## COMPUTER SCIENCE 311 SPRING 2019

# PROGRAMMING ASSIGNMENT 2 TRACING THE SPREAD OF A COMPUTER VIRUS DUE: 11:59 P.M., APRIL 21

#### 1. Overview

Suppose you are monitoring a collection of n networked computers, labeled  $C_1, C_2, \ldots, C_n$ , in order to track the spread of an online virus. You are given a collection of trace data indicating the times at which pairs of computers communicated. The data is provided as a sequence of m ordered triples  $(C_i, C_j, t_k)$ . Such a triple indicates that  $C_i$  and  $C_j$  communicated at time  $t_k$ . For simplicity, let us assume that each pair of computers communicates at most once during the interval under consideration. Your goal is to devise an efficient data structure to answer *infection queries*; that is, queries of the following form:

If the virus was inserted into computer  $C_a$  at time x, could it possibly have infected computer  $C_b$  by time y?

The mechanics of infection are simple: if an infected computer  $C_i$  communicates with an uninfected computer  $C_j$  at time  $t_k$  (in other words, if one of the triples  $(C_i, C_j, t_k)$  or  $(C_j, C_i, t_k)$  appears in the trace data), then computer  $C_i$  becomes infected as well, starting at time  $t_k$ . Infection can thus spread from one machine to another across a sequence of communications, provided that no step in this sequence involves a move backward in time. Thus, for example, if  $C_i$  is infected by time  $t_k$ , and the trace data contains triples  $(C_i, C_j, t_k)$  and  $(C_j, C_q, t_r)$ , where  $t_k \leq t_r$ , then  $C_q$  will become infected via  $C_i$ .

**Example 1.** Suppose that n = 4, the trace data consists of the triples

$$(C_1, C_2, 4), (C_2, C_4, 8), (C_3, C_4, 8), (C_1, C_4, 12),$$

and that the virus was inserted into computer  $C_1$  at time 2. Then  $C_3$  would be infected at time 8 by a sequence of three steps: first  $C_2$  becomes infected at time 4, then  $C_4$  gets the virus from  $C_2$  at time 8, and then  $C_3$  gets the virus from  $C_4$  at time 8.

**Example 2.** Suppose that n = 4, the trace data consists of

$$(C_2, C_3, 8), (C_1, C_4, 12), (C_1, C_2, 14),$$

and that the virus was inserted into computer  $C_1$  at time 2. Then  $C_3$  would not become infected during the period of observation: although  $C_2$  becomes infected at time 14,  $C_3$  only communicates with  $C_2$  before  $C_2$  was infected. There is no sequence of communications moving forward in time by which the virus could get from  $C_1$  to  $C_3$ .

**Project objective.** Write a program that takes a collection R of m triples, corresponding to the trace data for n computers, and preprocesses R in  $O(n + m \log m)$  time<sup>2</sup> to build a data structure G

<sup>&</sup>lt;sup>1</sup>We allow  $t_k$  to be equal to  $t_r$ . This just means that  $C_j$  had open connections to both  $C_i$  and  $C_q$  at the same time, and so a virus could move from  $C_i$  to  $C_q$ .

<sup>&</sup>lt;sup>2</sup>Worst-case or expected time bounds are equally acceptable for this project.

that, given the IDs for any two computers  $C_a$  and  $C_b$  and two times x, y, where  $x \le y$ , returns one of two things in O(m) time.

- If there is a sequence of communications such that a virus inserted into computer  $C_a$  at time x could have infected computer  $C_b$  by time y, then the program returns a list containing one such sequence.
- If no infection sequence exists, then the program returns null.

**Teamwork.** For this programming assignment, you should work in teams of two or three. It is your responsibility to assemble a team. Note that

- even if you signed up for a group for Project 1 on Canvas, you will need to sign up for a group for Project 2, and
- your group for Project 2 does not have to be the same as for Project 1.

Reference. This project is based on Exercise 11, Chapter 3, of

Jon Kleinberg and Éva Tardos, Algorithm Design, Addison-Wesley, 2006.

#### 2. The Data Structure

Our problem can be reduced to a graph connectivity problem that can be solved using breadth-first search or depth-first search. To do this, we use the triples in the trace data to construct a directed graph G as follows.

- Sort the triples by nondecreasing timestamp.
- Maintain a map where the keys are the computers and the associated values are lists. Initially, the list associated with each computer is empty.
- Next, scan the triples in sorted order.
- For each triple  $(C_i, C_i, t_k)$  we encounter in the scan do the following.
  - Create nodes  $(C_i, t_k)$  and  $(C_j, t_k)$ , if they do not already exist.
  - Add a directed edge from  $(C_i, t_k)$  to  $(C_i, t_k)$ .
  - Add a directed edge from  $(C_i, t_k)$  to  $(C_i, t_k)$ .
  - Append a reference to  $(C_i, t_k)$  to the list for  $C_i$ .
  - Append a reference to  $(C_i, t_k)$  to the list for  $C_i$ .
  - If  $(C_i, C_j, t_k)$  is not the first triple involving  $C_i$ , then add a directed edge from  $(C_i, t)$  to  $(C_i, t_k)$ , where t is the timestamp of the preceding element (the previously last one) in the list for  $C_i$ .
  - Do the analogous thing for  $C_i$ .

Figure 1 shows the graph for the trace data of Example 1 in the previous section.

Assume that G has been constructed. To determine whether a virus introduced at computer  $C_a$  at time x could have infected computer  $C_b$  by time y, we do the following.

- Walk through the list for  $C_a$  until we reach the first reference to a node  $(C_a, x')$  such that  $x' \ge x$ .
- Run BFS or DFS on G to determine all nodes reachable from  $(C_a, x')$ .
- If a node  $(C_b, y')$  with  $y' \le y$  is reachable from  $(C_a, x')$ , then we declare that  $C_b$  could have become infected by time y; otherwise, we declare that this is impossible.

You should verify for yourself that this algorithm indeed answers infection queries correctly in O(m) time.

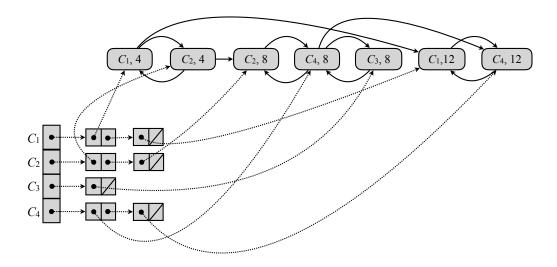


FIGURE 1. The graph *G* for the trace data of Example 1.

#### 3. Specifications

Your submission must be a Java program that contains the two classes described in this section: CommunicationsMonitor and ComputerNode. You *must* make these classes and their methods public. Each of the classes may contain additional methods apart from the required ones. Be judicious in designing the classes and the data structures.

- 3.1. **CommunicationsMonitor.** The CommunicationsMonitor class represents the graph G built to answer infection queries. It has the following methods.
  - CommunicationsMonitor(): Constructor with no parameters.
  - void addCommunication(int c1, int c2, int timestamp): Takes as input two integers c1, c2, and a timestamp. This triple represents the fact that the computers with IDs c1 and c2 have communicated at the given timestamp. This method should run in O(1) time. Any invocation of this method after createGraph() is called will be ignored.
  - void createGraph(): Constructs the data structure as specified in the Section 2. This method should run in  $O(n + m \log m)$  time.
  - List<ComputerNode> queryInfection(int c1, int c2, int x, int y): Determines whether computer c2 could be infected by time y if computer c1 was infected at time x. If so, the method returns an ordered list of ComputerNode objects that represents the transmission sequence. This sequence is a path in graph G. The first ComputerNode object on the path will correspond to c1. Similarly, the last ComputerNode object on the path will correspond to c2. If c2 cannot be infected, return null.

**Example 3.** In Example 1, an infection path would be  $(C_1, 4), (C_2, 4), (C_2, 8), (C_4, 8), (C_3, 8)$ .

This method can assume that it will be called only after createGraph() and that  $x \le y$ . This method must run in O(m) time. This method can also be called multiple times with different inputs once the graph is constructed (i.e., once createGraph() has been invoked).

• HashMap<Integer, List<ComputerNode>> getComputerMapping(): Returns a HashMap that represents the mapping between an Integer and a list of ComputerNode objects. The Integer represents the ID of some computer  $C_i$ , while the list consists of pairs

- $(C_i, t_1), (C_i, t_2), \ldots, (C_i, t_k)$ , represented by ComputerNode objects, that specify that  $C_i$  has communicated with other computers at times  $t_1, t_2, \ldots, t_k$ . The list for each computer must be ordered by time; i.e.,  $t_1 < t_2 < \cdots < t_k$ .
- List<ComputerNode> getComputerMapping(int c): Returns the list of ComputerNode objects associated with computer c by performing a lookup in the mapping.
- 3.2. **ComputerNode.** The ComputerNode class represents the nodes of the graph G, which are pairs  $(C_i, t)$ .
  - int getID(): Returns the ID of the associated computer.
  - int getTimestamp(): Returns the timestamp associated with this node.
  - List<ComputerNode> getOutNeighbors(): Returns a list of ComputerNode objects to which there is outgoing edge from this ComputerNode object.

#### 4. Guidelines on Code Submission

- Use the Java default package (unnamed package). While this is not good coding practice for larger applications, it is more convenient for testing.
- Make all the methods and constructors explicitly public.
- You may design helper classes and methods in addition to those listed in Section 3. However, every class and method listed in Section 3 must be implemented *exactly as specified*. This includes
  - the names of methods and classes (remember that Java is case-sensitive),
  - the return types of the methods, and
  - the types and order of parameters of each method/constructor.

If you fail to follow these requirements, you will lose a significant portion of the points, even if your program is correct.

- You are not allowed to have external JARs as dependencies. You may use inbuilt packages such as java.util.List.
- Please include all team members names as a JavaDoc comment in each of the Java files.
- Create the project folder as follows: <directory-name>/src/\*.java. This folder should nelude *only*.java files. Do not include any .class or other files.
- Create a zip file of your project folder.
- Upload your zip file on Canvas. Only one submission is necessary per team.

Your grade will depend on adherence to specifications, correctness, and efficiency.

Programs that do not compile will receive zero credit.

### **Important Note**

Some aspects of this specification are subject to change in response to issues detected by students or the course staff. *Check Canvas and Piazza regularly for updates and clarifications*.