# Advanced Programming Techniques

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### About myself...

- Associate Professor (*Conferentiar*) at the Univ. of Bucharest, Faculty of Mathematics and Computer Science.
- Research Scientist at the National Institute of Research and Development in Informatics (ICI).
- PhD in Nice-Sophia Antipolis, France.
- Various visits/internships in the US, Chile, Germany,...

Research in Graph Algorithms and Data Structures

#### Come to see me if...

• You want to learn more about (algorithmic aspects of) the class!

You need help for your Bachelor/Master thesis

 Plenty of my problems need good students to find good solutions, and/or implement some proposed solutions.

You want to discuss about internship/PhD/job opportunities...

#### Practical Information

- All materials (slides + documents) to be put on Moodle.
- My website with contact information

https://sites.google.com/view/guillaume-ducoffes-homepage/home

Additional documents can be put on it - upon request

- Attendance is NOT mandatory. But...
  - Bonus for seminars works (max. 1p)

## Your **grades**

- One practical test at the end of the semester.
  - Programming languages: either C++ or Java.
  - 4 to 5 sets of subjects proposed (students have to pick only one).

Final grade:  $min\{10, Practical Test + Bonus\}$ 

There shall be NO "punct din oficiu".

#### About the class

#### **Growing size** of networks



**Social networks** (Facebook  $\geq 1.79$  billion users)

**Data Centers** (Microsoft  $\geq 1$  million servers)

**the Internet** (≥ 55811 Autonomous Systems)

"Efficient" algorithms on these graphs?

polynomial → quasi-linear time

quadratic → (sub)linear space

sequential → parallel+distributed

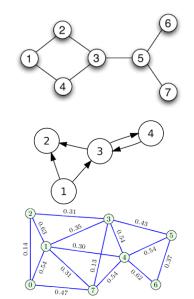
Focus on key procedures in graph & relational databases, and beyond.

#### Tentative schedule for this semester

- Graph basics (Today).
- Parallel search
- Parallel connected components & spanning forests
- Centrality indices
- Pattern detection and couting
- Labelling schemes (Hub labels, etc.)
- Spanners & emulators
- Graph sketches
- Map reduce Graph algorithms
- Acyclic Database schemes

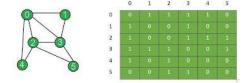
### Graphs

- (Undirected) Graph: G = (V, E)
- $V = {\sf vertices} = {\sf networks} \; {\sf units}$
- E = edges
- Directed Graph:  $\overrightarrow{G} = (V, A)$
- A = arcs
- Weighted Graph: G = (V, E, w)
- $w: E \to \mathbb{R}$  is a weight function (length, cost)

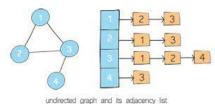


### Graph representations

Adjacency matrix



#### Adjacency list



### Basic operations on graphs

- Enumerate all edges
- Enumerate all neighbours of a vertex
- Output the degree of a vertex
- Decide whether two vertices are adjacent
- Addition/Removal of edges/vertices.
- **Ontraction** of an edge e = uv: replace u, v by some new vertex x whose neighbourhood equals  $N(u) \cup N(v) \setminus \{u, v\}$ .

**O** Enumerate all edges: scan the whole structure in O(n+m) time.

Better: Keep the lists of isolated vertices at the end, to avoid scanning them.  $\rightarrow \mathcal{O}(m)$  time.

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- lacktriangle Addition of a new vertex: insertion of a new empty list at the end in amortized  $\mathcal{O}(1)$  time (using a doubling array to store adjacency list).

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- lacktriangle Addition of a new vertex: insertion of a new empty list at the end in amortized  $\mathcal{O}(1)$  time (using a doubling array to store adjacency list).
- Removal of an <u>isolated</u> vertex v: Switch v with the last vertex, then remove the last vertex, in amortized  $\mathcal{O}(1)$  time (using a doubling array to store adjacency list).

#### Reminder: Hash Table

Store a collection of pairs (key, value).

- Three operations:
  - int lookup(e); Returns the value associated to some key e (if it is present in the table)
  - **void** insert(e, v); Adds a new pair (e,v) (if the key e is not already present in the table)
  - void delete(e); Deletes a pair (key, value) given its key e.

Each operation runs in expected  $\mathcal{O}(1)$  time.

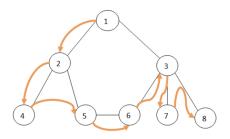
## Adjacency Testing

We store every edge in a Hash table. To every edge uv, we associate pointers its positions in the respective adjacency lists of u and v.

- lacktriangle Adjacency testing: Lookup in the table in expected  $\mathcal{O}(1)$  time.
- Addition of an edge uv: Insertion of u (v, resp.) at the bottom of the adjacency list of v (u, resp.). Insertion in expected  $\mathcal{O}(1)$  time in the Hash table, along with pointers to the bottom positions in the respective adjacency lists of u and v.
- lacktriangle Removal of an edge uv: Lookup in the Hash-table in order to find the positions of u, v in the respective adjacency lists of v, u. Then, removal in both adjacency lists and in the table.
- Removal of a vertex: removal of every incident edge + removal of an isolated vertex.

#### Serial DFS

Pick the **most recently visited** vertex with at least one neighbour unvisited. Then, go to an arbitrary unvisited neighbour of this vertex.



Equivalently: either continue the search to any neighbour of the current vertex (if possible) or backtrack to the father node in the search tree.

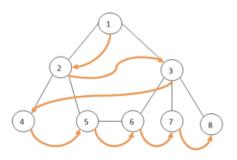
### Implementation

At any moment during the execution of the algorithm, we **keep in a stack** the path from the start vertex  $v_{n-1}$  to the current vertex  $v_i$ .

```
S := \{\}
S.push(v_{n-1})
Visit v_{n-1}
while !S.empty():
     u := S.top() //current vertex
     if there exists some v \in N(u) unvisited:
        S.push(v)
        Visit v
     else: S.pop()
Complexity: \mathcal{O}(n+m)
```

#### Serial BFS

Pick the **least recently visited** vertex with at least one neighbour unvisited. Then, go to an arbitrary unvisited neighbour of this vertex.

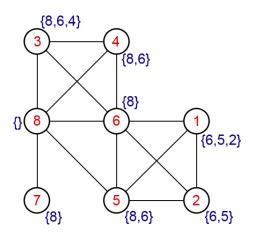


#### **Implementation**

We keep in a queue the next vertices to be visited, in order.  $Q := \{\}$ Q.enqueue( $v_{n-1}$ )  $visited[v_{n-1}] := True$ while !Q.empty(): u := Q.dequeue() //current vertex Visit u for all  $v \in N(u)$ : **if** !visited[v]: Q.enqueue(v)visited[v] := TrueComplexity:  $\mathcal{O}(n+m)$ 

#### **LexBFS**

The visited neighbours of each vertex are ordered by decreasing label. At every step, the next vertex to be visited must have a label which is **lexicographically** maximum.



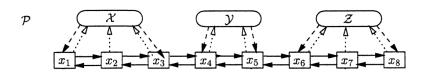
#### Partition Refinement

- Data Structure that maintains an ordered collection of pairwise disjoint sets, subject to the following basic operations:
  - init(V): initialize the structure with one set, equal to V.
  - refine(S): for each set X such that  $X \cap S \neq \emptyset$  and  $X \setminus S \neq \emptyset$ , we replace X by the two consecutive new sets  $X \cap S$  and  $X \setminus S$ .

• Operation init(V) is in worst-case  $\mathcal{O}(|V|)$ . Each operation refine(S) is in worst-case  $\mathcal{O}(|S|)$ . Note that these are optimal runtimes!

### Implementation

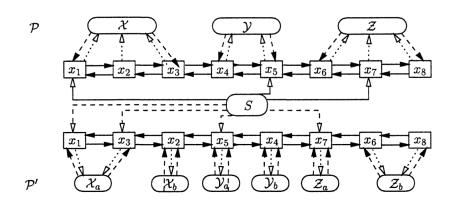
- Elements in V are maintained in a doubly-linked list  $\mathcal{L}$ , such that all elements in a same set X are consecutive.
- Each set X of the partition is represented by a structure with two fields: pointers to its first and last elements in  $\mathcal{L}$ .
- ullet Each node in the list  ${\mathcal L}$  stores a pointer to the set X which contains it.



#### Refinement

- To each set X, "lazily" associate an empty list L[X] (we effectively create the list only if it needs to be accessed at some point during the execution of the algorithm).
- For each  $s \in S$ , access to its set X and add a pointer to s in L[X]. Put a pointer to X in an auxiliary list  $\mathcal{H}$  (the sets of  $\mathcal{H}$  are those intersecting S).
- For each set X in  $\mathcal{H}$ , if  $L[X] \neq X$ , then:
  - Update the first and last element of X as its first and last elements not in S (forward/backward search in  $\mathcal{L}$ ).
  - Remove all elements in L[X] from  $\mathcal{L}$ ;
  - Reinsert L[X] immediately before the first element of X (or immediately after the last element of X);
  - Create a new set from L[X].

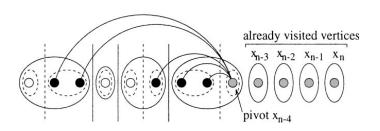
#### Refinement: illustration



## Application to LexBFS

Traverse backward the list of all vertices – with the start vertex  $v_{n-1}$  at the end.

Repeatedly visit the next vertex  $v_i$  and refine the unvisited vertices according to  $N(v_i)$ .



Complexity:  $\mathcal{O}(n+m)$ 

## Weighted graphs

If all edge-weights are positive, we can use Dijkstra's algorithm as a replacement for BFS:

```
H := empty heap
```

Insert every  $v \in V$  in H with infinite value.

$$H[v_{n-1}] := 0 //start vertex$$

while H is nonempty:

$$(v_i, d_i) := H.extract_min()$$

for all  $u \in N(v_i)$  such that u is in the heap:

if 
$$d_i + w(v_i, u) < H[u]$$
: H.decrease\_key(u,  $d_i + w(v_i, u)$ )

We need to perform on the heap: n insertions, n deletions, and  $\mathcal{O}(m)$  decrease key operations.

 $\rightarrow \mathcal{O}(n \log n + m)$  time by using Fibonacci heaps.

## Limitations of Dijkstra's algorithm

- Only applies to **positive** edge-weights (null-weights can be easily handled, but negative weights are a real issue).
- In some applications, we only need to compute a shortest path between a source vertex  $v_{n-1}$  and any vertex of some subset  $P \subseteq V$ . By contrast, Dijkstra's algorithm computes all distances from  $v_{n-1}$ .
- In real-life applications, the graph may be so large that we cannot even traverse it in full. It may even be infinite!
- $\implies$  An alternative search, proposed at the infant stages of AI: the  $A^*$  heuristic (Hart, Nilsson & Raphael, 1986).

#### A\* inputs

- A directed + arc-weighted graph G = (V, A, w).
- $\rightarrow$  Weights may be negative. However, we assume that there is no negative circuit (Assumption A1).
- A start vertex s + a destination subset  $P \subseteq V$
- $\rightarrow$  Since G may be infinite, we need to make the minimal assumption that there exists a path from s to P (Assumption A2)
- A heuristic function  $h: V \to \mathbb{R}$ , which must represent a lower bound estimate on the distance from a vertex to P, i.e.,  $d(x, P) \ge h(x)$ .
- $\rightarrow$  e.g., distances "as the crow flies" for road/street networks. If all weights are nonnegative, then we may simply set h(x) = 0.

### A\* output

Compute d(s, P), and a shortest path from s to a closest vertex in P.

• The  $A^*$  heuristic will stop as soon as it reaches a vertex in P.

• For this halting condition to make sense, we need to make the minimal assumption that  $d(p, p') \ge 0$  for every two vertices  $p, p' \in P$  (Assumption A3).

#### Initialization

The presentation mostly follows that proposed by Habib and Simonet in their research report (1991).

- We maintain two subsets: Closed (already visited vertices), and Open (vertices to be visited)).
- $\rightarrow$  Initially, Closed =  $\emptyset$  and Open =  $\{s\}$ .

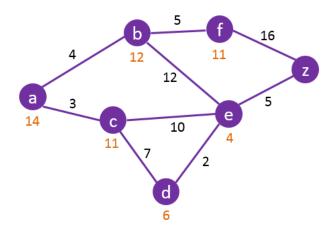
As we shall see, we may reopen closed vertices if a better path from s has been found. This is different from Dijkstra's algorithm, where a visited vertex will never need to be visited again.

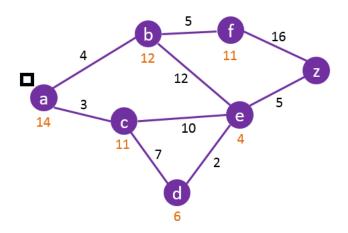
- We maintain an in-arborescence ("shortest-path tree") with root s. The predecessor of each vertex  $x \neq s$  is stored in a variable  $\operatorname{pred}(x)$ . The best-known distance from s to x is stored in variable g(x).
- $\rightarrow$  We need g(x) to be defined only if x is in Open. Initially, g(s) = 0.

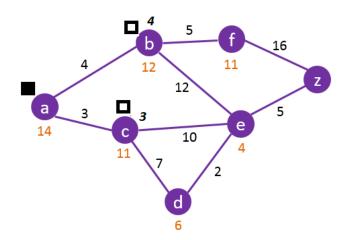
### Main loop

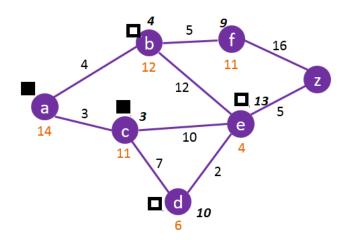
The following procedure runs while Open is nonempty:

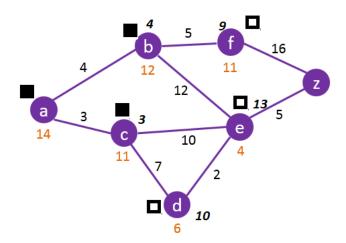
- 1) We compute a vertex  $x \in \text{Open}$  such that f(x) := g(x) + h(x) is minimized (estimated distance from s to P going by x).
- 2) We remove x from Open, then we insert x in Closed.
- 3) If  $x \in P$ , then we stop. Else, we do as follows for every  $y \in N^+(x)$  (out-neighbours):
  - If  $y \in \text{Open and } g(y) > g(x) + w(x, y)$ , then: g(y) := g(x) + w(x, y), pred(y) := x.
  - If  $y \in \text{Closed}$  and g(y) > g(x) + w(x, y), then: g(y) := g(x) + w(x, y), pred(y) := x. Furthermore, we remove y from Closed, then we insert y in Open.
  - If  $y \notin \text{Open} \cup \text{Closed}$ , then: g(y) := g(x) + w(x, y), pred(y) := x. Furthermore, we insert y in Open.

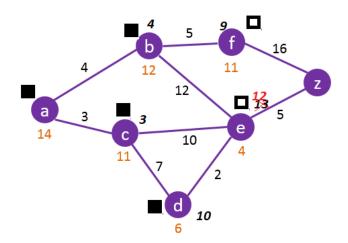


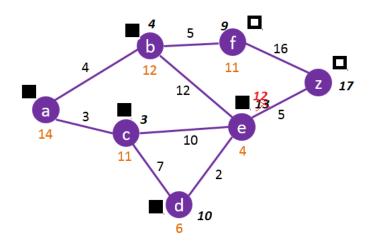


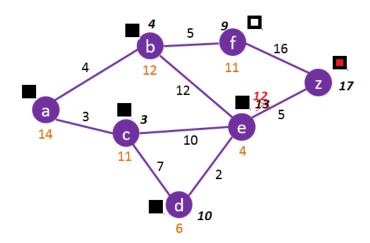








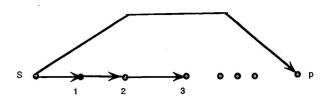




#### No termination

Set w(s, p) = 2, w(s, 1) = 1, w(i, i + 1) = 1/i(i + 1) for every positive integer i.

Set h(i) = 1/i for every positive integer i.



After *i* steps: g(j) = 1 - 1/j for every  $1 \le j \le i$ .

 $\rightarrow s, 1, 2, \dots, i-1$  are closed, i is open and f(i) = 1.

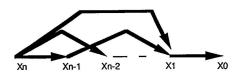
Consequence: we stay blocked on an infinite path!

## The case of finite graphs

**Proposition**: If G is finite, then  $A^*$  always halts.

<u>Proof</u>: every time a vertex x is reopen, we found a better path from s to x. There are only a finite (but exponential in n) number of sx-paths.

Unfortunately, the runtime may be exponential (Martelli):



• In the special case where all weights are nonnegative, and there is no estimate (i.e., h(x) = 0), then we retrieve Dijkstra's algorithm. However, the larger h, the less vertices (without counting repetitions) are visited.

#### The B heuristic

• First proposed by Martelli as a variation of  $A^*$  for graphs with nonnegative arc weights.

<u>Intuition</u>: Recall that f(x) = g(x) + h(x) is a lower bound estimate on the shortest path from s to P that goes by x. Let F be the maximum value f(x) amongst all the *closed* vertices.

- If some open vertices x' satisfy f(x') < F, then amongst those we pick one minimizing g(x') (i.e., we ignore h, or equivalently we perform Dijkstra's algorithm).
- Otherwise, we pick any open vertex x' that minimizes f(x') (the same as for  $A^*$ ).

**Proposition**: the number of iterations is at most  $\mathcal{O}(n^2)$ . Furthermore, it is never worse than for  $A^*$ .

## Questions

