

# **DSA3 seminar assignment**

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# 1 Introduction

## 2 BFS and DFS

### 2.1 Purpose and theoretical background

Both BFS (breadth-first search) and DFS (depth-first search) are tree-search algorithms. They are both used to determine whether a graph is connected.

#### Theorem

A graph is connected if and only if, you can not split it into two such that there are no edges between them.

An approach to check whether a graph is connected is by checking all the partitions (the problem is that there are  $2^{n-1}$  of them so it is very inefficient). Another way to check whether a graph is connected is by checking all pairs of vertices - if there is a path connecting them (problem here again is that there are  $n^2$  pairs of vertices and for each of them we have to check if there is a path which is also very inefficient on top of number of partitions that we would need to check).

#### Definition

A graph is connected if and only if it has a spanning tree.

Checking whether a graph is connected in such a way (by determining whether it has a spanning tree) is far and away the most efficient way to check graph connectedness.

Both DFS and BFS work in a similar way (they both produce spanning trees), but with differences that impact the priority of edges selected. Depending on the way we used to choose edges we get different trees.

A **tree-search** algorithm on  $G$ :

- > Start with a single root vertex  $r \in V(G)$ . This is our initial tree (with just one vertex).
- > Repeat (for as long as possible):
  - Do we have a spanning tree?
  - Is the tree edge cut empty?
  - If not, add one edge from the cut to the tree.

If we want to check *connectedness* of  $G$ , that is the whole algorithm. As noted above, depending on the way we use to choose edges, different spanning tree will result (if the graph is connected of course).

Define: edge cut, rooted tree (also called r-tree), levels (distance from root to a specific vertex), ancestor, descendant, parent or predecessor and children.

$$v \in V$$

Predecessor function:  $p(v)$ , for all  $\{r\}$

## 2.2 BFS

### 2.2.1 BFS introduction

In **breadth-first search** the adjacency lists of vertices are considered on a first-come-first serve basis. **BFS** is implemented with a *queue* data structure.

A *queue* data structure operates in a FIFO (first-in, first-out) principle. Meaning first element that entered the *queue*, will be the first one to leave the *queue* (elements are removed in the same order in which they were inserted).

In **BFS** algorithm we will first consider all of the neighbors (one-by-one) before we look through the neighbours of any of them.

Therefore, first edges incident to  $r$  are selected, and only after we have used (added them to the tree) we are looking at the neighbors of the neighbors of  $r$ .

This is called **breadth-first search** algorithm. This approach expands (spreads) the tree as much as possible.

We start with just the root  $r$  in the *queue* and we repeatedly pop the head of the *queue*, and push all its new neighbors to the *queue*.

For a connected graph, the algorithm will return:

- A spanning tree (**BFS** spanning tree) given by its predecessor function,
- the level function,
- the time function (order in which vertices are added to the tree)

### 2.2.2 BFS algorithm

> INPUT: a connected graph  $G$ , a vertex  $r \in V(G)$

> OUTPUT: an  $r$ -tree  $T \subseteq G$ , its predecessor function  $p$ , its level function  $l$ , the time function  $t$

#### BFS algorithm

$Q := \emptyset, Q \leftarrow r, l(r) := 0, t(r) := 1, \text{ mark } r, i := 1$

**while**  $Q \neq \emptyset$

    consider the head  $x$  of  $Q$

**if**  $x$  has unmarked neighbor  $y$  **then**

$i++$

$Q \leftarrow y, \text{ mark } y, p(y) := x, l(y) := l(x) + 1, t(y) := i$

**else**

        remove head of  $Q$

**end if**

**end while**

**return everything**

### 2.2.3 BFS properties

**BFS**-trees have two basic properties, the first of which justifies our referring to  $l$  as a level function.

**Theorem** Let  $T$  be a BFS tree of  $G$ , with root  $r$ .

a.)  $l(v) = d_T(r, v),$  , for every  $v \in V$ ,

b.)  $|l(u) - l(v)| \leq 1$ , for every  $uv \in E(G)$ .

Level of  $v$  is exactly the distance from root  $r$  to  $v$ .

Every edge of the graph connects only vertices of the same level of the tree or difference by most 1.

**Theorem** Let  $T$  be a BFS tree of  $G$ , with root  $r$ . Then

$l(v) = d_G(r, v)$ , for every  $v \in V$

## 2.3 DFS

### 2.3.1 DFS introduction

In contrast to BFS, where we first scan the whole adjacency list of the vertex on top of the *queue*, in **depth-first search** we scan the adjacency list of the most recent vertex  $x$  added to the *stack* and we look for its neighbour not in  $T$ .

If there is such a neighbor, we add it to  $T$ . If not, we backtrack to the vertex which was added to  $T$  just before  $x$  and examine its neighbours, and so on.

For DFS to be implemented, we use a **stack** data structure. A *stack* is a linear data structure (like a simple list) which has a top element and basic operations such as placing a new item on top of the *stack*, or removing the top element from the *stack*. In contrast to *queue*, a *stack* operates in LIFO (last-in, first-out) principle, meaning elements are inserted and removed exclusively at the designated end of the structure, referred to as the top.

In **depth-first search**, the *stack*  $S$  is initially empty. We pick a root vertex  $r$  and scan its neighbours in its adjacency list. We pick one element from that list and place it on top of the *stack*  $S$ . We then look at this new vertex, now acting as the new top element of *stack*  $S$  and we inspect its adjacency list. If in that list exists a vertex  $y$  which is not already in our tree  $T$  we select it and add it to the top of *stack*  $S$  again. And so on, we continue until there are no suitable vertices in the top element of *stack*  $S$ . If there indeed aren't any, we remove the top element of *stack*  $S$  and check the vertex that is the new top of the *stack*  $S$ .

Again for a connected graph, the algorithm will return:

- A spanning tree (**DFS spanning tree**) given by its predecessor function,
- the level function,
- the time function (order in which vertices are added to the tree)

### 2.3.2 DFS algorithm

Completely the same as BFS, except that we use a **stack** instead of a queue.

> INPUT: a connected graph  $G$ , a vertex  $r \in V(G)$

> OUTPUT: an  $r$ -tree  $T \subseteq G$ , its predecessor function  $p$ , its level function  $l$ , the time function  $t$

#### DFS algorithm

$S := \emptyset, S \leftarrow r, l(r) := 0, t(r) := 1, \text{ mark } r, i := 1$

**while**  $S \neq \emptyset$

    consider the top vertex  $x$  of  $S$

**if**  $x$  has unmarked neighbor  $y$  **then**

$i++$

        move  $y$  to the top of  $S$ , mark  $y$ ,  $p(y) := x$ ,  $l(y) := l(x) + 1$ ,  $t(y) := i$

**else**

        remove  $x$  from  $S$

**end if**

**end while**

**return everything**

### 2.3.3 DFS properties

**Theorem** Let  $T$  be a DFS tree of  $G$ . Every edge of  $G$  joins vertices related in  $T$ .

Like we had for **BFS** a theorem that says every edge of the graph skips at most one level, which is the most important property of BFS. Here we have this very important property of **DFS** trees. Every edge goes vertically, from ancestor to predecessor.

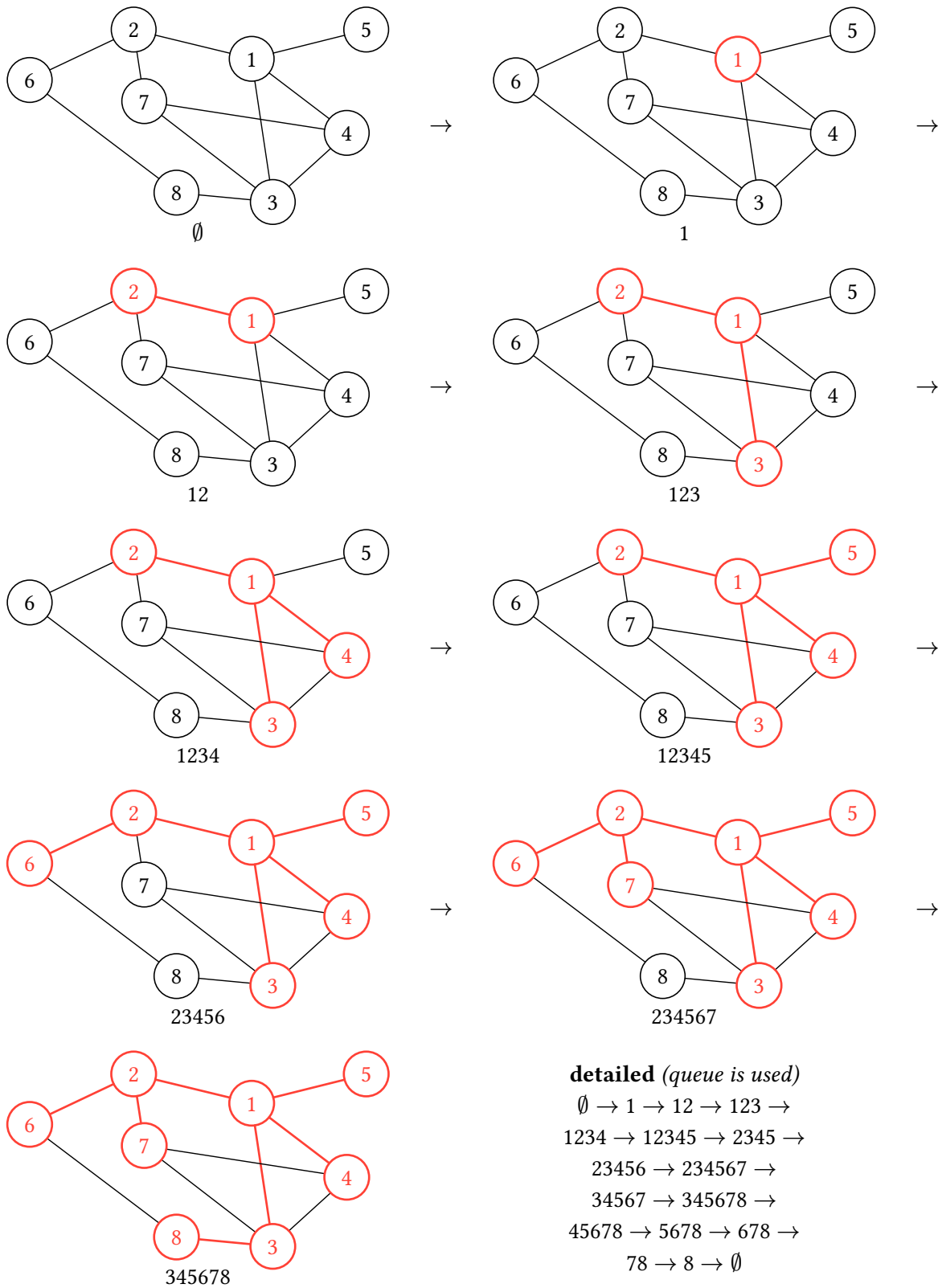
The following proposition provides a link between the input graph  $G$ , its **DFS**-tree  $T$ , and the two time functions  $t$  and  $l$  returned by **DFS**.

**Proposition** Let  $u$  and  $v$  be two vertices of  $G$ , with  $f(u) < f(v)$ .

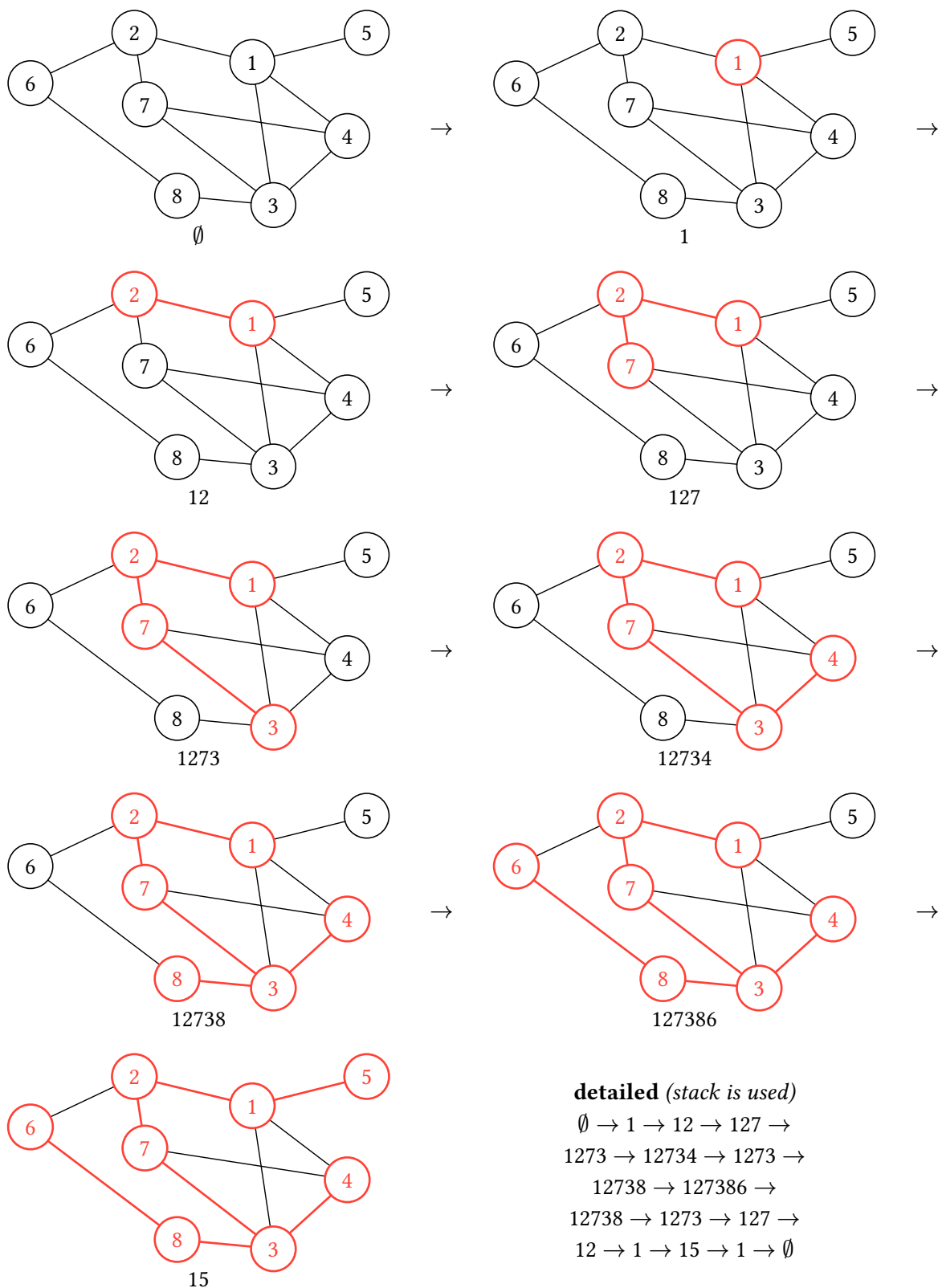
- a) If  $u$  and  $v$  are adjacent in  $G$ , then  $l(v) < l(u)$ .
- b)  $u$  is an ancestor of  $v$  in  $T$  if and only if  $l(v) < l(u)$ .

## 2.4 Algorithm examples

### 2.4.1 BFS example



### 2.4.2 DFS example





### **3 Applications**

## **4 Conculsion**