

## Theory Problems

Course: Coursera Algorithms Specialization  
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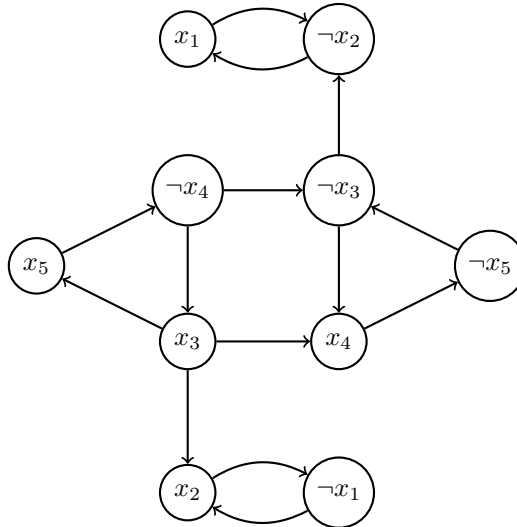
### Problem 1

In the 2SAT problem, you are given a set of clauses, where each clause is the disjunction of two literals (a literal is a Boolean variable or the negation of a Boolean variable). You are looking for a way to assign a value "true" or "false" to each of the variables so that all clauses are satisfied — that is, there is at least one true literal in each clause. For this problem, design an algorithm that determines whether or not a given 2SAT instance has a satisfying assignment. (Your algorithm does not need to exhibit a satisfying assignment, just decide whether or not one exists.) Your algorithm should run in  $O(m+n)$  time, where  $m$  and  $n$  are the number of clauses and variables, respectively. [Hint: strongly connected components.]

Each clause, a disjunction of 2 literals, can be represented as 2 implications. These implications can be used to create an implication graph. If the implication graph has any cycles containing a variable and its negation, a contradiction exists and therefore no satisfying assignment exists. Kosaraju's algorithm can be used to find the strongly connected components of a graph. It uses two depth-first searches, so it has  $O(m + n)$  complexity. If a variable and its negation are present in the same strongly connected component, no satisfying assignment exists.

Example: Given  $(x_1 \vee x_2) \wedge (\neg x_2 \vee x_3) \wedge (x_3 \vee x_4) \wedge (\neg x_3 \vee x_5) \wedge (\neg x_4 \vee \neg x_5) \wedge (\neg x_3 \vee x_4) \wedge (\neg x_1 \vee \neg x_2)$

$$\begin{aligned} (x_1 \vee x_2) &\leftrightarrow (\neg x_1 \rightarrow x_2) \wedge (\neg x_2 \rightarrow x_1) \\ (\neg x_2 \vee x_3) &\leftrightarrow (x_2 \rightarrow x_3) \wedge (\neg x_3 \rightarrow \neg x_2) \\ (x_3 \vee x_4) &\leftrightarrow (\neg x_3 \rightarrow x_4) \wedge (\neg x_4 \rightarrow x_3) \\ (\neg x_3 \vee x_5) &\leftrightarrow (x_3 \rightarrow x_5) \wedge (\neg x_5 \rightarrow \neg x_3) \\ (\neg x_4 \vee \neg x_5) &\leftrightarrow (x_4 \rightarrow \neg x_5) \wedge (x_5 \rightarrow \neg x_4) \\ (\neg x_3 \vee x_4) &\leftrightarrow (x_3 \rightarrow x_4) \wedge (\neg x_4 \rightarrow \neg x_3) \\ (\neg x_1 \vee \neg x_2) &\leftrightarrow (x_1 \rightarrow \neg x_2) \wedge (x_2 \rightarrow \neg x_1) \end{aligned}$$



A satisfying assignment exists.

## Problem 2

In lecture we define the length of a path to be the sum of the lengths of its edges. Define the bottleneck of a path to be the maximum length of one of its edges. A minimum-bottleneck path between two vertices  $s$  and  $t$  is a path with bottleneck no larger than that of any other  $s - t$  path. Show how to modify Dijkstra's algorithm to compute a minimum-bottleneck path between two given vertices. The running time should be  $O(m \log n)$ , as in lecture.

Dijkstra's algorithm can be modified so that the value stored in the priority queue (minimum heap) is the bottleneck instead of the total path length. Dijkstra's running time is  $O(m \log n)$ .

## Problem 3

We can do better. Suppose now that the graph is undirected. Give a linear-time  $O(m)$  algorithm to compute a minimum-bottleneck path between two given vertices.

Camerini's algorithm finds the minimum bottleneck spanning tree in  $O(m)$ . The minimum bottleneck path between any two vertices will follow the minimum bottleneck spanning tree between those two vertices. See Wikipedia article on Minimum Bottleneck Spanning Trees.

## Problem 4

What if the graph is directed? Can you compute a minimum-bottleneck path between two given vertices faster than  $O(m \log n)$ ?

Camerini's algorithm for directed graphs finds the minimum bottleneck spanning arborescence, a directed graph from source vertex to each of the other connected vertices, in  $O(m \log m)$ . Gabow and Tarjan's algorithm for minimum bottleneck spanning arborescence uses the modified version of Dijkstra's algorithm mentioned in Problem 2 and implements the priority queue using a Fibonacci heap. It runs in  $O(m + n \log n)$ . See Wikipedia article on Minimum Bottleneck Spanning Trees.

## Problem 5

Recall that a set  $H$  of hash functions (mapping the elements of a universe  $U$  to the buckets  $0, 1, 2, \dots, n - 1$ ) is universal if for every distinct  $x, y \in U$ , the probability  $\text{Prob}[h(x) = h(y)]$  that  $x$  and  $y$  collide, assuming that the hash function  $h$  is chosen uniformly at random from  $H$ , is at most  $\frac{1}{n}$ . In this problem you will prove that a collision probability of  $\frac{1}{n}$  is essentially the best possible. Precisely, suppose that  $H$  is a family of hash functions mapping  $U$  to  $0, 1, 2, \dots, n - 1$ , as above. Show that there must be a pair  $x, y \in U$  of distinct elements such that, if  $h$  is chosen uniformly at random from  $H$ , then  $\text{Prob}[h(x) = h(y)] \geq \frac{1}{n} - \frac{1}{|U|}$ .

Let  $\text{pairs}_{\text{total}}$  be the number of pairs of distinct elements in  $U$ . Let  $\text{pairs}_{\text{collide}}$  be the number of pairs in  $U$  that will collide when hashed. Let  $e_i$  be the number of elements in hash bucket  $i$ .

$$\begin{aligned}
pairs_{total} &= \binom{U}{2} = \frac{|U|(|U| - 1)}{2} \\
pairs_{collide} &= \sum_{i=1}^n \frac{e_i(e_i - 1)}{2} \\
pairs_{collide} &\geq \sum_{i=1}^n \frac{|U|}{2n} \left( \frac{|U|}{n} - 1 \right) = \frac{|U|}{2} \left( \frac{|U|}{n} - 1 \right) \\
Prob[h(x) = h(y)] &= \frac{pairs_{collide}}{pairs_{total}} \geq \frac{\frac{|U|}{n} - 1}{|U| - 1} > \frac{1}{n} - \frac{1}{|U|}
\end{aligned}
\qquad e_i = \frac{|U|}{n} \text{ best case}$$