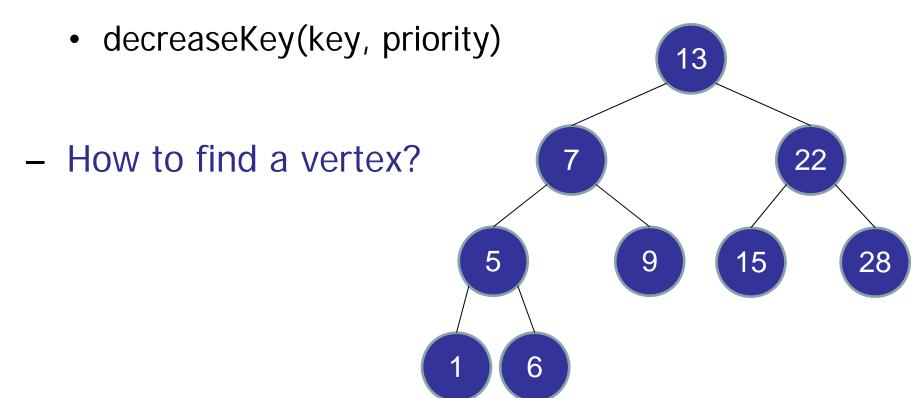
#### **AVL** Tree

- Indexed by: priority
- Existing operations:
  - deleteMin()
  - insert(key, priority)



#### **AVL** Tree

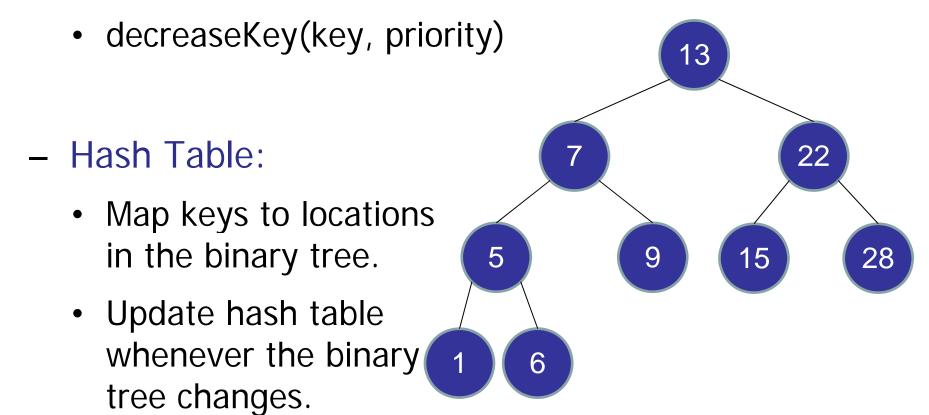
- Other operations:
  - contains()



### **Priority Queue**

### **AVL** Tree

- Other operations:
  - contains()

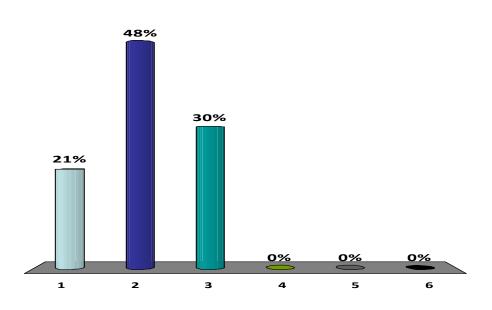


### Priority Queue by AVL tree:

- insert(key, priority): O(log n)
- deleteMin(): O(log n)
- decreaseKey(key, priority): O(log n)
- contains(key): O(1)

## What is the running time of Dijkstra's Algorithm, using an AVL tree Priority Queue?

- 1. O(V + E)
- ✓2. O(E log V)
  - 3. O(V log V)
  - 4.  $O(V^2)$
  - 5. O(VE)
  - 6. None of the above



```
public Dijkstra{
     private Graph G;
     private MinPriQueue pq = new MinPriQueue();
     private double[] distTo;
      searchPath(int start) {
           pq.insert(start, 0.0);
           distTo = new double[G.size()];
           Arrays.fill(distTo, INFTY);
           distTo[start] = 0;
while (!pq.isEmpty()) {
                                     / How many times?
                 int w = pq.deleteMin();
                 for (Edge e : G[w].nbrList)
                       relax(e);

How many times?
```

```
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;
          if (pq.contains(w))
               pq.decreaseKey(w, distTo[w]);
          else
               pq.insert(w, distTo[w]);
```

### **Analysis:**

- insert / 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:  $O((V+E)\log V) = O(E \log V)$

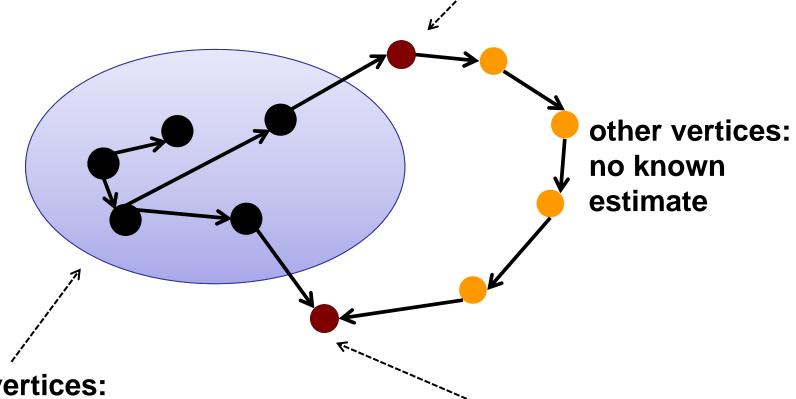
Why does it work?

### 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.

fringe vertices: neighbor of a finished vertex.

Every edge crossing the boundary has been relaxed.



finished vertices: distance is accurate.

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

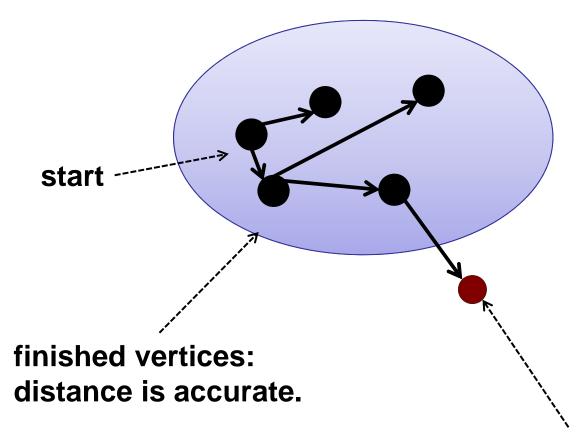
### Proof by induction:

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

### Proof by induction:

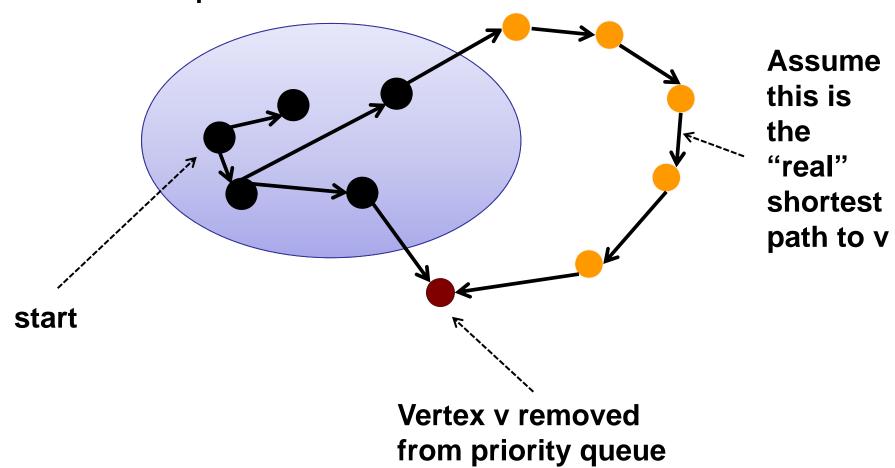
- Every "finished" vertex has correct estimate.
- Initially: only "finished" vertex is start.

- Inductive step:
  - Remove vertex from priority queue.
  - Relax its edges.
  - · Add it to finished.
  - Claim: it has a correct estimate.

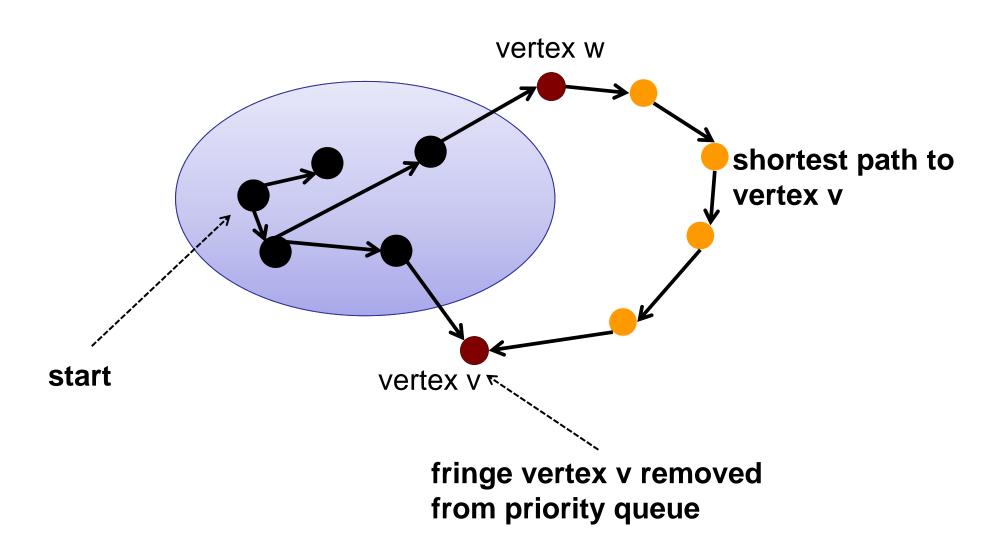


Vertex v going to be removed from priority queue next. Thus, with minimum distance amount the unfinished

Assume NOT. The current estimate is not the shortest path.

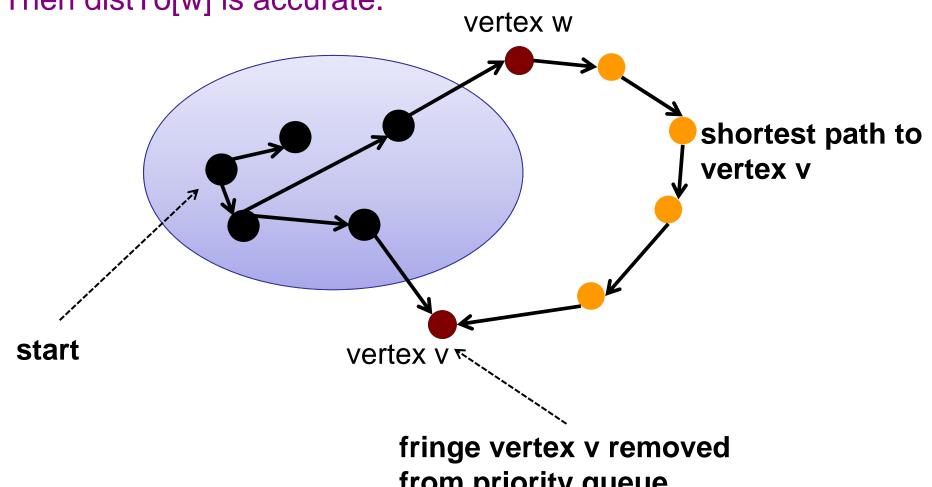


There must be a vertex w in the current PQ on this "real" path.

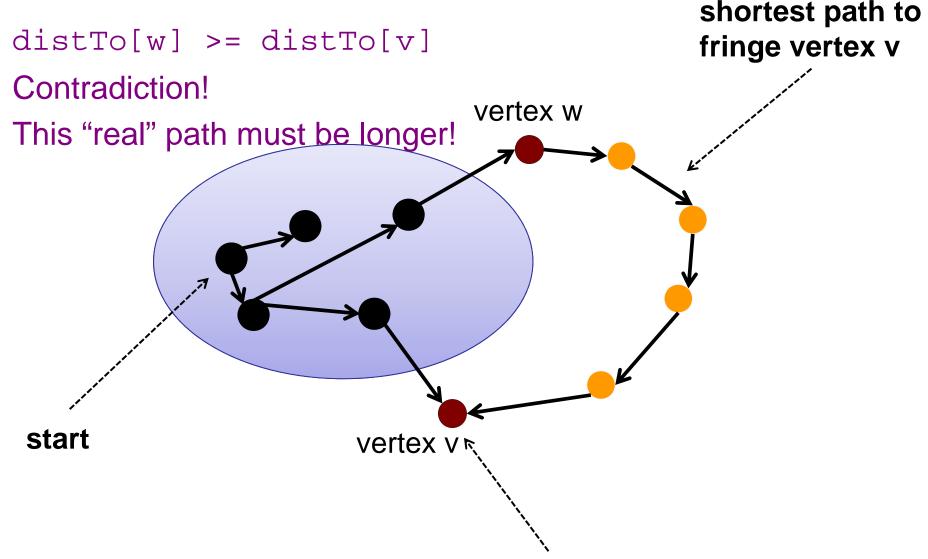


If P is shortest path to v, then prefix of P is shortest path to w.

Then distTo[w] is accurate.



from priority queue



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

### 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.

```
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;
          if (pq.contains(w))
               pq.decreaseKey(w, distTo[w]);
          else
               pq.insert(w, distTo[w]);
```

### **Analysis:**

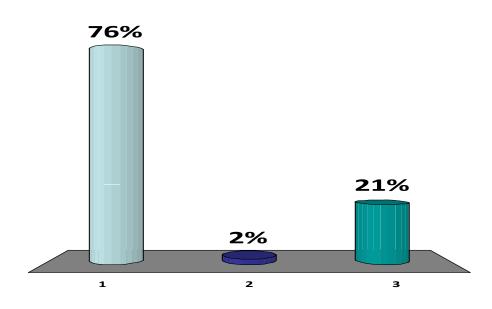
- insert / deleteMin: |V| times each
  - Each node is added to the priority queue once.

- decreaseKey: |E| times
  - Each edge is relaxed once.
- Priority queue operations: O(log V)

- Total:  $O((V+E)\log V) = O(E \log V)$ 

# Source-to-Destination Dijkstra Can we stop as soon as we dequeue the destination?

- ✓1. Yes.
  - 2. Only if the graph is sparse.
  - 3. No.



### 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

### Basic idea:

- Maintain distance estimates.
- Repeat:
  - Find unfinished vertex with smallest estimate.
  - Relax all outgoing edges.
  - Mark vertex finished.

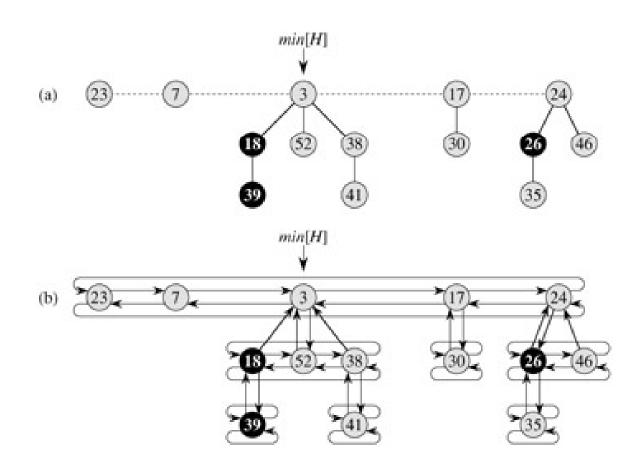
O(E log V) time (with AVL tree).

## Dijkstra's Performance

PQ Implementation	insert	deleteMin	decreaseKey	Total
Array	1	V	1	O(V <sup>2</sup> )
AVL Tree	log V	log V	log V	O(E log V)
d-way Heap	dlog <sub>d</sub> V	dlog <sub>d</sub> V	log <sub>d</sub> V	O(Elog <sub>E/V</sub> V)
Fibonacci Heap	1	log V	1	O(E + V log V)

## Fibonacci Heap

• Not in CS2020

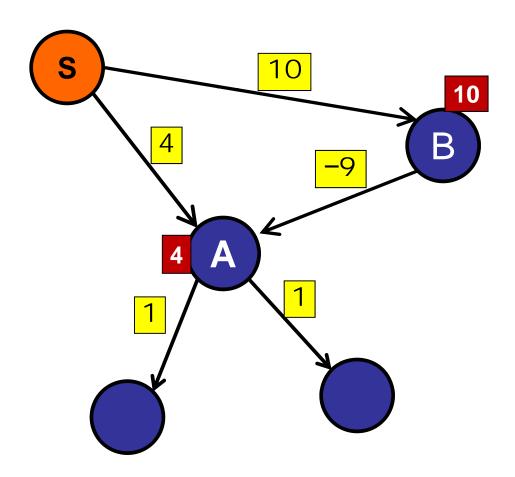


## Dijkstra Summary

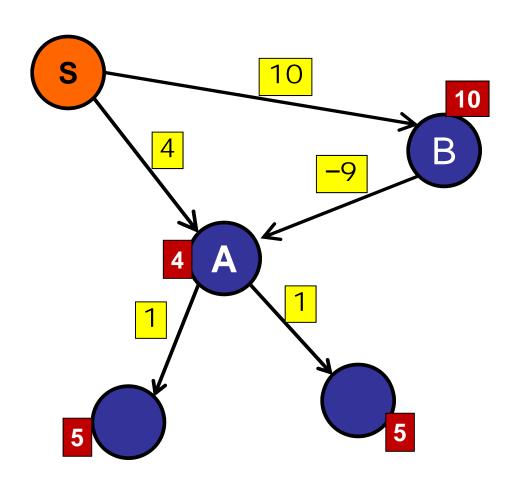
Edges with negative weights?

shortest path to What goes wrong with negative weights? fringe vertex v vertex w start vertex v fringe vertex v removed from priority queue

Edges with negative weights?

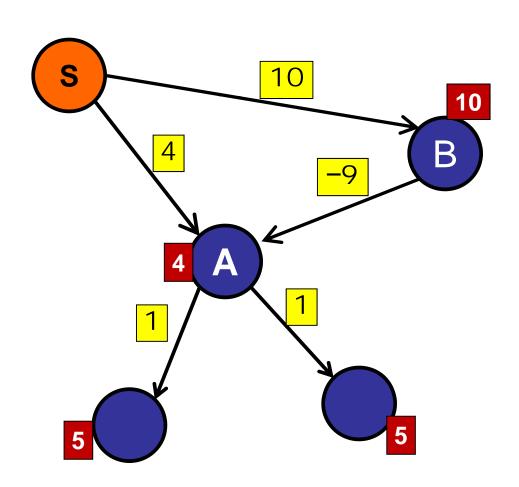


Edges with negative weights?



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

### Edges with negative weights?



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

. . .

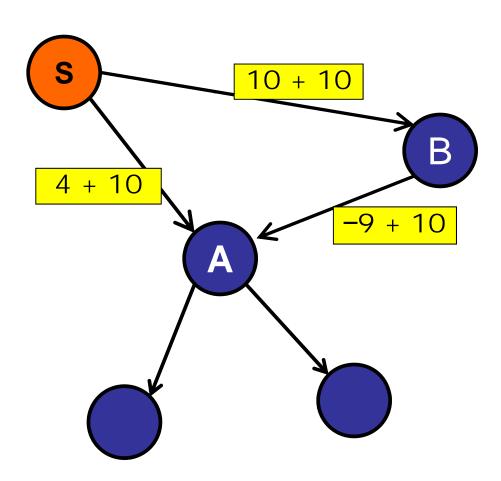
Step 4: Remove B.
Relax B.
Mark B done.

Oops: We need to update A.

shortest path to What goes wrong with negative weights? fringe vertex v vertex w start vertex v fringe vertex v removed from priority queue

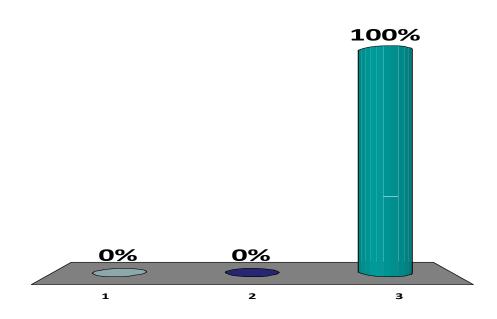
Can we reweight?

e.g.: weight +=10

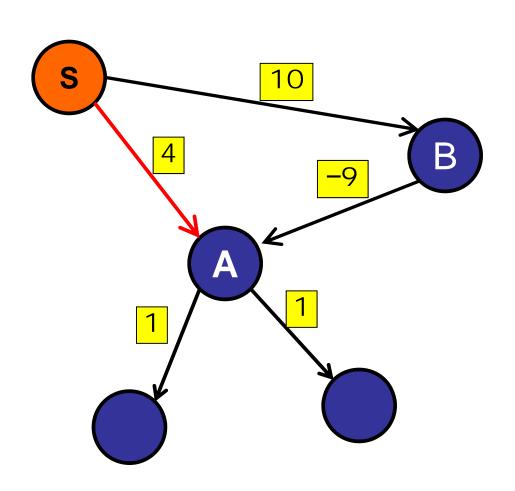


### Can we reweight the graph?

- 1. Yes.
- 2. Only if there are no negative weight cycles.
- **✓**3. No.



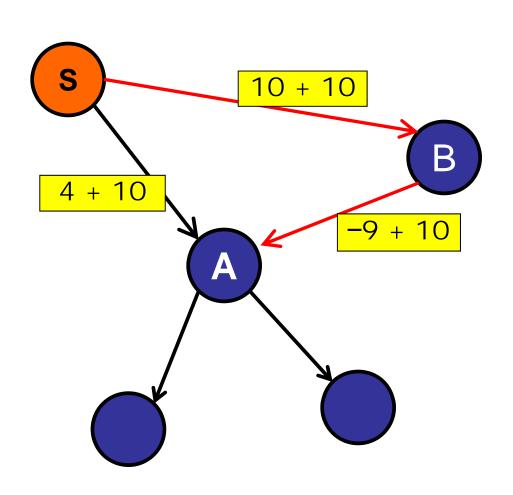
Can we reweight?



Path S-B-A: 1

Path S-A: 4

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.

## Roadmap

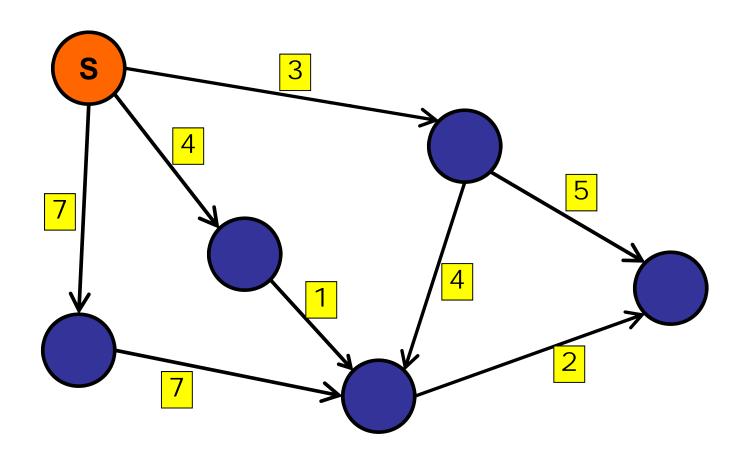
#### Part I: Shortest Paths

- Special Case: Tree
- Special Case: Non-negative weights (Dijkstra's)
- Special Case: Directed Acyclic Graphs

#### Part II: Applications of Shortest Paths

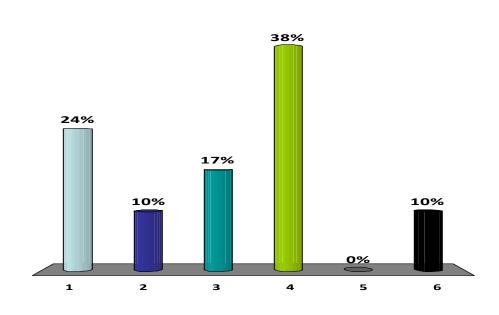
- DNA Alignment
- Constraint Systems

Acyclic Graph: Suppose the graph has no cycles.

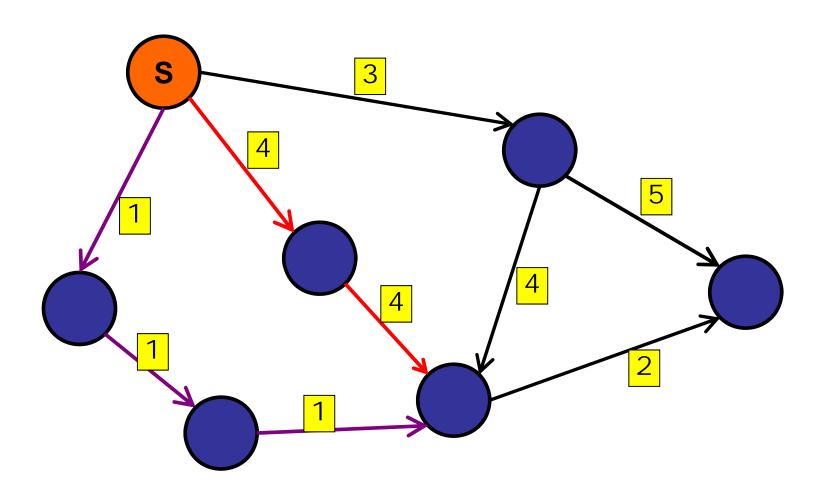


#### What order should we relax the nodes?

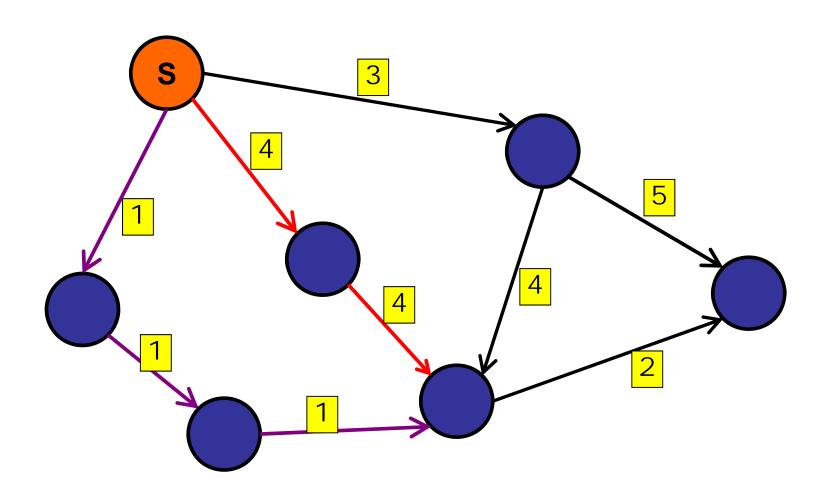
- 1. BFS
- 2. DFS pre-order
- ✓3. DFS post-order
  - 4. Shortest edge
  - 5. Longest edge
  - 6. Other

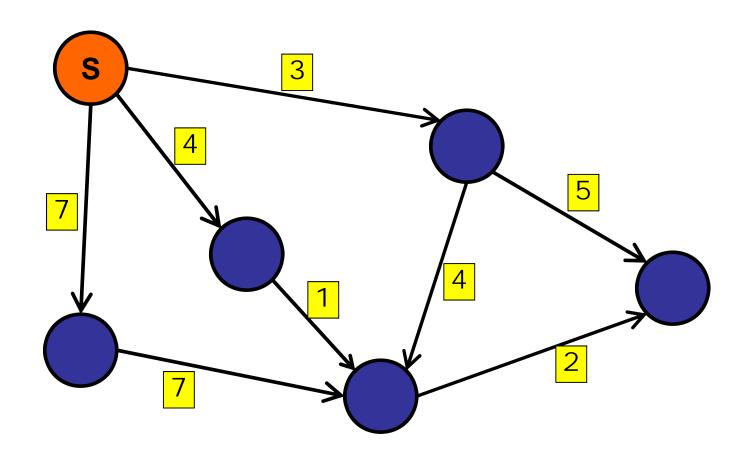


Acyclic Graph: Not BFS.



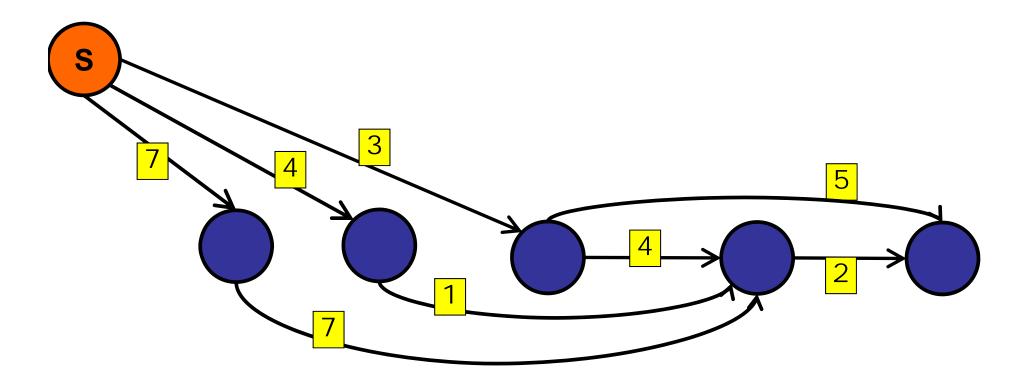
Acyclic Graph: Not DFS-preorder.



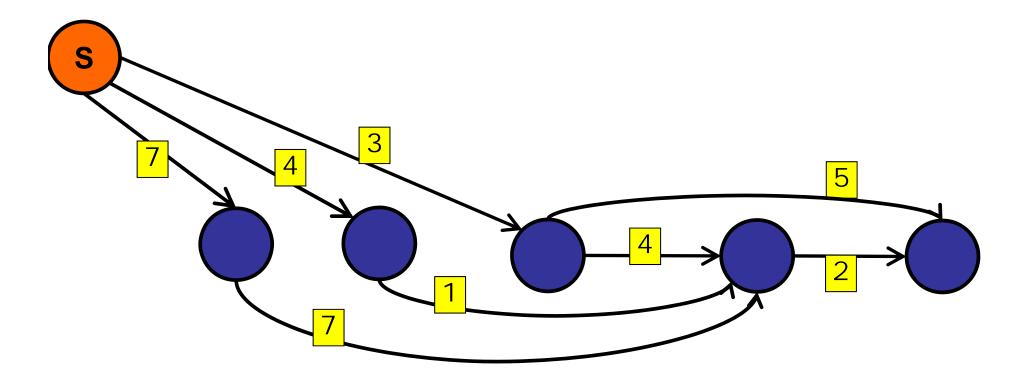


Acyclic Graph: has no cycles.

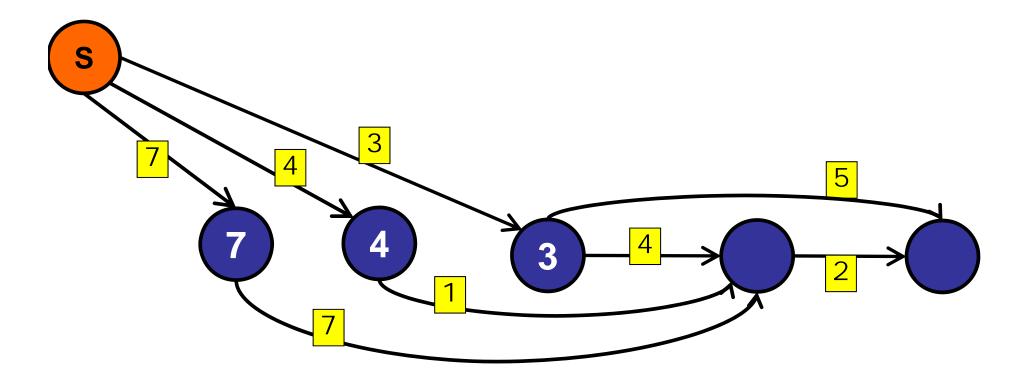
1. Topological sort



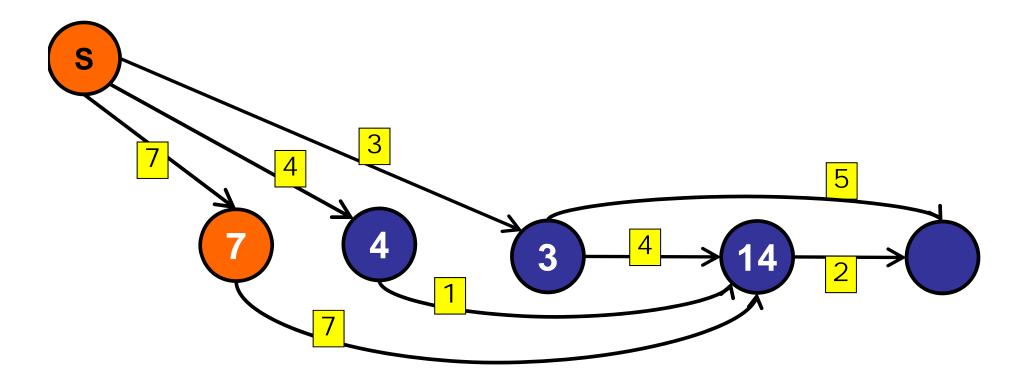
- 1. Topological sort
- 2. Relax in order.



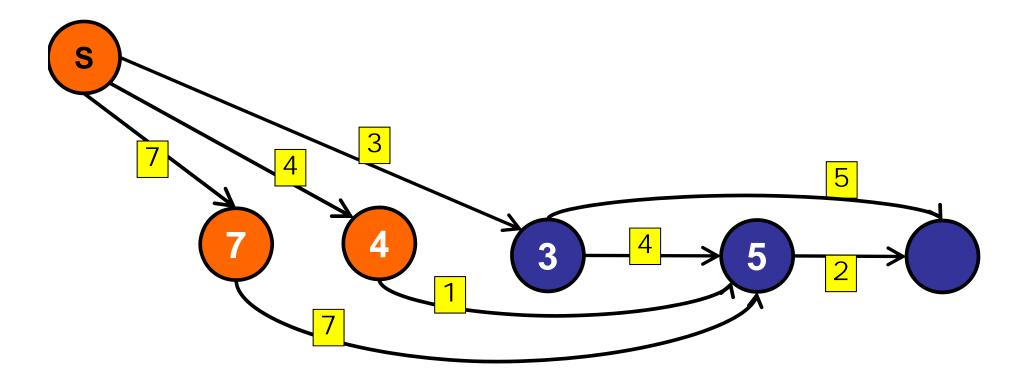
- 1. Topological sort
- 2. Relax in order.



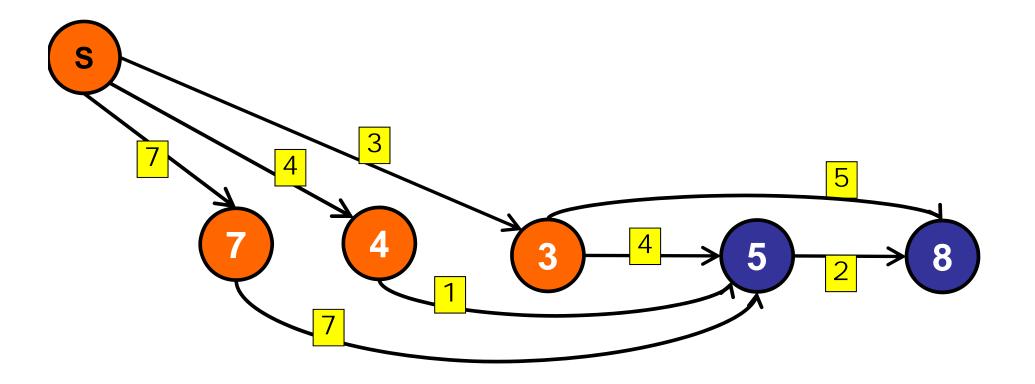
- 1. Topological sort
- 2. Relax in order.



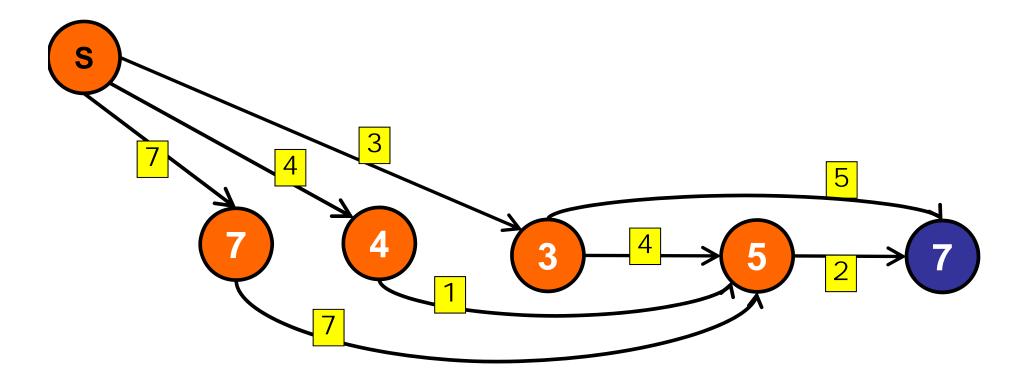
- 1. Topological sort
- 2. Relax in order.



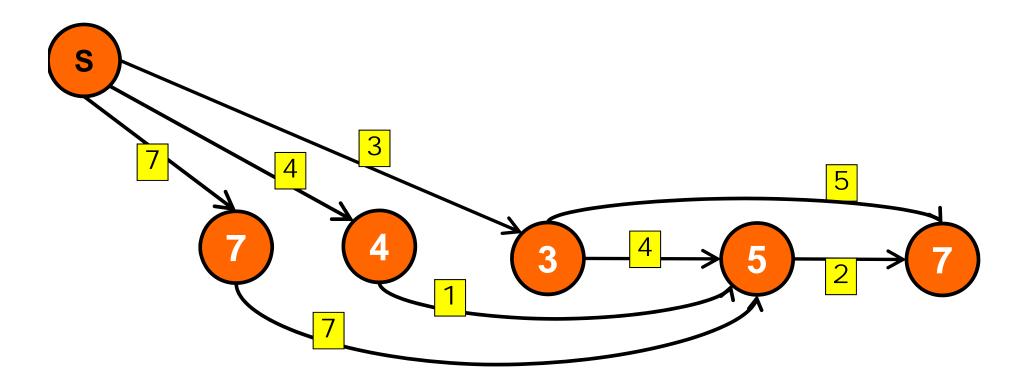
- 1. Topological sort
- 2. Relax in order.

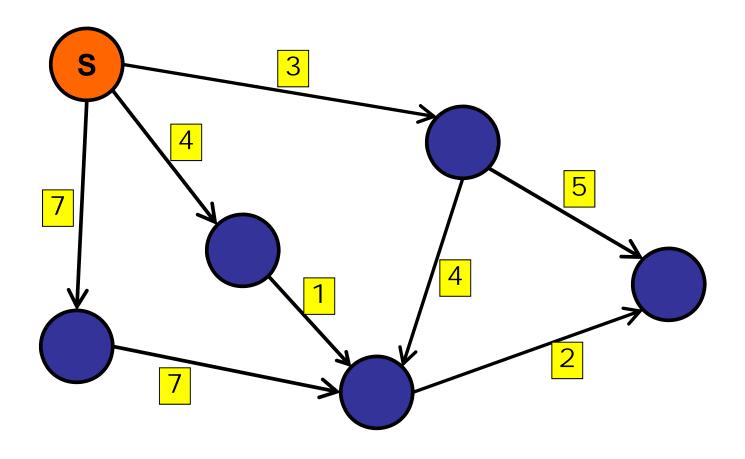


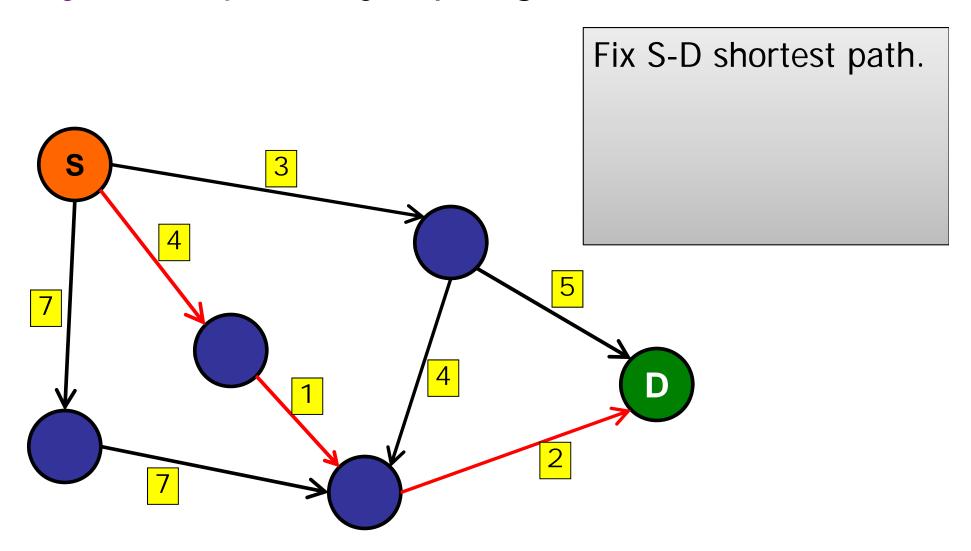
- 1. Topological sort
- 2. Relax in order.

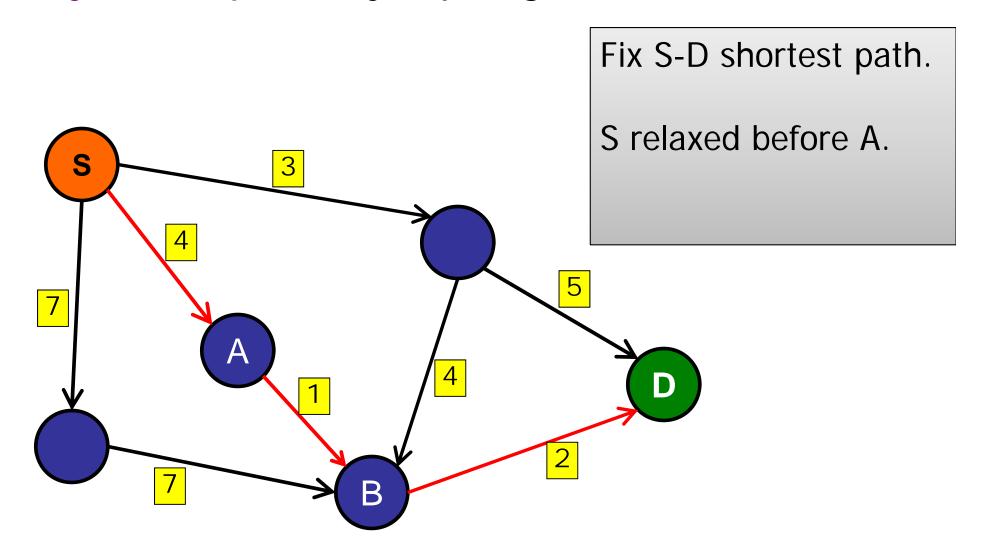


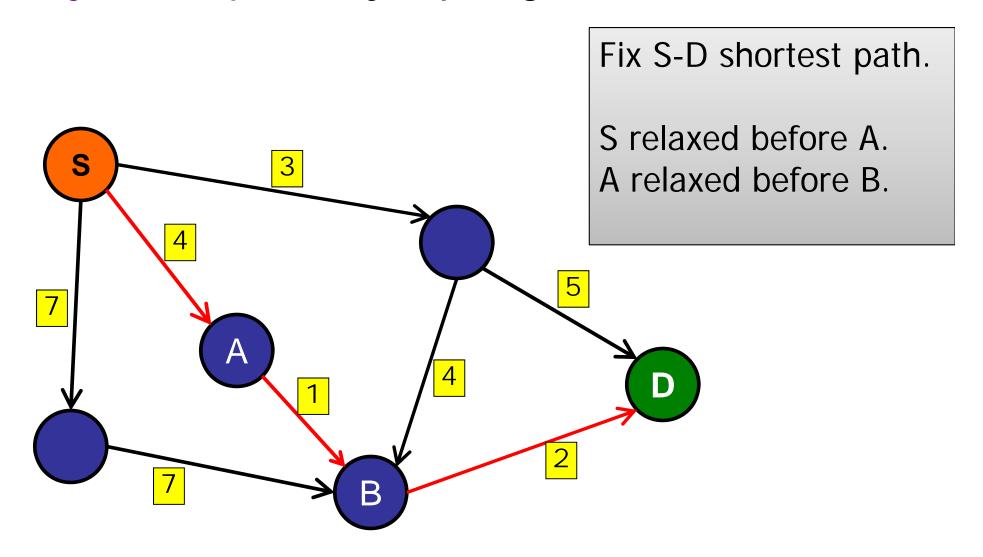
- 1. Topological sort
- 2. Relax in order.

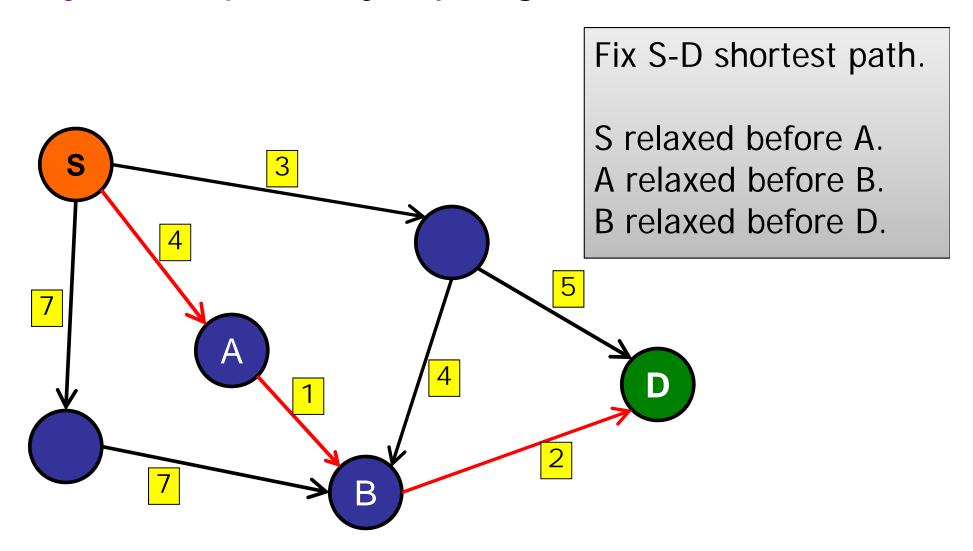


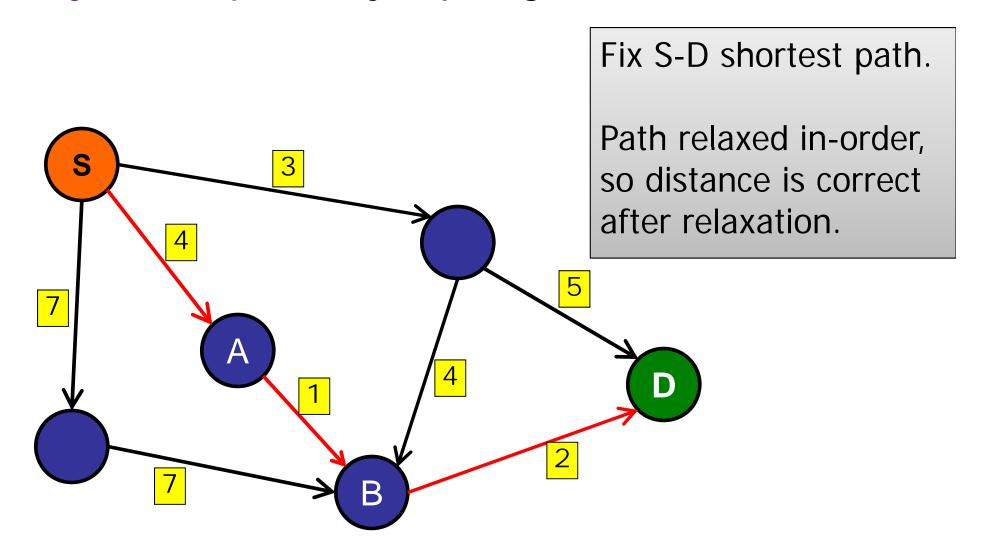






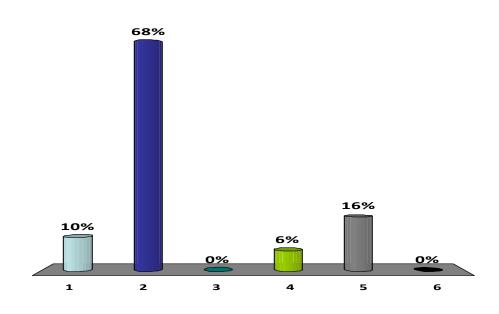




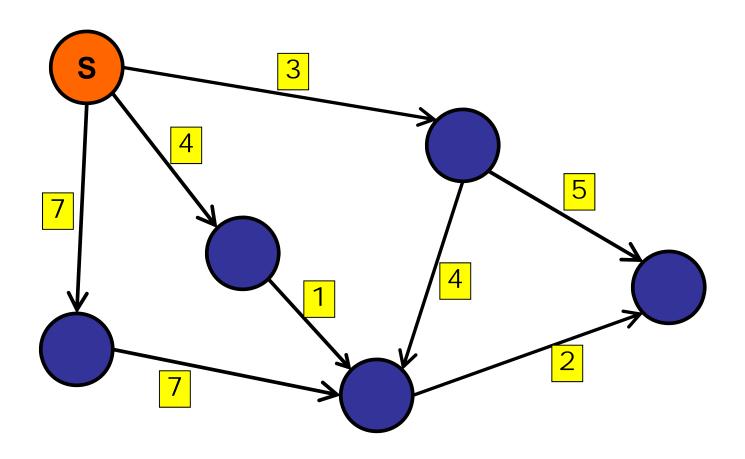


# What is the running time of shortest paths on a DAG?

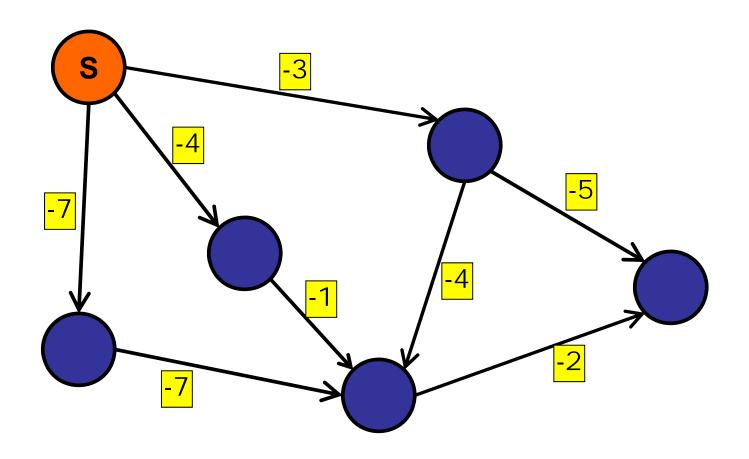
- 1. O(V)
- **✓**2. O(E)
  - 3.  $O(V^2)$
  - 4. O(E log V)
  - 5. O(V log E)
  - 6. O(VE)



Acyclic Graph: Any ideas?

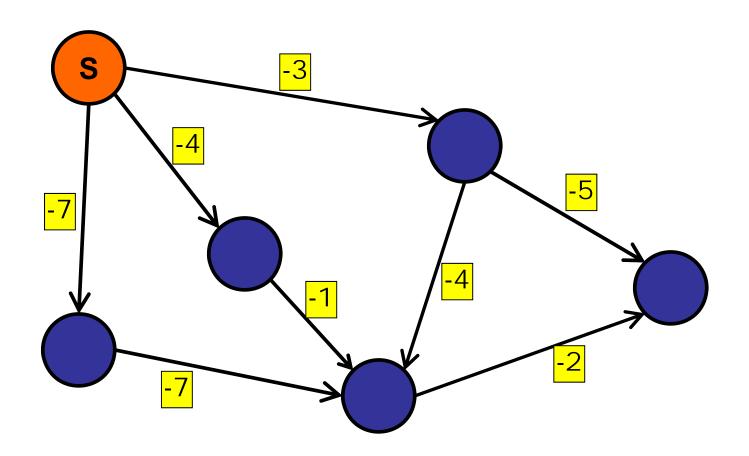


Acyclic Graph: Negate the edges!

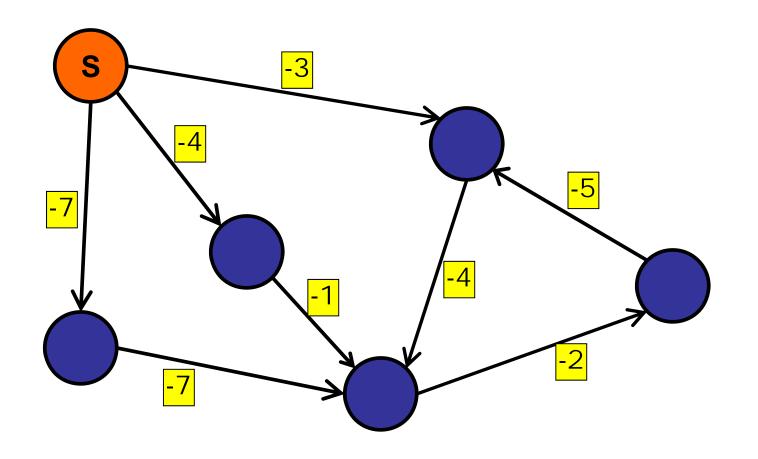


#### Acyclic Graph:

shortest path in negated=longest path in regular



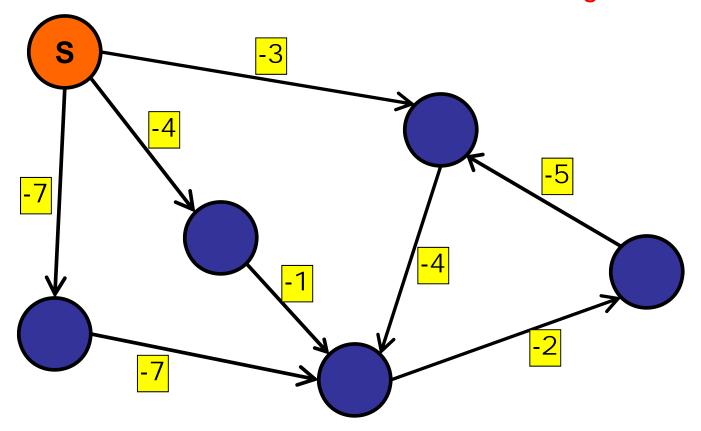
General (cyclic) Graph: (positive weights)
Can we use the same trick?



General (cyclic) Graph: (positive weights)

Can we use the same trick? NO

Negative weight cycles!

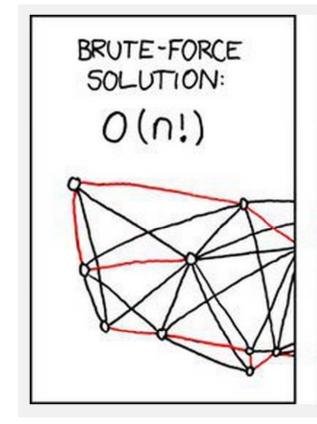


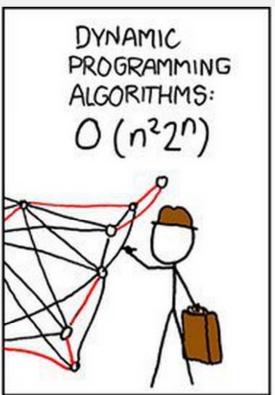
#### Directed Acyclic Graph:

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.

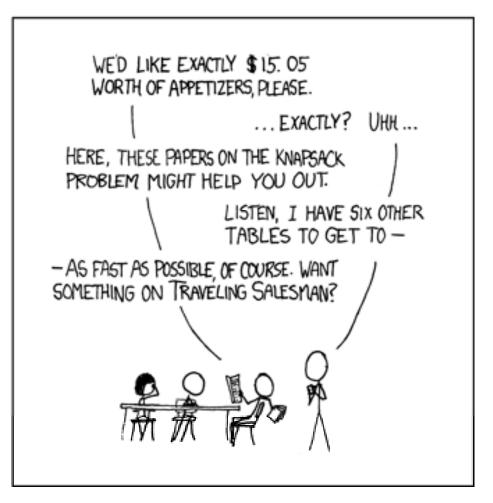






MY HOBBY: EMBEDDING NP-COMPLETE PROBLEMS IN RESTAURANT ORDERS

CHATAIL	
CHOTCHKIES RESTAURANT	
APPETIZER	N
MUXED FRUIT	2.15
FRENCH FRIES	2.75
SIDE SALAD	3.35
HOT WINGS	3.55
MOZZARELLA STICKS	4.20
SAMPLER PLATE	5.80
→ SANDWICHES →	
RARRECHE	6 55



## Roadmap

#### Part I: Shortest Paths

- Special Case: Tree
- Special Case: Non-negative weights (Dijkstra's)
- Special Case: Directed Acyclic Graphs

#### Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

#### Input: two DNA strings:

- AGGAACCGTA
- AGAATCCGAA

### How similar are they?

Metric: edit distance

How many operations to transform one DNA string into another?

#### Input: two DNA strings:

- AGGAACCGTA ← delete G, delete T
- AGAATCCGA ← add T

#### Three operations:

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

#### Input: two DNA strings:

- AGGAACCGTA ← delete G, delete T
- AGAATCCGA ← 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?

#### 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

Transform: CG to AGT

start ACG AGCG AGTCG CG **AGTG** AG AGG G G AG **AGT** A

Vertical: delete character

Horizontal: add character

Diagonal: transform character

#### CG to AGT

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

start ACG AGCG CG **AGTG** AG AGG G G AG **AGT** A

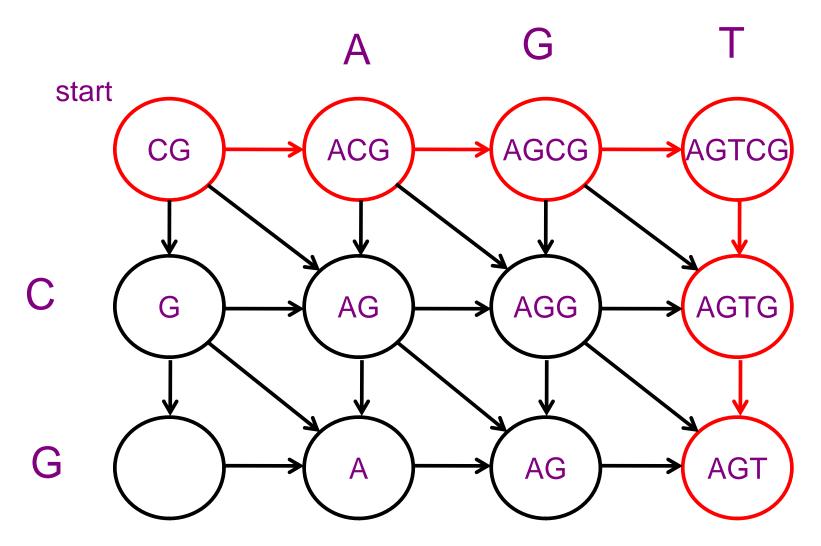
Vertical: delete character

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

Transform: CG to AGT

start ACG AGCG CG AGTCG **AGTG** AG AGG G G AG **AGT** A

Vertical: delete character

Horizontal: add character

Diagonal: transform character

#### CG to AGT

Edge costs:

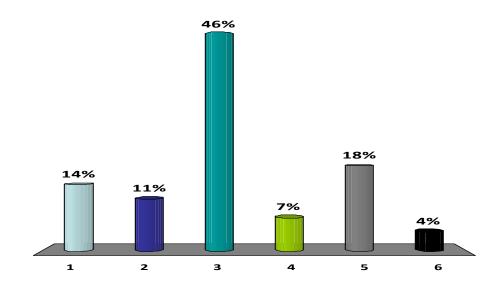
start a a a ACG AGCG CG AGTCG a a AGG AGTG AG G G AG **AGT** A a a a

Vertical: delete character

Horizontal: add character

Diagonal: transform character What is the running time to find the minimum edit distance from a string of length n to a string of length n?

- 1. O(n)
- 2. O(n log n)
- **✓**3. O(n²)
  - 4. O(n<sup>2</sup>log n)
  - 5.  $O(n^3)$
  - 6.  $O(n^4)$



# Example: DNA Alignment

Transform: CG to AGT

start ACG AGCG CG AGTCG **AGTG** AG AGG G G AG **AGT** A

Vertical: delete character

Horizontal: add character

Diagonal: transform character

### Roadmap

#### Part I: Shortest Paths

- Special Case: Tree
- Special Case: Non-negative weights (Dijkstra's)
- Special Case: Directed Acyclic Graphs

#### Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems

#### Input:

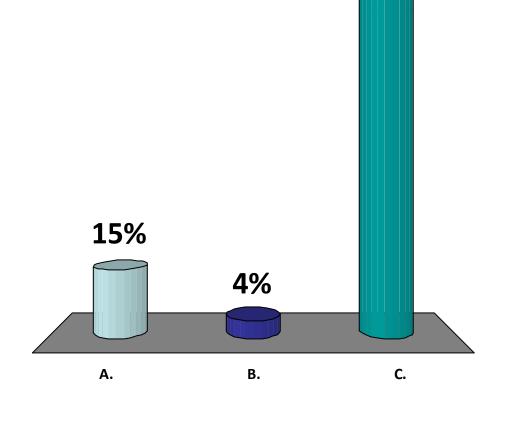
- Set of tasks: A, B, C, D, E, F
- Constraints:
  - A must be done at least 10 minutes before C
  - D must be done at most 20 minutes after E
  - B must be done after F

#### Output:

- Feasible?
- Schedule?

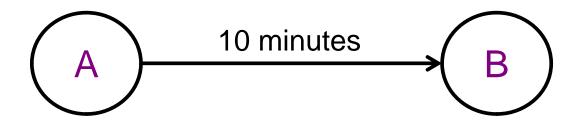
We can assume there is no negative cycle because...

- A. we said so
- B. our algorithms cannot solve it
- C. Negative cycles make the scheduling problem meaningless



81%

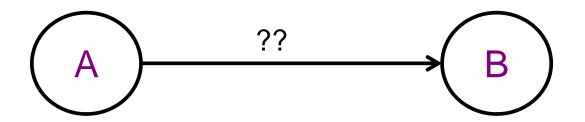
B must be executed at most 10 minutes after A



Shortest path = schedule time

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

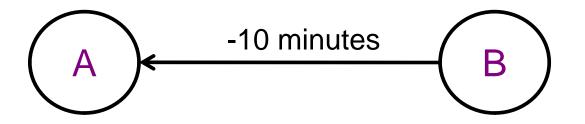
B must be executed at least 10 minutes after A



Shortest path = schedule time

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

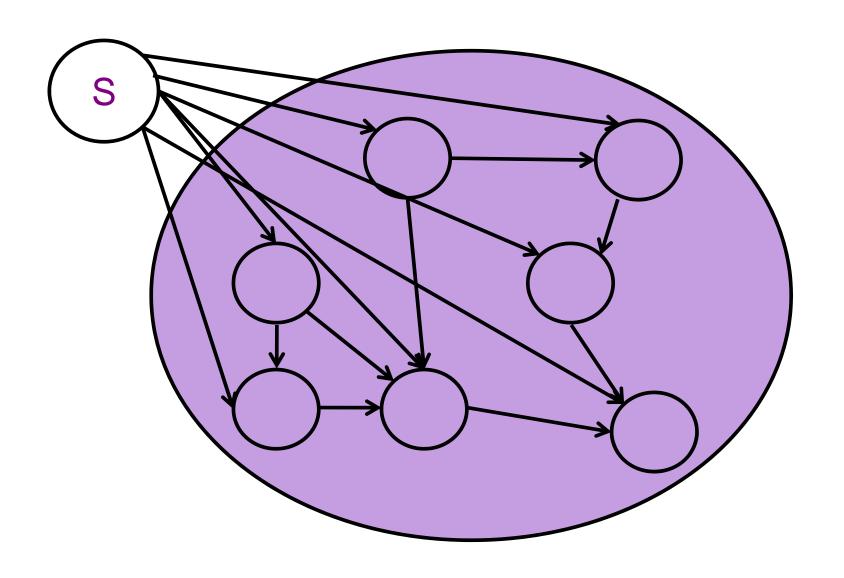
B must be executed at least 10 minutes after A



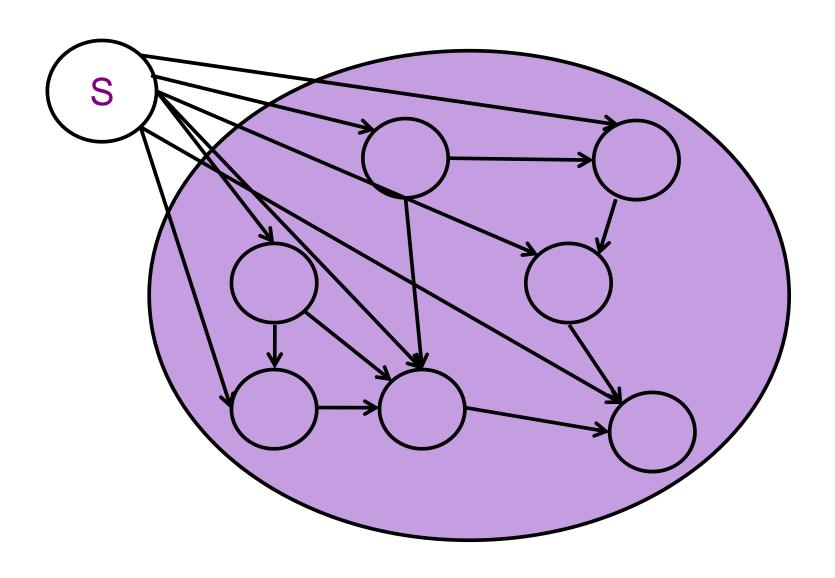
Shortest path = schedule time

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

Add source S connected by 0 weight edges to all.

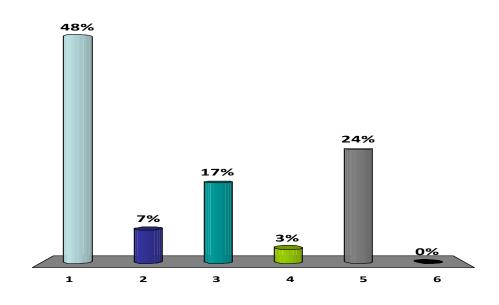


Solve shortest paths.

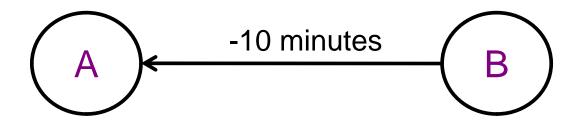


# What is the running time to find the schedule for n jobs and m constraints?

- 1. O(n + m)
- 2. O(n log m)
- 3. O(m log n)
- 4.  $O(n^2)$
- **✓**5. O(nm)
  - 6. O(n<sup>m</sup>)



B must be executed at least 10 minutes after A



Negative edges: use Bellman-Ford!

Running time: O(nm)

#### 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

#### Part I: Shortest Paths

- Special Case: Tree
- Special Case: Non-negative weights (Dijkstra's)
- Special Case: Directed Acyclic Graphs

#### Part II: Applications of Shortest Paths

- DNA Alignment
- Constraint Systems