

(1)

- Given an arbitrary graph G that may or may not be a DAG.
- While we have not visited all the nodes in the graph G or we have not found a cycle in the graph G .
- Try to find a node V with no incoming edges
- If we find a node V with no incoming edges, then we will find a topological order
 - Order the node V first
 - Delete V from G
 - Recursively compute a topological ordering of $G - \{V\}$ and append this order after V
- Graph G is DAG
 - Output the topological order of the reordered list
- If we fail to find a node V with no incoming edges, which means every node has at least one incoming edge, this shows that Graph G has a cycle
 - We are currently at an arbitrary node b
 - We will repeatedly follow an edge and pick the first adjacency edge among all the incoming edges
 - Record down every new visited node
 - Repeat this until we re-visit node b again
 - Output all the visited nodes between the two follow up visits of node b , this is our cycle for G . Graph G is not a DAG

① cont.

The algorithm takes $O(n)$ to search for a cycle because we are following the edges of the current node we are at until we reach back to the original node we start off

(2)

- We are using the algorithm to build an undirected and not connected graph because we will perform breadth first search and each component requires a starting node.
 - All our specimen are represented as vertex while any edge between two arbitrary vertex V_x and V_y are represented as the judgements between the two specimens.
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- For every component B of the graph G
 - Appoint a starting node n and give a letter label A
 - Make a note that node n as being visited
 - Create a output list for the breadth first search and record down node n . The output list is represented as Z
 - For the first layer of the BFS graph, label it with the starting node as $L(0) = n$
 - Let y be the index of traversing between different layers such that $y = 0, 1, 2, 3, 4, \dots$
 - For every node x within layer y which is $L(y)$
 - Examine every edge between node x and an arbitrary node w , which are edges that are incident to node w
 - If node w has not been visited which is has not been labeled yet
 - Label node w as being visited
 - If the judgement comes out as the same, then, it means node w has the same label as node x

- else, the judgement come out as different,
then it means node v has the opposite label of node x
- Endif
- Insert node v to the output list Z . Also, append
node v to the next layer which is $L(y+1)$

Endif
 Endfor
 Endfor
 Endfor

} refer to the starting statement
from the previous page

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- We are gonna check whether the m judgements are consistent
by running the following pseudocode.
 - Examine every single edge between node x and node w .
 - If the judgement comes out to be the same
 - If node x and node w don't have the same label
 - Output inconsistency message
 - Endif
 - Else the judgement comes out to be different
 - If node x and node w have the same label
 - Output inconsistency message
 - Endif
 - Endif
 - Endfor

③ - According to the problem statement, we know that the distance b between node s and t is greater than $n/2$ which means $b > n/2$

- We will use BFS algorithm to follow the path from the starting node s to the ending node t
- Let assume we will find node t at layer Y
- Let further assume that within layer $Y_1, Y_2, Y_3, \dots, Y_{b-1}$, there exists a single node.
- We want to eliminate the possibility that each layer Y_i could have at least size two to show there is at least one layer that has only one node.
- If the size is two for each layer, the distance will be included at least $2(\frac{n}{2}) = n$ nodes
- But, we only have n nodes and both node s and t are not within $Y_1, Y_2, Y_3, \dots, Y_{b-1}$ layers
- Therefore, there is a layer Y_i such that it only contains one node which is node v .
- The problem statement says that deleting node v from graph G will destroy all the paths from node s to node t
- Let set up a set of nodes that does not include node t within Y_1, Y_2, \dots, Y_{i-1} layers
This means that our set A is the union between node s and all the layers Y_1, Y_2, \dots, Y_{i-1}

- Any edge that are not included in set A should be included in the layer Y_i
- In order to move to node t , we will somehow get out from set A .
- This means that node v is the only node at layer Y_i .

④

- We will create a directed graph as our algorithm's solution.
- We first examine all the possible sorted triples.
- We create an array and each of the slot will point to a separate linked list
- Each of the linked list is related to computer C_a
- Throughout the scan, for every triple (C_i, C_j, t_k) we have examined, we gonna separate the triple into 2 separate nodes which are (C_i, t_k) & (C_j, t_k)
- We will connect the two nodes with edges that could go into either direction
- We will create a list for C_i and a list for C_j .
- The list for C_i will be constructed as linking all the nodes that has C_i but different communication time, such as $(C_i, 1)$ & $(C_i, 2)$
- The list for C_j will be constructed as linking all the nodes that has C_j but different communication time.
- If we have already passed the first triple that include C_i , we will draw a directed edge from (C_i, t_a) to (C_i, t_b) , t_a just refer to any previously last one node within the list of C_i .
- We will repeat the last bullet point for the list of C_j .
- Both C_i and C_j let us to keep track of each triple's nodes and edges by making the time constant.

- In order to determine whether a virus introduced at computer C_a at time x have infected computer C_b by time y , we need to examine the list of C_a
- We will traverse the list of C_a up to the point when we reach the last node in the list.
- Let assume that the first node is (C_a, x') and the last node is (C_a, x'') such that $x' \leq x$
- Starting from (C_a, x') , we will use BFS and trace all the possible reachable nodes from there.
- Throughout the journey, if there is a node (C_b, y') that is reachable such that $y' \leq y$
- Then, we will conclude that computer C_b might have been infected by virus by time y .
- If there isn't a node (C_b, y') that is reachable, then C_b might have not been infected by the virus.

- ⑤
- For this algorithm, we will represent the algorithm by a directed graph A .
 - People from the village will be denoted by P_x
 - Every single villager will have their own birth dates and death dates. They will be denoted by B_x and D_x
 - The edges in the graph will be represented as links that connect the current fact with the previous fact.
 - We will link up the birth dates and the death dates of each person. It is denoted by the set (B_x, D_x)
 - According to the problem statement, we will create an edge for (D_i, B_j) under the fact that "person P_i died before person P_j was born for some i and j "
 - According to the problem statement, we will create an edge for between two sets (B_i, D_j) and (B_j, D_i) under the fact that "the life spans of P_i and P_j overlapped at least partially for some i and j "
 - However, there are two possible situations that could have happened for the graph: whether the graph has a cycle or not

- First, if the graph has a cycle, this means that each event has to be in the order that correctly place before the next event.
- However, a cycle also means that there isn't an event which could be considered as the starting node. So, there does not exist an beginning or an end.
- Therefore, if our graph exists with cycles, the facts are not internally consistent
- Second, if the graph does not have a cycle, then we can represent the order of each person's birth date and death date in a topological order
- Therefore, if our graph does not exist with cycles, the facts are internally consistent.

6a)

- According to the problem statement, the array is sorted, therefore we could use binary search.
- We are looking for the first occurrence of array $G[x] = x + 1$ which is our missing number at the lowest index
- We will recursively search the first half of array G when $G[\frac{n}{2}] = \frac{n}{2} + 1$ and $x \leq \frac{n}{2}$
- We will recursively search the second half of array G
- In both cases, the time analysis is
$$O(n) = O(\frac{n}{2}) + \Theta(1)$$
- So, the worst case running time is
$$\boxed{O(\log n)}$$

6b)

- According to the problem statement, the array is not sorted, we need to check every slot of the array G for the missing element
- The algorithm works as follows:
- We will create a helper array W with index from 1 to $n+1$ & each slot of the array will be initialize to zero.
- For x from 1 to n , we scan through both array G & W based on $W[G[x]] = 1$
- Then, we scan through array W to search for the slot that contains a zero, the slot index will indicate our missing element.

Time Analysis:

Construct array $W \rightarrow O(n)$

Scan array $G \rightarrow O(n)$

Scan array $W \rightarrow O(n) +$

$$3O(n) \approx \boxed{O(n)}$$