# Efficient algorithms and data structures

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#### Course

- What the course is:
  - a selection of topics on the design and analysis of algorithms
  - with emphasis on rigorous analysis (Ph.Flajolet: "mathematically oriented engineering")
  - dealing with basic data structures (graphs, strings, trees, tables, ...)
  - including programming assignments and in-class projects
- What the course is not:
  - a programming course
  - a course oriented to a specific programming language (an imperative programming language is assumed, one of Python, C, C++, Java)
  - a course oriented to a specific application area
  - a math course

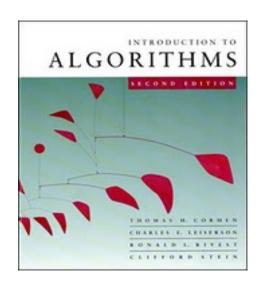
#### Course

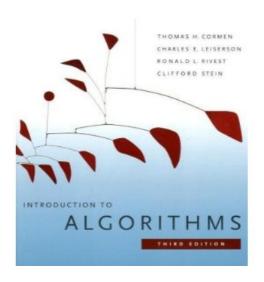
- Varying level of difficulty
- Prerequisites:
  - ▶ imperative programming (C, C++, Java, ...)
  - Basic data structures: lists, arrays, stacks, queues
  - Recursion, Big-Oh notation
  - Sorting, ...
- "Free-style" pseudo-code
- Having a laptop assumed

# Grading

- participation in class 10%
  - full attendance is expected
  - in-class projects
- programming exercises 40%
  - ▶ one every ~2 weeks
  - plagiarism is not tolerated
- exam 50%

#### Useful books







CLRS = Cormen & Leiserson & Rivest & Stein

#### Some other good algorithm textbooks:

- Steven Skiena, The Algorithm Design Manual, 2nd Edition, Springer, 2008 [a bit advanced?]
- Jon Kleinberg and Éva Tardos, Algorithm Design, MIT Press 2005
- Robert Sedgewick and Kevin Wayne, Algorithms, Addison-Wesley, 4th Edition, 2011
   [for beginners, Java-oriented]

#### How to measure the efficiency of algorithms?

- Efficiency (in this course) = TIME and SPACE
  - other possible measure of efficiency: accuracy
- Classical model: RAM model of computation
  - all memory accesses have equal cost
  - no parallel execution
  - unit cost (O(1)) of basic operations (unless we want to explicitly count individual bits operations)
  - > space = # of computer words (unless bit complexity is considered); each computer word contains  $\Theta(\log n)$  bits
  - other possible measures can be considered: disk accesses, cache misses, probe model, query complexity ...

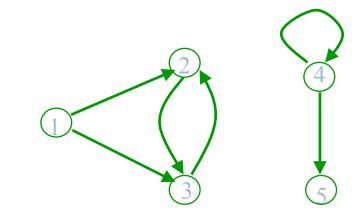
#### How to measure the efficiency of algorithms?

- Algorithms solve mass problems
  - n: input size (in computer words or bits)
  - time/space as a function of *n*
- Different complexity analyses:
  - worst-case complexity
  - average-case complexity
  - smoothed analysis
  - query (probe) complexity
  - • •

# Graphs

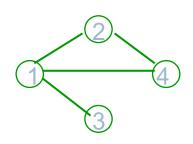
# Graphs

Directed graph G = (V, E) V finite set of nodes (vertices)  $E \subseteq V \times V$  set of edges (arcs), i.e., a relation on V



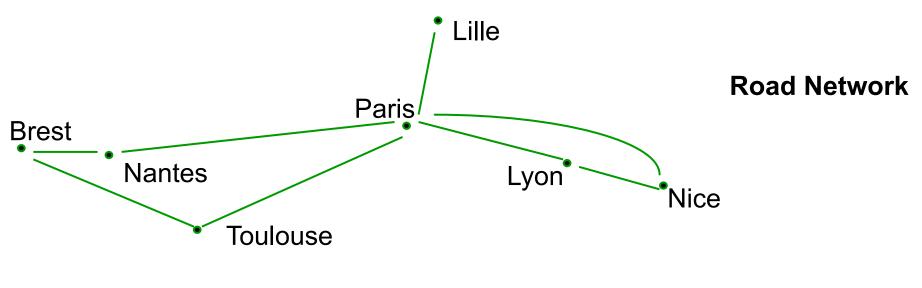
$$V = \{ 1, 2, 3, 4, 5 \}$$
  
 $E = \{ (1, 2), (1, 3), (2, 3), (3, 2), (4, 4), (4, 5) \}$ 

Undirected graph G = (V, E) E set of edges (arcs), symmetric relation



$$E = \{ 1, 2, 3, 4 \}$$
  
 $V = \{ \{1, 2\}, \{1, 3\}, \{1, 4\}, \{2, 4\} \}$ 

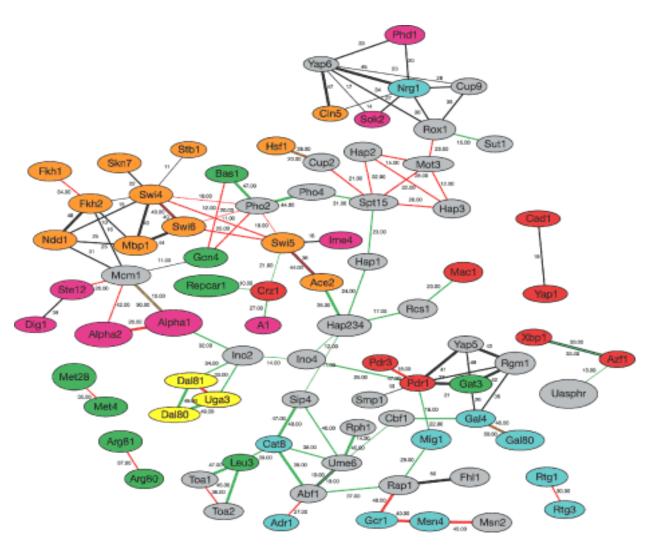
# Graphs are everywhere



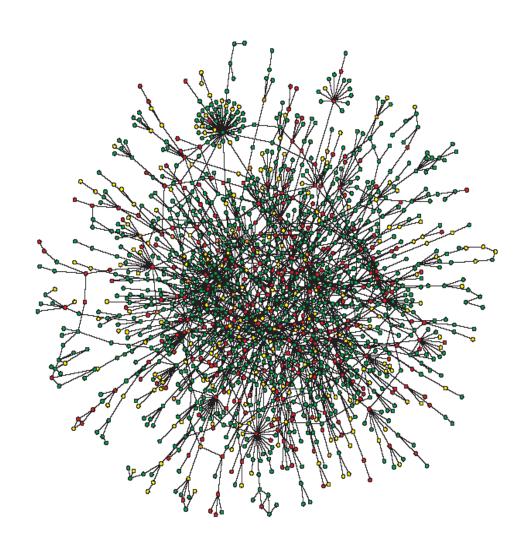
Acyclic graph of an expression (DAG)

$$((a+b)*c+d/(b*(a+b)))*(a+b)*c$$

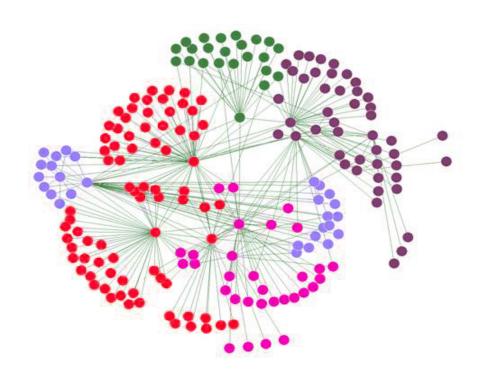
#### Gene regulation network in biology



#### **Protein-protein interaction network (in yeast)**



#### **Social networks**



### Graph representations

$$G = (V, E)$$
  $V = \{1, 2, ..., n\}$ 

#### Adjacency list

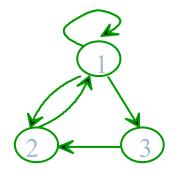
reduces the size if  $|E| << (|V|)^2$  reading time : O(|V| + |E|)

#### Adjacency matrix

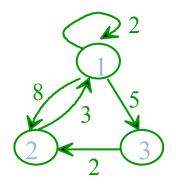
using matrix operations reading time  $O(|V|)^2$ 

Other representations possible

### Adjacency lists

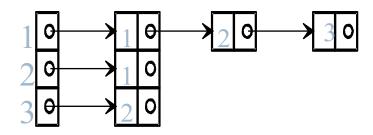


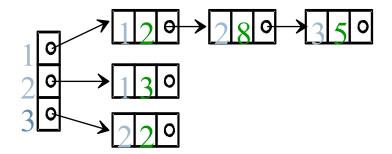
Lists of A(s)



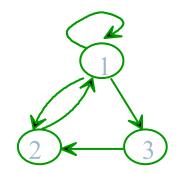
weight:  $w: A \longrightarrow X$ 

$$V = \{ 1, 2, 3 \}$$
  
 $E = \{ (1,1), (1, 2), (1, 3), (2, 1), (3, 2) \}$ 

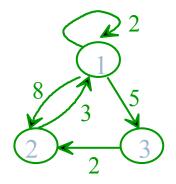




### Adjacency matrix



M[i, j] = 1 iff j is adjacent to i



weight:  $w: A \longrightarrow X$ 

$$V = \{ 1, 2, 3 \}$$
  
 $E = \{ (1,1), (1, 2), (1, 3), (2, 1), (3, 2) \}$ 

$$M = \left(\begin{array}{ccc} 1 & 1 & 1 \\ 1 & 0 & 0 \\ 0 & 1 & 0 \end{array}\right)$$

$$W = \left(\begin{array}{ccc} 2 & 8 & 5 \\ 3 & 0 & 0 \\ 0 & 2 & 0 \end{array}\right)$$

# Graph algorithms

- Exploration
  - Depth-first or breadth-first search
  - Topological sorting
  - Strongly connected components
- Path computation
  - Shortest path
  - Transitive closure
  - Eulerian and Hamiltonian paths

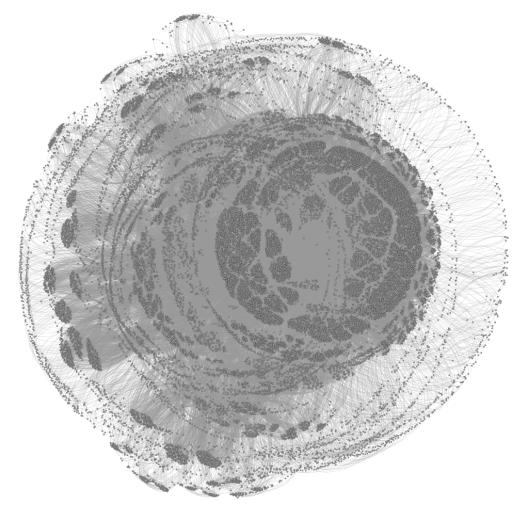
- Minimum spanning trees
  - Kruskal's and Prim's algorithms
- Networks
  - Maximum flow
- Others
  - Graph coloring
  - Planarity testing
  - ...

# Shortest paths in graphs

#### Single-source shortest path: unweighted case

- Path length = number of edges
- Distance between two nodes = length of the shortest path
- ▶ Problem: given a (directed or undirected) graph G = (V, E) and a source node  $s \subseteq V$ , compute the distance from s to each reachable node

#### Single-source shortest bath: unweighted case

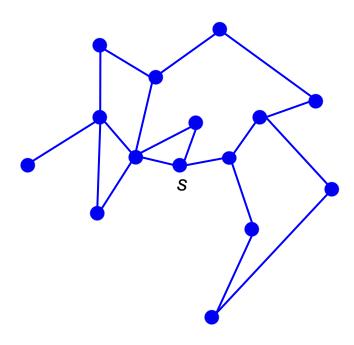


a subgraph (29,160 nodes) of the graph of Rubik's mini cube (2x2x2) configurations (3,674,160 nodes)

https://miscellaneouscoder.wordpress.com/2014/07/28/working-with-rubiks-group-cycle-graphs/

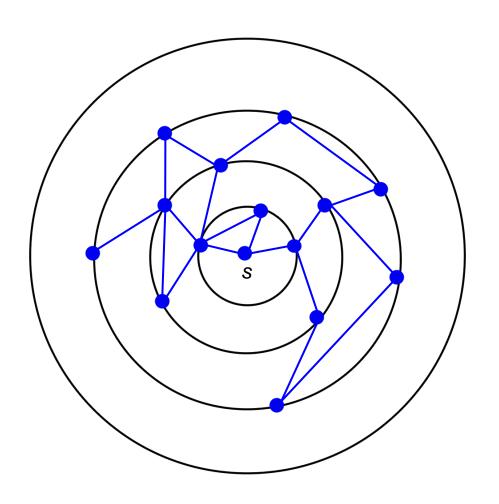
Given a source node s,

Discovers all nodes reachable from s



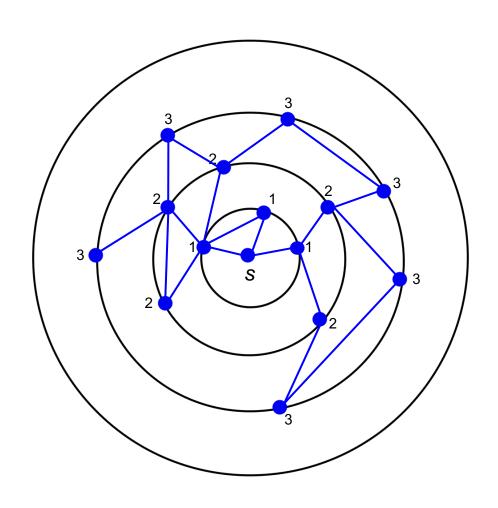
#### Given a source node s,

- Discovers all nodes reachable from s
- Proceeds by "concentric circles"
- Discovers all nodes at distance d from s before discovering any nodes at distance d+1



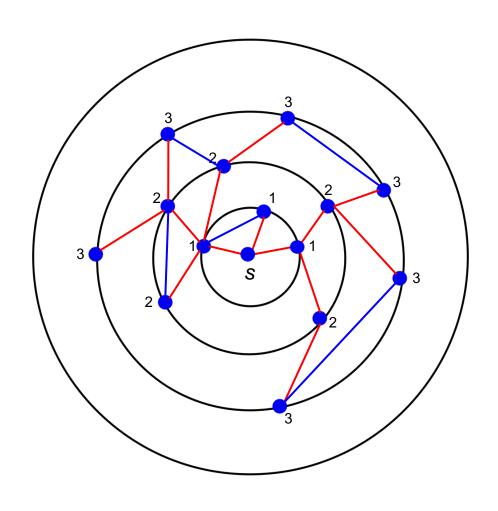
#### Given a source node s,

- Discovers all nodes reachable from s
- Proceeds by "concentric circles"
- Discovers all nodes at distance d from s before discovering any nodes at distance d+1
- Computes the distances from s
- Computes a breadth-first tree encoding one shortest path for each node



#### Given a source node s,

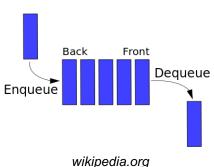
- Discovers all nodes reachable from s
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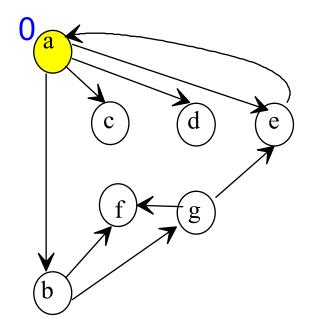


#### How it works?

- colors every node white (not yet discovered), yellow (discovered but may have white adjacent nodes), or red (discovered and all adjacent nodes discovered)
- yellow nodes = "active frontier" (nodes under processing)
- when processing a (yellow) node, determine all white neighbors, set their distance to be larger by 1, color them yellow. After that, color the node red.

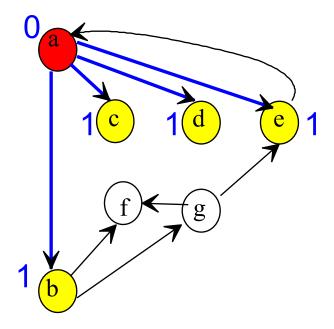
```
procedure BFT (s node of V);
begin
for each node v of V do {
       visited[v] = false ; //s is white
      d[v] = \infty; \pi(v) = nil
visited[s]=true ; //s becomes yellow
d[s]=0;
Queue = enqueue (empty-queue, s);
while not empty (Queue) do {
       u = dequeue (Queue);
       for t = first to last successor of u do
              if not visited [ t ] then
                     visited[ t ]=true ; //t becomes yellow
                     d[t] = d[u] + 1; \pi(t) = u
                     Queue = enqueue (Queue, t);
       //s' becomes red
end
```





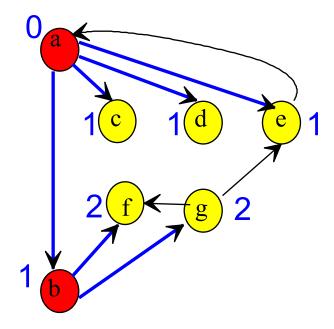
Queue: a

**Order of traversal:** 



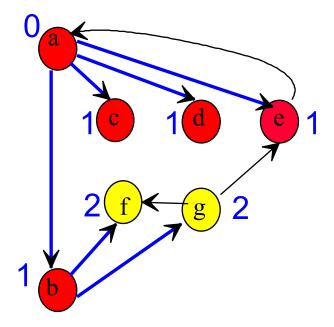
Queue: a b c d e

Order of traversal: a



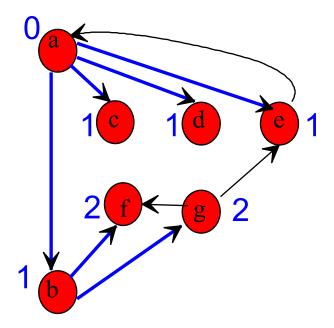
Queue:a b c d efg

Order of traversal: a b



Queue: a b c d e f g

Order of traversal: a b c d e



Queue: a b c d efg

Order of traversal: a b c d e f g

#### Questions

▶ Show that BFS runs in time O(n+m) (assuming the graph is represented by adjacency lists), n=|V|, m=|E|

- ▶ Show that if  $(v_1, v_2, ..., v_r)$  is the state of the Queue, then  $d[v_r] \le d[v_1] + 1$  and  $d[v_i] \le d[v_{i+1}]$  for all i
- Show that upon termination  $d[v]=\delta(s,v)$ , where  $\delta(s,v)$  is the length of the shortest path from s to v

# $d[v]=\delta(s,v)$ : sketch of the proof

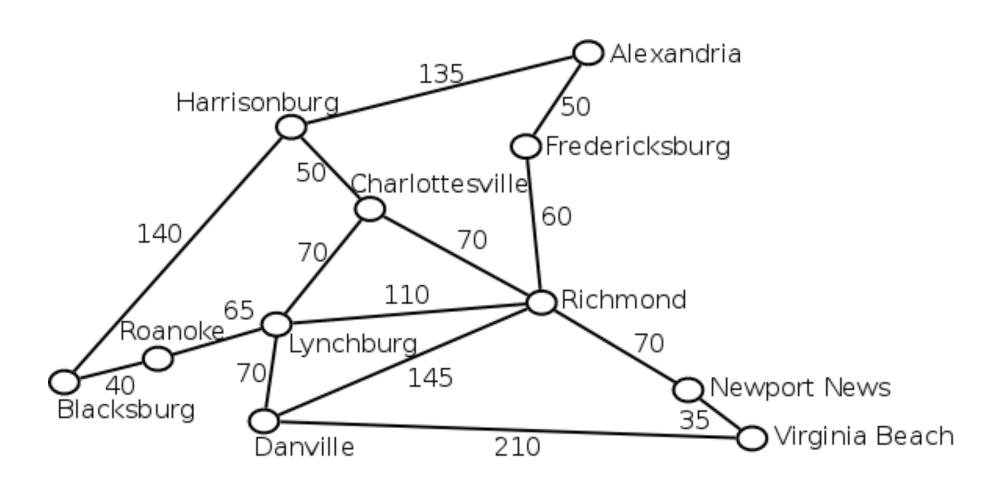
- by contradiction, let v be the closest to s node with  $d[v] > \delta(s,v)$
- consider a shortest path from s to v, and let u be the node preceding v in this path
- $\delta(s,v)=\delta(s,u)+1$  (by properties of shortest paths)
- rightharpoonup consider the moment when u was dequeued  $(d[u] = \delta(s,u))$
- ightharpoonup if v was white then, we have  $d[v] = \delta(s,v) \Rightarrow contradiction$
- if v was yellow then, it was visited earlier by exploring the successors of some w with  $d[w] \le d[u]$ . Then  $d[v] = d[w] + 1 \le d[u] + 1 \Rightarrow contradiction$
- ▶ if v was red, then  $d[v] \le d[u] \Rightarrow$  contradiction

### Space efficient BFS

- **BFS** stores the queue which (in the worst case) can contain O(n) nodes, i.e.  $O(n \log n)$  bits
- Can we implement BFS with o(n log n) bits?
- **Example of a result:** There exists an algorithm that outputs vertices in the BFS order in time O(n+m) and uses 2n+o(n) bits
  - [N. Banerjee, S. Chakraborty, V. Raman, and S. R. Satti. Space efficient linear time algorithms for BFS, DFS and applications. Theory of Computing Systems, Jan 2018]

# Single-source shortest path: weighted case

#### Single-source shortest path: weighted case



# Shortest path problem

Weighted (directed or undirected) graph: G = (V, E, w) where

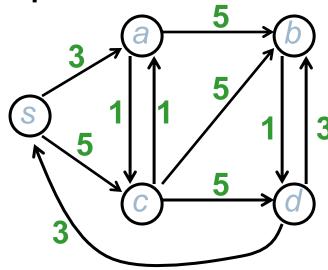
 $w: E \rightarrow \mathbf{R}$  (weight/cost)

Source :  $s \in V$ 

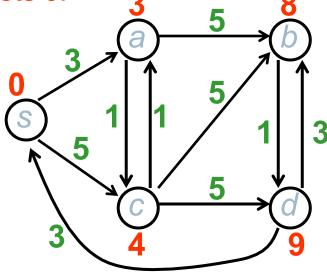
**Problem**: for all  $t \in V$ , compute

 $\delta(s, t) = \min \{\{ w(c) ; c \text{ path from } s \text{ to } t \} \cup \{+\infty\} \}$ 

#### **Example**:



#### Costs $\delta$ :



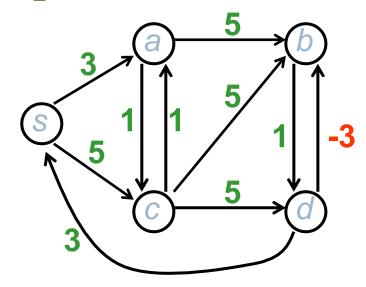
## Properties of the shortest paths

#### **Proposition 1 (existence):**

shortest paths are well-defined (i.e. for all  $t \in V$ ,  $\delta(s, t) > -\infty$ ) **iff** the graph does not have a cycle of cost < 0 reachable from s

**Proposition 2:** if there exists a shortest path from *s* to *t*, then there exists one without a cycle

**Proposition 3:** if there exists a shortest path from s to t, then there exists one with no more than |V|-1 edges



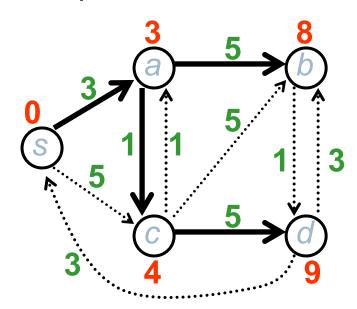
# Main properties

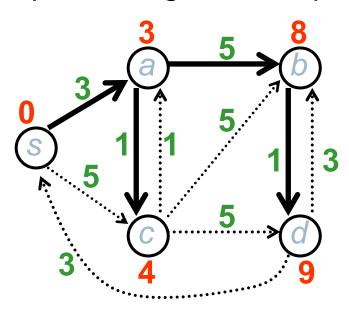
**Property 1**: G = (V, E, w)let c be a shortest path from p to rand q be the node preceding r in c. Then  $\delta(p, r) = \delta(p, q) + w(q, r)$ .



**Property 2**: A subpath of a shortest path is a shortest path

Shortest path tree: tree rooted at s representing shortest paths





# Main properties (cont)

**Property 3**: G = (V, E, w) let c be a path from p to r and q be the node preceding r in c. Then  $\delta(p, r) \leq \delta(p, q) + w(q, r)$ .



### Relaxation

```
Compute \delta(s,t) by successive approximations
t \in V \ d[t] = \text{estimate (from above) of } \delta(s, t)
       \pi[t] = predecessor of t on
              a path from s to t of cost d[t]
Initialization of d and \pi
INIT
       for all t \in V do
       { d[t] = \infty ; \pi[t] = \text{nil} }
       d[s] = 0;
                                                               d(q)
Relaxation of the edge (q, r)
RELAX(q, r)
       if d[q] + w(q, r) < d[r]
```

then  $\{d[r] = d[q] + w(q, r) ; \pi[r] = q\}$ 

# Relaxation (cont)

#### **Proposition**:

the following property is an invariant of **relax**: for all  $t \in V$ ,  $d(t) \ge \delta(s, t)$ 

*Proof*: by induction on the number of executions of relax

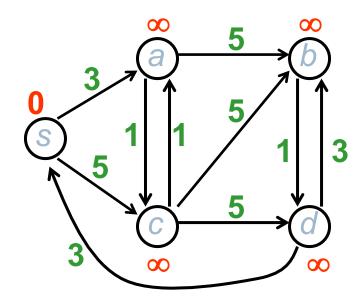
# Dijkstra's algorithm

**Assumption**:  $w(p, q) \ge 0$  for all edges (p, q)

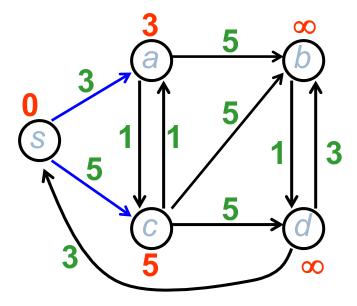
```
begin  \begin{array}{c} \textbf{INIT};\\ S = \varnothing \;;\;\; Q = V \;;\\ \textbf{while}\;\; Q \neq \varnothing \;\;\; \textbf{do}\;\; \{\\ q = \textbf{MIN}_d(Q) \;;\;\; Q = Q \setminus \{q\} \;;\;\; S = S \cup \{q\} \;;\\ \textbf{for all}\;\; r \;\; \textbf{successor}\;\; \textbf{of}\;\; q \;\;\; \textbf{do}\\ \textbf{RELAX}(q,\,r) \;;\\ \} \\ \textbf{end} \end{array}
```

- At each iteration, the algorithm extracts a node from Q that is never returned to Q
- **RELAX**(q, r) may change d[r]

# Example

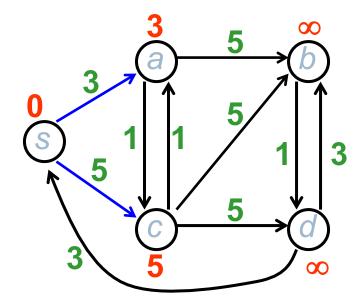


$$S = \{\emptyset\}$$
  
 $Q = \{s, a, b, c, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = \text{nil}$   
 $\pi[b] = \text{nil}$   
 $\pi[c] = \text{nil}$   
 $\pi[d] = \text{nil}$ 

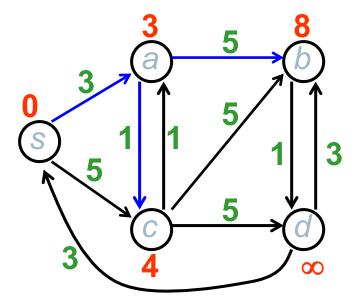


$$S = \{s\}$$
  
 $Q = \{a, b, c, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
 $\pi[b] = \text{nil}$   
 $\pi[c] = s$   
 $\pi[d] = \text{nil}$ 

## Example (cont)

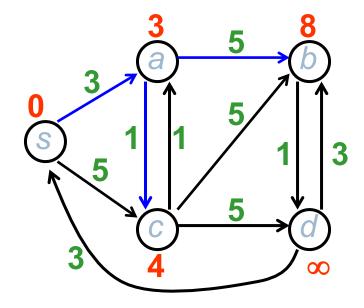


$$S = \{s\}$$
  
 $Q = \{a, b, c, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
 $\pi[b] = \text{nil}$   
 $\pi[c] = s$   
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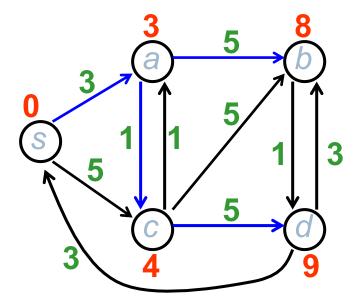


$$S = \{s, a\}$$
  
 $Q = \{b, c, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
 $\pi[b] = a$   
 $\pi[c] = a$   
 $\pi[d] = \text{nil}$ 

## Example (cont)

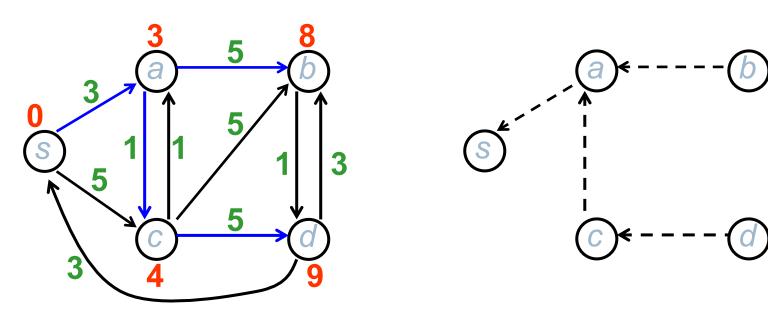


$$S = \{s, a\}$$
  
 $Q = \{b, c, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
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$$S = \{s, a, c\}$$
  
 $Q = \{b, d\}$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
 $\pi[b] = a$   
 $\pi[c] = a$   
 $\pi[d] = c$ 

# Example (cont)

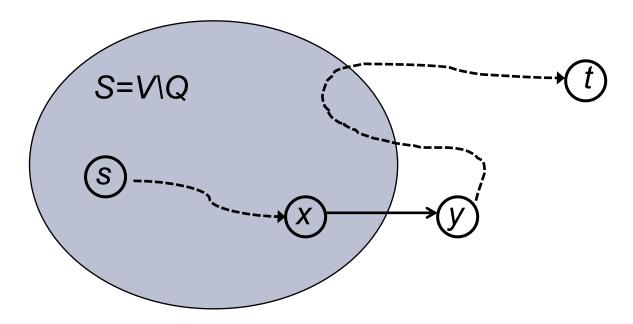


$$S = \{s, a, c\}$$
  
 $Q = \{b, d\}, Q = \{d\} \text{ then } Q = \emptyset$   
 $\pi[s] = \text{nil}$   
 $\pi[a] = s$   
 $\pi[b] = a$   
 $\pi[c] = a$   
 $\pi[d] = c$ 

# Correctness of Dijkstra's algorithm

**Proposition**: After the execution of Dijkstra's algorithm on a graph G = (V, E, w),  $d[t] = \delta(s, t)$  for all  $t \in V$ .

Proof by contradiction: let  $d[t] \neq \delta(s, t)$ 



Properties of Dijkstra's algorithm

▶ Algorithm maintains three sets:

S: finished nodes, for which  $d[t] = \delta(s, t)$  (red)

▶ S': nodes of Q with  $d[t] < \infty$  (yellow)

▶ nodes of Q with  $d[t]=\infty$  (white)

 Algorithm can be seen as expanding a ball centered at s following a greedy strategy

# Implementation

#### With adjacency matrix

time  $O(n^2)$  (where n=|V|)

#### With adjacency lists

depends on the data structure for Q

we need to support operations:

- insert an element to Q
- extract an element with minimum d value
- modify (decrease) the *d* value of an element (when relaxing)
- ⇒ (min-)priority queue

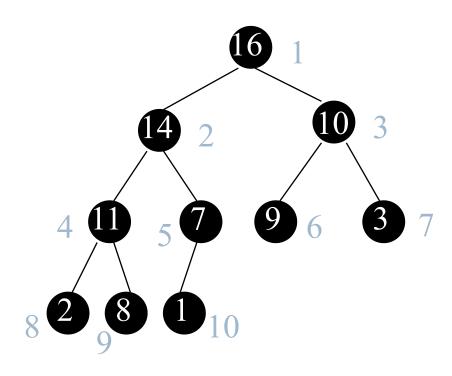
# Priority Queues

- (max-)Priority Queue is a data structure that supports operations
  - ► INSERT(S,x)
  - MAX(S)
  - EXTRACT-MAX(S)
  - INCREASE-KEY(S,x,k): increase the key of x to k
- Priority Queues are used in
  - Dijkstra's algorithm for shortest paths
  - Prim's algorithm for minimum spanning tree
  - other greedy algorithms
- Implemented using heaps

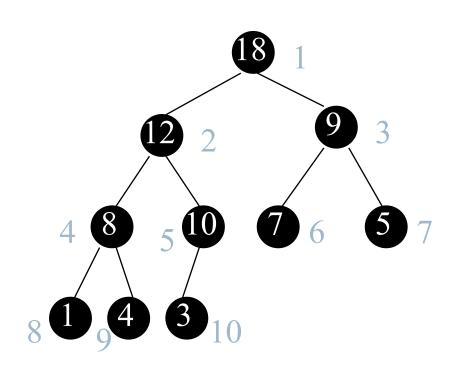
# Binary Heaps

- Binary heap:
  - ▶ a binary tree that is
  - **complete**: every level except possibly the bottom one is completely filled and the leaves in the bottom level are as far left as possible
  - > satisfies the (max-)heap property: the key stored in every node is greater than or equal to the keys stored in its children
    - If the key at each node is smaller than or equal to the keys of its children, then we have a min-heap

# Binary (max-)heap: example



# Binary heaps stored in arrays



Due to their regular structure, binary heaps are easily stored in arrays

Given index i of a node,

- the index of its parent is [i / 2]
- the indices of its children are 2i and 2i+1



# Binary heaps: some properties

▶ The height of a heap is [log(n)]

Not every array represents a heap

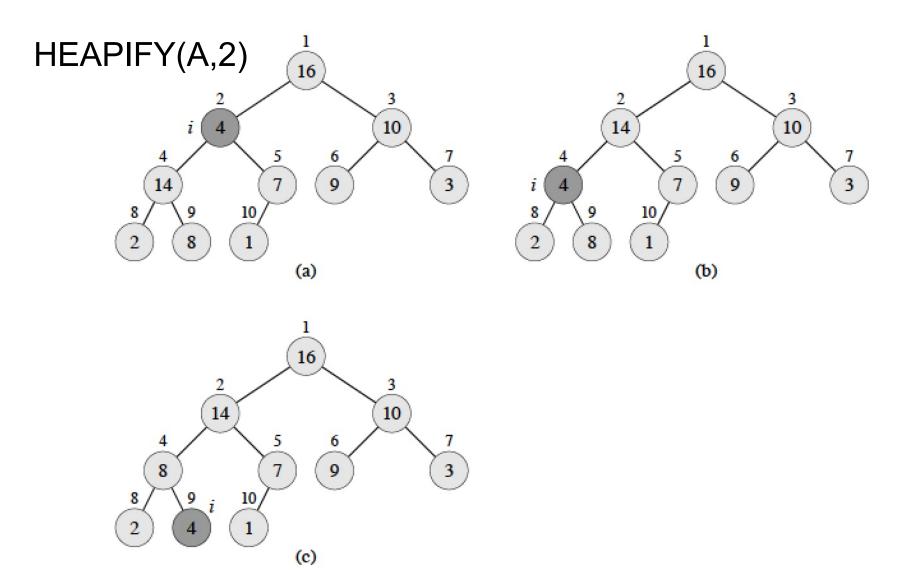
In a max-heap, the largest element is at the root and the smallest element is in a leaf

# Heapify

Assume that node i violates the heap property, but the children nodes 2i and 2i+1 (if exist) are heaps.

```
HEAPIFY(A,i)
  if A[2i]>A[i] or A[2i+1]>A[i] then
      if A[2i+1]>A[2i] then
             exchange A[i] and A[2i+1];
             HEAPIFY(A, 2i+1)
      else
             exchange A[i] and A[2i];
             HEAPIFY(A, 2i)
      end
  end
```

# Heapify: example



# Building a binary heap

• Given an array A[1..n], build a binary heap for array elements

```
BUILD-HEAP(A,n)

for i=\lfloor n/2 \rfloor downto 1 do HEAPIFY(A,i);
```

**Exercise:** build the heap for A=[4,1,3,2,16,9,10,14,8,7]

# BUILD-HEAP: complexity

Straightforward estimation O(n · log(n))

#### Refined analysis:

- Cost of a call to HEAPIFY at a node depends on the height, h, of the node O(h).
- Height of most nodes smaller [log(n)]
- $\blacktriangleright$  Height of nodes h ranges from 0 to  $\lfloor \log(n) \rfloor$
- number of nodes of height h is at most  $[n/2^{h+1}]$ ?

# Heap Characteristics

- $\blacktriangleright$  Height =  $\lfloor \log n \rfloor$
- Number of leaves = [n/2]
- Number of nodes of height  $h \leq \lceil n/2^{h+1} \rceil$

#### Proof by induction:

- remove all leaves from the heap
- there remains  $n \lfloor n/2 \rfloor = \lfloor n/2 \rfloor$  nodes
- ▶ the height of each node is decremented by 1
- ▶ nb of nodes of height h-1 is (by induction)  $\lceil \lfloor n/2 \rfloor / 2^h \rceil \le \lceil n/2^{h+1} \rceil$

# Tighter bound for BUILD-HEAP: O(n)

time of BUILD-HEAP is 
$$\sum_{h=0}^{\lfloor \log n \rfloor} \left[ \frac{n}{2^{h+1}} \right] O(h) = O\left(n \sum_{h=0}^{\lfloor \log n \rfloor} \frac{h}{2^h}\right)$$

note that 
$$\sum_{h=0}^{\lfloor \log n \rfloor} \frac{h}{2^h} \le \sum_{h=0}^{\infty} \frac{h}{2^h} = \frac{1/2}{(1-1/2)^2} = 2$$

therefore the time is O(n)

# Priority Queue

- ▶ MAX(A): return the heap root
- ► EXTRACT-MAX(A):

# Priority Queue

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- ► INCREASE-KEY(A,i,k):

# Priority Queue

- ▶ MAX(A): return the heap root
- EXTRACT-MAX(A): exchange A[1] and A[n], discard element n, and apply HEAPIFY(A,1)
- ► INCREASE-KEY(A,i,k):  $A[i] \leftarrow k;$ while A[[i/2]] < A[i] do

exchange A[[i/2]] and A[i];

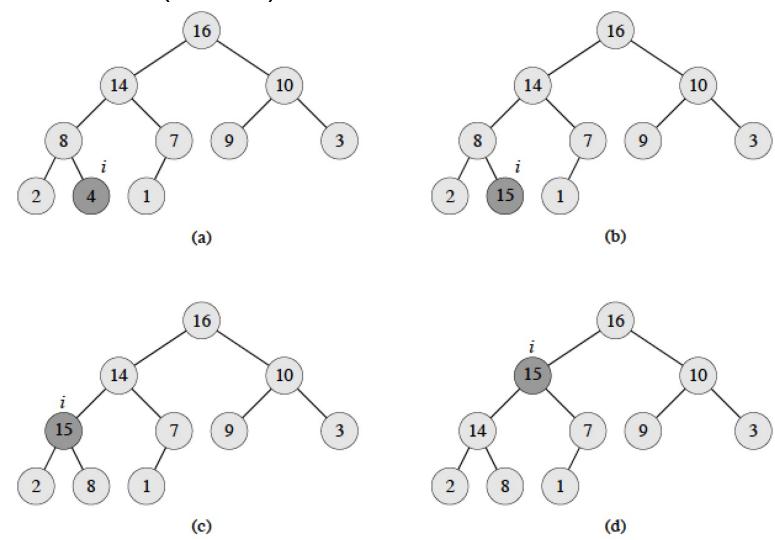
 $i\leftarrow parent(i)$ 

#### end

► INSERT(A,i): insert a new leaf n+1 with key  $-\infty$ ; call INCREASE-KEY(A,n+1,k)

# INCREASE-KEY: example

INCREASE-KEY(A,9,15)



## Priority Queues: time bounds

- ▶ MAX: *O*(1)
- ► EXTRACT-MAX, INCREASE-KEY, INSERT: O(log(n))

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- ▶ MAX: *O*(1)
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#### Various improvements have been proposed

- ▶ Fibonacci heaps take O(1) amortized time for INSERT and INCREASE-KEY
- ▶ if keys are integers bounded by C, van Emde Boas trees support INSERT, DELETE, MAX, MIN, SUCC, PRED in time O(log log(C))

# Back to Dijkstra's algorithm

```
With adjacency matrix time O(n^2)
```

#### With adjacency lists

```
Q: priority queue if implemented by binary heaps: n building a heap of n elements: O(n) n operations \min_{d} : O(n \cdot \log n) m operations \min_{d} : O(m \cdot \log n) total time O((n+m) \cdot \log n): improves over O(n^2) if m=o(n^2/\log n)
```

time can be improved to  $O(n \cdot \log n + m)$  using Fibonacci heaps, as decreasing the key takes O(1) amortized