Solutions of Introduction to Algorithms

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## Chapter 1

# **Dynamic Programming**

## 1.1 Rod cutting

**Exercise 2** No it cannot always produce an optimal solution. Consider the following example.

For n=3 the greedy approach cut the rod in 2 pieces. The length of one of them is 2 and the other's is 1. So the profit is 50\$ + 1\$ = 51\$. But the optimal solution is to keep the rod intact so the profit is 72\$.

Exercise 3 We can keep the rod intact so we don't need to incur the fixed cost c or we can have at least one cut. We need to choose the best solution among all of them:

$$r(i) = \begin{cases} \max_{1 \le k < n} (p_i, r(i-k) + p_k - c) & i > 0 \\ 0 & i = 0 \end{cases}$$

So the solution is r(n). We have n distinct subproblem. In each step we need to choose between keeping the rod intact or have at least one cut which divide the rod into two pieces. The length of one of them is k and the other's n-k. We don't know the exact value of k so we need to try all possible values. This can be done in O(n). Therefore the overall running time is  $O(n^2)$ 

```
1: function F(p, n, c)
       let r[0..n] be a new array
2:
       r[0] \leftarrow 0
3:
       for j from 1 to n do
4:
5:
           q \leftarrow p[j]
           for i from 1 to j-1 do
6:
               q = max(q, r[j-i] + p[i] - c)
7:
           end for
8:
           r[j] = q
9:
       end for
10:
       return r[n]
11:
12: end function
```

## 1.2 Matrix-chain multiplication

**Exercise 4** I've used the following equations:

$$\sum_{i=1}^{n} i = \frac{n(n+1)}{2} \tag{1.1}$$

$$\sum_{i=1}^{n} i^2 = \frac{n(n+1)(2n+1)}{6} \tag{1.2}$$

Each node of the graph represents a distinct sub-problem. Suppose we have two nodes v and u. There is an edge from v to u, if the solution of subproblem v is depended on subproblem u. In other words, there is an edge from m[i, j] to all m[i, k] and m[k + 1, j] for  $i \le k < j$ .

Usually |V| determines space complexity and |V| + |E| time complexity. we know for every subproblem m[i, j],  $j \ge i$ . Hence we have n - i + 1 subproblems which starts with  $A_i$ . So the number of vertices is:

$$|V| = \sum_{i=1}^{n} n - i + 1$$

$$= \sum_{i=1}^{n} i$$

$$= \frac{n(n+1)}{2}$$
(1.3)

Hence the space complexity is is  $O(n^2)$ . We don't use all of the array cells when j < i. So we waste  $\frac{n^2-n}{2}$  of allocated array. By analyzing lines 5 - 10 of MATRIX-CHAIN-ORDER pseudocode in the text book we can compute the number of edges. As you can see in line 10, m[i, j] is depends on two subproblem m[i, k] and m[k + 1, j]. We visit each distinct subproblem exactly once. So by counting the outdegree of each node we can calculate the number of edges in a

5 **for** 
$$l = 2$$
 **to**  $n$  //  $l$  is the chain length  
6 **for**  $i = 1$  **to**  $n - l + 1$   
7  $j = i + l - 1$   
8  $m[i, j] = \infty$   
9 **for**  $k = i$  **to**  $j - 1$   
10  $q = m[i, k] + m[k + 1, j] + p_{i-1}p_kp_j$ 

directed graph:

$$|E| = \sum_{l=2}^{n} \sum_{i=1}^{n-l+1} \sum_{k=i}^{i+l-2} 2$$

$$= \sum_{l=2}^{n} \sum_{i=1}^{n-l+1} 2(l-1)$$

$$= 2 \sum_{l=2}^{n} (n-l+1)(l-1)$$

$$= 2 \sum_{l=2}^{n} (n-(l-1))(l-1)$$

$$= 2 \sum_{l=1}^{n-1} (n-l)l$$

$$= 2 (\sum_{l=1}^{n-1} l - \sum_{l=1}^{n-1} l^{2})$$

$$= 2 (n \sum_{l=1}^{n-1} l - \sum_{l=1}^{n-1} l^{2})$$

$$= 2 [n \frac{(n-1)n}{2} - \frac{(n-1)(n)(2n-1)}{6}]$$

$$= n^{2} (n-1) - \frac{n(n-1)(2n-1)}{3}$$

$$= \frac{3n^{2}(n-1) - n(n-1)(2n-1)}{3}$$

$$= \frac{n(n-1)(3n-2n+1)}{3}$$

$$= \frac{n(n-1)(n+1)}{3}$$

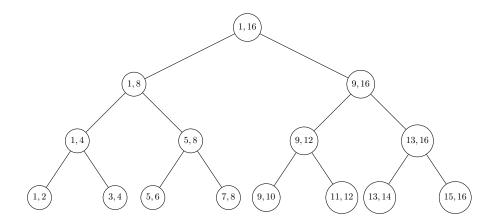
$$= \frac{n(n^{2}-1)}{3}$$

$$= \frac{n^{3}-n}{3}$$

So the running time is  $|V| + |E| = \frac{n^2 + n}{2} + \frac{n^3 - n}{3} = O(n^3)$ 

## 1.3 Elements of dynamic programming

Exercise 2 Each node is filled with (p, r). p is the index of leftmost element and r is the index of rightmost element of array which the subprolem wants to sort. As you can see there is no overlapping between subproblems so dynamic programming is not a good idea for merge sort. In other words, we don't see a previously solved subproblem again and we only waste memory. As a general rule if the subproblem graph is a tree, dynamic programming cannot be applied.



# Chapter 2

# Amortized Analysis

## 2.1 Aggregate analysis

**Exercise 1** No it doesn't hold. The maximum number of pops, including multipop, is proportional to the number of previous push operations. If we can only push one item, the number of pushed elements is at most n. If we add a new operation named multipush, then the number of pushed items is at most  $n \times k$ . So the amortized cost is O(k). For example we can have two operations. One is multipushing  $10^9$  items and the other is multipopping  $10^9$  items. It is obvious the total cost is not O(n) = O(2) and is  $O(nk) = O(2 \times 10^9)$ .

**Exercise 2** The following pseudo-code explains how to implement DECRE-MENT. The worst case happens when we start with 0 and then decrements it

```
1: function DECREMENT(A)
      while i < A.length and A[i] == 0 do
3:
          A[i] = 1
4:
         i = i + 1
5:
      end while
6:
      if i < A.length then
7:
          A[i] = 0
8:
      end if
9:
10: end function
```

to get  $2^k - 1$  which all bits are set to 1 and then increments it to get 0. We

repeat this loop until we have n operations. For n=4 and k=3 we have:

000

111

000

111

## 2.2 Accountant method

For another example of "accountant method" see exercise 5 of 3.1.

# Chapter 3

# Elementary Graph Algorithms

## 3.1 Representation of graphs

**Exercise 1** We know that adj[u] is a list. Depends on the list implementation, it can take O(1) to determine its size. In that case the running time for finding the out-degree of each vertex is O(V). If we cannot determine size of the list in O(1), then the overall running time of algorithm is O(V + E). The running time for finding in-degree of each vertex is O(V + E).

**Exercise 3** For adjacency-matrix it takes  $O(V^2)$  and for adjacency-list it takes O(V+E).

## **Algorithm 1** G' using adjacency matrix

```
1: function TransposeGraph(G)
      Let G' be a new graph
      G' \leftarrow G
3:
      for all u \in V do
4:
          for all v \in V do
5:
              G'.A[v][u] = G.A[u][v]
6:
7:
          end for
       end for
8:
      return G'
10: end function
```

## **Algorithm 2** G' using adjacency list

```
1: function TransposeGraph(G)
     Let G' be a new graph
2:
      G'.V = G.V
3:
      for all u \in G.V do
4:
         for all v \in G.Adj[u] do
5:
             G'. Adj[v].insert(u)
6:
7:
         end for
      end for
8:
9: end function
```

**Exercise 4** We create a new adjacency-list for G' called adj. For each vertex u in G, suppose v is its neighbor. If  $u \neq v$ , then adj[u].insert(v) and adj[v].insert(u). If there are multiple edges between u and v, we see v as u's neighbor more than once. So if the last element if adj[v] is u, it means there are more than one edges between them so we shouldn't insert v again. Traversing G takes O(V+E). Finding out there are more than one edge between two vertices is O(1). So the overall running time is O(V+E). Note that I supposed G is also undirected.

```
1: function F(G)
       let G' be a new graph
2:
3:
       G'.V = G.V
       for all u \in G.V do
4:
           for all v \in G.adj[u] do
5:
               if u \neq v \land G'.adj[v].last() \neq u then
6:
                  G'.adj[v].insert(u)
7:
               end if
8:
           end for
9:
10:
       end for
       return G'
11:
12: end function
```

**Exercise 5** The running time of matrix-list implementation is  $O(V^3)$ . For analyzing the running time of adjacency-list implementation we can use amortized analysis. We use "accountant method".

```
in_u: The number of edges that enter u out_u: The number of edges that leave u e_u: And edge from u to an arbitrary vertex v \neq u
```

We assign to all  $e_u$  cost  $c_{e_u} = 1 + in_u$ . Because by traversing the graph, we visit  $e_u$  at least once (line 6). For each edge that enters u we visit or revisit  $e_u$ 

(lines 7 - 8). We know that  $\sum_{u=1}^{|V|} in_u + \sum_{u=1}^{|V|} out_u = 2|E|$ . So we can easily calculate the total cost.

$$\sum_{e_u \in E} c_{e_u} = \sum_{e_u \in E} 1 + in_u$$

$$= \sum_{e_u \in E} 1 + \sum_{u=1}^{|V|} in_u$$

$$= |E| + \sum_{u=1}^{|V|} in_u$$

$$\leq 3|E|$$

We execute line 6 at most |E| times and lines 7 - 8 at most 2|E| times. So the total running time of algorithm using adjacency-list is O(|V|+3|E|) = O(V+E).

### Algorithm 3 Finding square graph using matrix-list

```
1: function MakeSquareGraph(G)
       Let G' be a new Graph
                                                                \triangleright G.A[1..|V|, 1..|V|]
 2:
       for all u \in G.V do
 3:
           for all v \in G.V do
 4:
               G'.A[u][v] = G.A[u][v]
                                                                     ▷ 1-edge paths
 5:
               if G.A[u][v] = 1 then
 6:
 7:
                  for all k \in G.V do
                      G'.A[u][k] = G.A[v][k]
                                                                     \triangleright 2-edge paths
 8:
                  end for
9:
               end if
10:
           end for
11:
       end for
13: end function
```

## Algorithm 4 Finding square graph using adjacency-list

```
1: function MakeSqureGraph(G)
       Let G' be a new graph
2:
       G'.V = G.V
3:
4:
       for all u \in G.V do
           for all v \in G.Adj[u] do
5:
              G'. Adj[u].insert(v)
                                                                   ▷ 1-edge paths
6:
              for all w \in G.Adj[v] do
7:
                  G'.Adj[u].insert(w)
                                                                   \triangleright 2-edge paths
8:
              end for
9:
          end for
10:
       end for
11:
12: end function
```

**Exercise 6** Suppose A is an adjacency matrix for G.

$$A[i,j] = \begin{cases} 1 & \text{i cannot be a universal sink} \\ 0 & \text{j cannot be a universal sink} \end{cases}$$

The following algorithm find the universal sink in O(V). In each step we remove one vertex from all candidates for "universal sink". It takes O(V) to have only one candidate. To determine that candidate is indeed a universal sink we need O(2V) operations. So the overall running time of algorithm is O(V) + O(2V) = O(V).

```
1: function GETUNIVERSALSINK(G)
                                                                          \triangleright A[1..|V|, 1..|V|]
        A = G.A
        u \leftarrow 1
 3:
        while u \leq |V| do
 4:
 5:
            v \leftarrow u + 1
            sink \leftarrow u
                                    \triangleright Vertices from sink to |V| can be universal sink
 6:
            while v \leq |V| \wedge A[u,v] = 0 do
                v \leftarrow v + 1
                                                          \triangleright v cannot be a universal sink
 8:
            end while
 9:
            u \leftarrow v
10:
                                               \triangleright u to v-1 cannot be a universal sink
        end while
11:
        for c from 1 to sink - 1 do
12:
            if A[sink, c] \neq 0 then
13:
                return "No universal sink"
14:
            end if
15:
        end for
16:
        for r \in V - \{sink\} do
17:
            if A[r, sink] \neq 1 then
18:
                return "No universal sink"
19:
            end if
20:
        end for
21:
        return sink
22:
23: end function
```

**Exercise 7** We know that B is an  $V \times E$  matrix which we show it as  $B_{V \times E}$ . By definition  $B^T$  is an  $E \times V$  matrix which we show it as  $B_{E \times V}^T$ . We define  $P_{V \times V} = B_{V \times E} \times B_{E \times V}^T$ .

$$p[i,j] = \sum_{k=1}^{E} b[i,k] \times b^T[k,j]$$

We consider two cases.

- 1.  $i \neq j$ : It is impossible that both b[i,k] and b[k,j] have the value of "1". Because the k<sup>th</sup> edge cannot enter both vertices i and j. With the same argument we can prove that both of them cannot have value of "-1". If the k<sup>th</sup> edge connect i to j, then b[i,k] = -1 and  $b^T[k,j] = 1$ . Otherwise both have value of zero. In other words, for  $i \neq j$  the value of p[i,j] is the number of edges between i and j.
- 2. i = j: It is obvious both b[i, k] and b[k, i] should have the same value. In this case p[i, i] is the sum of all edges that enter and leave the vertex i.

$$p[i,j] = \begin{cases} \text{number of edges between i and j} & i \neq j \\ indegree(i) + outdegree(i) & i = j \end{cases}$$

### 3.2 Breadth-first search

**Exercise 7** We need to determine whether an undirected graph is bipartite or not. We can paint the vertices of a bipartite graph with two colors in such a way that no two adjacent vertices share the same color.

We can easily prove that if there is a cycle in graph in which the number of edges is odd, then the graph cannot be bipartite.

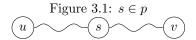
We can use BFS. We know that in an undirected graph we can only have tree and back edges (see 3.3). We run BFS on an arbitrary vertex s. Suppose u is reachable from s. If u.d is even we color that vertex "blue" otherwise we color it "red". For tree edges we don't have any problem. We need to think about back edges.

Suppose (u,v) is a back edge. Since we discover v first we can say  $v.d \leq u.d$ . We know that (v,u) is also an edge. According to BFS properties  $u.d \leq v.d+1$ . Hence  $0 \leq u.d - v.d \leq 1$ . If u.d = v.d+1, then u and v have different colors. So we only need to consider u.d = v.d. In that case both u and v have the same color and we need to prove that this graph cannot be bipartite. When we have a back edge it means that we have a cycle. The number of edges in this cycle is  $u.d+v.d+1=2\times u.d+1$  which is odd. So the graph cannot be bipartite. Note that the graph can have more than one connected component so it is possible we need to run BFS more than once. We use a new attribute u.bColor which we use for bipartite color as described.

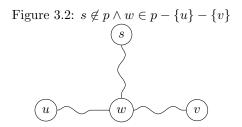
Algorithm 5 Determining whether a graph is bipartite or not

```
1: function IsBipartiteGraph(G)
       for all u \in G.V do
2:
          u.color = WHITE
 3:
          u.d = \infty
 4:
          u.\pi = NIL
 5:
       end for
 6:
       for all u \in G.V do
 7:
          if u.color == WHITE \land BFS(G, u) == FALSE then
 8:
9:
              return FALSE
          end if
10:
       end for
11:
       {\bf return}\ TRUE
12:
13: end function
 1: function BFS(G, s)
       s.color = GRAY
 2:
       s.bColor = 0
                              ▶ The color which we use for bipartite algorithm
3:
       s.d = 0
 4:
       Q=\emptyset
 5:
       ENQUEUE(Q, s)
 6:
       while Q \neq \emptyset do
 7:
          u = \text{Dequeue}(Q)
 8:
          for all v \in G.adj[u] do
9:
              if v.color == WHITE then
                                                      \triangleright edge (u, v) is a tree edge
10:
                 v.color = GRAY
11:
                 v.bColor = 1 - u.bColor
12:
                 v.d = u.d + 1
13:
                 v.\pi = u
14:
15:
                 ENQUEUE(Q, v)
              else if u.bColor == v.bColor then \triangleright edge (u, v) is a back edge
16:
                  return FALSE
17:
              end if
18:
          end for
19:
          u.color = BLACK
20:
21:
       end while
       {\bf return}\ TRUE
22:
23: end function
```

**Exercise 8** Suppose the maximum distance is path  $p = (v_0, v_1, \ldots, v_k)$  in which  $u = v_0$  and  $v = v_k$ . Consider an arbitrary vertex s. We know that there is exactly one path between every two vertices in a tree. We have two cases.



2.  $s \notin p$ : In this case there is exactly one path between s and  $w \in p - \{u\} - \{v\}$ . For example if w = u then the diameter is between s and v.



If we run BFS on s, then  $\max_{x \in G.V}(x.d)$  belongs to either u or v. Otherwise the diameter is not between u and v. Without loss of generality suppose it is  $u.d = \max_{x \in G.V}(x.d)$ . Then we run another BFS on u to get v in a similar manner. The running time of algorithm is O(2V+2E) = O(V+E). Since in a tree |E| = |V| - 1 the running time is O(V).

- 1: **function** FINDDIAMETER(G)
- 2: Let s be an arbitrary vertex such that  $s \in G.V$
- 3: INITBFS(G)
- 4: u = BFS(G, s)
- 5: INITBFS(G)
- 6: v = BFS(G, u)
- 7: **return** u, v, v.d
- 8: end function

```
1: function BFS(G, s)
       s.color = GRAY
2:
       s.d = 0
3:
       Q = \emptyset
 4:
       Engueue(Q, s)
 5:
 6:
       max = -\infty
       while Q \neq \emptyset do
 7:
           u = \text{Dequeue}(Q)
 8:
           for all v \in G.adj[u] do
9:
              \mathbf{if}\ v.color == WHITE\ \mathbf{then}
10:
11:
                  v.color = GRAY
                  v.d=u.d+1
12:
                  if v.d > max then
13:
                      max = v.d
14:
                      z = v
15:
                  end if
16:
                  v.\pi = u
17:
                  ENQUEUE(Q, v)
18:
              end if
19:
           end for
20:
           u.color = BLACK
21:
       end while
22:
       return z
23:
24: end function
```

**exercise 9** This undirected graph is equivalent to a directed graph which for all  $u, v \in V$ ,  $(u, v), (v, u) \in E$ . We can use a modified version of DFS. Because we have both edges (u, v) and (v, u), we don't have "cross edges". We need to choose between "forward edges" or "back edges". In the following algorithm we use "forward edges" and skip "back edges".

```
1: function DFS(G, u)
        u.color \leftarrow Gray
 2:
        paths \leftarrow \phi
 3:
        for all v \in G.Adj[u] do
 4:
 5:
            if v.color = White then
                                                                                ▶ Tree edge
 6:
                paths \leftarrow \{(u, v)\} \cup DFS(G, v) \cup \{(v, u)\}
            else if v.color = Black then
                                                                            ⊳ Forward edge
 7:
                paths \leftarrow paths \cup \{u, v\} \cup \{v, u\}
 8:
            end if
 9:
        end for
10:
11:
        u.color \leftarrow Black
        return paths
12:
13: end function
```

## 3.3 Depth-first search

Edge classification Suppose s is the root of DFS or BFS tree.

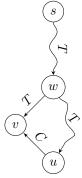
- Tree edge
  - Directed graph
    - \* DFS: We can have tree edges
    - $\ast\,$  BFS: We can have tree edges
  - Undirected graph
    - \* DFS: We can have tree edges
    - \* BFS: We can have tree edges
- Forward edge
  - Directed graph
    - \* DFS: We can have forward edges
    - \* BFS: We can't have forward edges. Suppose we can have forward edge (u,v). According to forward edge properties we can say u.d < v.d (if u.d = v.d it's cross edge) and according to BFS properties  $v.d \le u.d+1$ . Hence  $0 < v.d-u.d \le 1$ . If v.d = u.d+1 it is a tree edge, unless we have a multigraph. So (u,v) cannot be a forward edge.
  - Undirected graph
    - \* DFS: We can't have forward edges
    - \* BFS: We can't have forward edges
- Back edge
  - Directed graph

- \* DFS: We can have back edges
- \* BFS: We can have back edges. Suppose (u, v) is a back edge. Since we discover v first we can say  $v.d \le u.d$ . According to BFS properties we can say  $v.d \le u.d + 1$ . Hence  $v.d u.d \le 1$ . Note that we have only upper bound. So it is possible v.d u.d < 0
- Undirected graph
  - \* DFS: We can have back edges
  - \* BFS: We can have back edges. Suppose (u,v) is a back edge. Since we discover v first we can say  $v.d \le u.d$ . We know that (v,u) is also an edge. According to BFS properties  $u.d \le v.d+1$ . Hence  $0 \le u.d v.d \le 1$

#### • Cross edge

- Directed graph
  - \* DFS: We can have cross edges
  - \* BFS: We can have cross edges. Suppose (u,v) is a cross edge. Since we discover v first we can say  $v.d \leq u.d$ . According to BFS properties we can say  $v.d \leq u.d + 1$ . Hence  $v.d u.d \leq 1$ . Note that we only have upper bound. It is possible that v.d u.d < 0. See figure 3.3
- Undirected graph
  - \* DFS: We can't have cross edges
  - \* BFS: We can't have cross edges

Figure 3.3: BFS – cross edge in a directed graph.  $v.d \leq u.d + 1$ 



**Exercise 1** You can use the following facts. Suppose we have edge (u, v) and we consider loops as back edges.

Tree edge: u.d < v.d < v.f < u.fForward edge: u.d < v.d < v.f < u.fBack edge:  $v.d \le u.d < u.f \le v.f$ Cross edge: v.d < v.f < u.d < u.f

Note that when the graph is undirected we don't have "forward edge" and "cross edge". Because they are equivalent to "back edge" and "tree edge" respectively.

Table 3.1: Directed graph

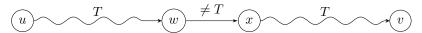
	white	gray	black
white	tree, back, forward, cross	back, cross	cross
gray	tree, forward	tree, back, forward	tree, forward, cross
black	impossible	back	tree, back, forward, cross

Table 3.2: Undirected graph

	G of				
	white	gray	black		
white	tree, back	tree, back	impossible		
gray	tree, back	tree, back	tree, back		
black	impossible	tree, back	tree, back		

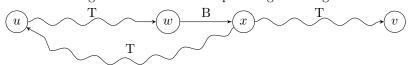
**Excercise 8** We need to show some examples which there is only one path from u to v and at least of one the edges in this path is non-tree edge. Without loss of generality, suppose in this path except e = (w, x) which is a non-tree edge, all other edges are tree ones. We consider all possible types.

Figure 3.4: Edge e = (w, x) is a non-tree edge



- Forward Edge: If (w, x) is forward ege, then w is an ancestor of x which leads to v be a descendant of u. So it cannot be a forward edge
- Cross edge: If (w, x) is a cross edge, then x finishes before the discovery of w. In other words, all reachable vertices from x, including v, will be discover before w and u. So it cannot be a cross edge
- Back edge: Consider the following example which the root of DFS tree is vertex x and it discover u before v.

Figure 3.5: Counterexample using back edge



Excercise 9 Suppose we have edge (u, v) and we consider loops as back edges.

Tree edge: u.d < v.d < v.f < u.fForward edge: u.d < v.d < v.f < u.fBack edge:  $v.d \le u.d < u.f \le v.f$ Cross edge: v.d < v.f < u.d < u.f

As you can see only in cross edge the discovery of one endpoint is after the other finished. We need to prove that if there is a path from u to v, it is possible to have v.d > u.f. In other words, edge (v,u) is a cross edge and there is a path from u to v in which there is at least one edge which is not a tree edge. We can make counterexample even simpler by removing the cross edge. Note that the root of DFS tree is vertex s. In general if there is a cycle from s to u and u to s, then it is possible v.d > u.f.

Figure 3.6: Counterexample using cross edge

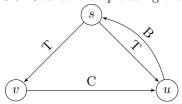


Figure 3.7: Counterexample without cross edge

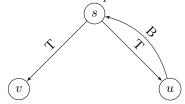
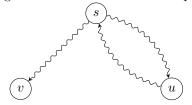


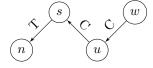
Figure 3.8: General counterexample



Exercise 10 In an undirected graph forward edges are equivalent to back edges and cross edges are equivalent to tree edges. Although the following modification works for both directed and undirected graph, you can remove portion of code that is related to "forward" and "cross" edges to save space.

```
1: function DFS-VISIT(G, u)
2:
      time = time + 1
      u.d = time
3:
      u.color = GRAY
4:
      for all v \in G.adj[u] do
5:
          if v.color == WHITE then
                                                   \triangleright edge (u, v) is a tree edge
6:
             PRINT-EDGE(u, v, TREE)
7:
8:
             v.\pi = u
             DFS-VISIT(G, v)
9:
          else if v.color == GRAY then
                                                    \triangleright edge u, v is a back edge
10:
             PRINT-EDGE(u, v, BACK)
11:
          else if u.d < v.d then
12:
13:
             PRINT-EDGE(u, v, FORWARD)
14:
             PRINT-EDGE(u, v, CROSS)
15:
          end if
16:
      end for
17:
      u.color = BLACK
18:
      time = time + 1
19:
      u.f = time
20:
21: end function
```

**Exercise 11** If both incoming and outgoing edges are cross, that happens. Consider the following example. Suppose DFS starts at s, then u and finally at w



end for

14: end function

5:

1: function DFS(G) 2: for all  $u \in G.V$  do 3: u.color = WHITE4:  $u.\pi = NIL$ 

**Algorithm 6** Connected components in an undirected graph

```
6: time = 0

7: ccn = 0 \triangleright ccn is the number of connected components

8: for all u \in G.V do

9: if u.color == WHITE then

10: ccn = ccn + 1

11: DFS-VISIT(G, u)

12: end if

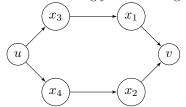
13: end for
```

```
1: function DFS-VISIT(G, u)
      time = time + 1
2:
3:
      u.d = time
4:
      u.cc = ccn
      u.color = GRAY
5:
      for all v \in G.adj[u] do
6:
          if v.color == WHITE then
7:
8:
             v.\pi = u
9:
             DFS-VISIT(G, v)
          end if
10:
      end for
11:
      u.color = BLACK
12:
      time = time + 1
13:
      u.f = time
15: end function
```

Exercise 13 It is obvious that we should only have tree and back edges. It is important to start from the right vertex. Consider figure 3.9. There is exactly

two distinct paths between u and v. If we start the DFS from u, in the first run we can detect that the graph is not singly connected. I thought I can design an

Figure 3.9: non singly connected graph



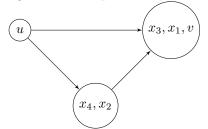
O(V+E) algorithm to solve this problem. I was wrong.

Wrong idea Start DFS from an arbitrary vertex s. If you found "forward" or "cross" edges then it is not singly connected so the algorithm terminates. After DFS finished, it is possible we have unvisited vertices.

**component:** All undiscovered vertices which will be discover in one DFS run. For example if we run DFS from  $x_3$  in figure 3.9,  $x_3$ ,  $x_1$  and u are belong to the same component.

We can determine and number the components (similar to connected components). If we find "forward" or "cross" edges within each component, the algorithm terminates. Note that "cross" edges between components is not trivial. We can have singly connected graph which has at least one cross edge between its components. We can create a new graph which its vertices are the components of input graph and its edges are the cross edges between components. Suppose in figure 3.9 we run DFS first on  $x_3$ , then  $x_4$  and finally u. You can see the result in figure 3.10. This algorithm is not always correct. Suppose we run DFS first on v,  $x_1$ ,  $x_2$ ,  $x_3$ ,  $x_4$  and finally u. As you can see we don't have any tree edges and the graph of components is same as original one.

Figure 3.10: Graph of components



# Chapter 4

# Minimum Spanning Trees

For questions similar to Prim's minimum spanning tree see "Variants" in chapter 5.

## 4.1 Problems

### Problem 3 Bottleneck spanning tree

- **a.** Consider an arbitrary MST T. Suppose the maximum-weight edge in T is e. If we remove that edge we have a forest of two trees  $C_1$  and  $C_2$ . Now consider MST T' whose maximum-weight edge, e', is less that e. There should be an edge in T' that connects  $C_1$  to  $C_2$ . The weight of that edge should be less than e. In other words, e is not a light edge which contradicts T is an MST.
- **b.** We remove all edges in G.E which their weight are higher than b. We called the modified graph  $G_b$ . It is obvious that  $G_b.V = G.V$  and  $G_b.E = \{(u,v) \in G.E : w(u,v) \leq b\}$ . If  $G_b$  remains connected then every spanning tree of  $G_b$  doesn't have an edge whose weight is greater than b. The running time of this algorithm is O(V+E). Since G should be a connected graph,  $|E| \geq |V| 1$ . So we can say the running time is O(E).

```
1: function VALID-BST-VALUE(G, b)
      G' = REMOVE-EDGES(G, b)
2:
      INITDFS(G')
3:
                                   ▶ The number of connected components
      c = 0
4:
      for all u \in G'.V do
5:
         if u.color == WHITE then
6:
            c = c + 1
7:
            DFS(G', u)
8:
         end if
9:
      end for
10:
      return (c == 1)
11:
12: end function
```

```
1: function REMOVE-EDGES(G, b)
        G_b.V = G.V
2:
        G_b.E = \emptyset
3:
        for all (u, v) \in G.E do
4:
           if (u, v).c \le b then
                                                                    \triangleright (u,v).c \equiv w(u,v)
5:
               G_b.E = G_b.E \cup \{(u,v)\}
6:
            end if
7:
        end for
8:
        return G_b
9:
10: end function
```

- **c.** We need some definitions:
- $G_b$ : a new sub-graph of G which  $G_b.V = G.V$  and  $G_b.E = \{(u,v) \in G.E : w(u,v) \leq b\}$
- comp(u): Suppose connected components  $c_1, c_2, \ldots, c_k$  make the Graph  $G_b$  and vertex u belongs to the i<sup>th</sup> component. Then comp(u) = i
- $G'_b$ : Suppose  $C = \{c_1, c_2, \ldots, c_k\}$  is the set of connected components of  $G_b$ . Then  $G'_b.V = C$  and  $G'_b.E = \{(u,v) \in G.E : comp(u) \neq comp(v)\}$ . It is possible that there are more than one edge between  $c_i$  and  $c_j$ . In this case we choose the light edge (minimum weight)

We want to find  $b_m = \min_{b \in G.E \land |G'_b,V|=1}(b)$ . In other words, we want to find the minimum of  $b \in G.E$  which  $G_b$  is a connected graph. Consider an arbitrary edge (u,v). Suppose w(u,v) = b There are two cases:

- 1.  $G_b$  is connected: It means  $b_m < b$ . More precisely,  $b_m \in G_b.E$
- 2.  $G_b$  is not connected: It means  $b_m \geq b$ . More precisely,  $b_m \in G_b'.E$ .

Consider set  $W = w(u, v) : (u, v) \in G.E$ . We can find the median of W in O(E) by "median of medians" algorithm. After finding the median, we can divide

4.1. PROBLEMS 25

the edges into two equal sets:  $G_b.E$  and  $G'_b.E$ . In each step we eliminate half of edges. For simplicity we assume edge (u, v) has an attribute c such that (u, v).c = w(u, v).

```
1: function BST(G)
       if |G.E| == 1 then
 2:
 3:
            return G.E
        end if
 4:
       m = \text{FIND-MEDIAN}(G.E)
                                                                                 \triangleright O(E)
 5:
       if VALID-BST-VALUE(G, m) then
 6:
           G_b = REMOVE-EDGES(G, m)
                                                                                 \triangleright O(E)
 7:
            R = BST(G_b)
 8:
9:
       else
           G_b' = \mathsf{MAKE}\text{-}G_b'(\mathsf{G})
                                                                                 \triangleright O(E)
10:
            R = BST(G'_b)
11:
        end if
12:
       {\bf return}\ R
13:
14: end function
```

```
1: function MA\overline{KE-G'_b(G)}
       C, comp = CC(G)
2:
       G_b'.V = C
3:
       for all (u, v) \in G.E do
 4:
           if comp[u] \neq comp[v] then
 5:
               if (comp[u], comp[v]) \not\in G'_b.E then
 6:
                   (comp[u], comp[v]).c = (u, v).c
 7:
                  G'_b.E = G'_b.E \cup \{(comp[u], comp[v])\}
 8:
               else if (u, v).c < (comp[u], comp[v]).c then
9:
                   (comp[u], comp[v]).c = (u, v).c
10:
               end if
11:
           end if
12:
       end for
13:
       return G'_b
14:
15: end function
```

```
1: function CC(G)
      Let comp[1..G.V] be a new array
2:
       C = \emptyset
                                           ▶ The set of connected components
3:
       c = 0
                                      ▶ The number of connected components
4:
       INIT-DFS(G)
5:
6:
       for all u \in G.V do
          if u.color == WHITE then
7:
              c = c + 1
8:
              C = C \cup \{c\}
9:
              DFS(G, u, c, comp)
10:
11:
          end if
       end for
12:
       return C, comp
13:
14: end function
```

```
1: function DFS(G, u, c, comp)
2: u.color = GRAY
3: comp[u] = c
4: for all v \in G.adj[u] do
5: if v.color == WHITE then
6: DFS(G, v, c, comp)
7: end if
8: end for
9: end function
```

So the total running time of algorithm is O(E):

$$T(E) = T(\frac{E}{2}) + O(E)$$

$$= O(E) + O(\frac{E}{2}) + O(\frac{E}{4}) + \dots + O(\frac{E}{2^{i}}) + \dots + O(1)$$

$$= O(\frac{E}{2^{0}}) + O(\frac{E}{2^{1}}) + \dots + O(\frac{E}{2^{\log_{2} E}})$$

$$= \frac{E(\frac{1}{2})^{\log_{2} E + 1} - E}{\frac{1}{2} - 1}$$

$$= 2E - 1$$

For the analysis of run-time we assumed G is connected so  $|E| \ge |V| - 1$ .

## Chapter 5

# Single-Source Shortest Path

#### Variants

#### 1. Single-source shortest-path problem:

(a) acm-icpc World Finals 2002 question C, Crossing the Desert: You can see the problem statement in "DESERT Problem in SPOJ" and "Problem 1011 in UVa" online judges.

In this problem, you will compute how much food you need to purchase for a trip across the desert on foot.

At your starting location, you can purchase food at the general store and you can collect an unlimited amount of free water. The desert may contain oases at various locations. At each oasis, you can collect as much water as you like and you can store food for later use, but you cannot purchase any additional food. You can also store food for later use at the starting location. You will be given the coordinates of the starting location, all the oases, and your destination in a twodimensional coordinate system where the unit distance is one mile. For each mile that you walk, you must consume one unit of food and one unit of water. Assume that these supplies are consumed continuously, so if you walk for a partial mile you will consume partial units of food and water. You are not able to walk at all unless you have supplies of both food and water. You must consume the supplies while you are walking, not while you are resting at an oasis. Of course, there is a limit to the total amount of food and water that you can carry. This limit is expressed as a carrying capacity in total units. At no time can the sum of the food units and the water units that you are carrying exceed this capacity.

You must decide how much food you need to purchase at the starting location in order to make it to the destination. You need not have any food or water left when you arrive at the destination. Since the general store sells food only in whole units and has only one million

food units available, the amount of food you should buy will be an integer greater than zero and less than or equal to one million.

Input The first line of input in each trial data set contains n  $(2 \le n \le 20)$ , which is the total number of significant locations in the desert, followed by an integer that is your total carrying capacity in units of food and water. The next n lines contain pairs of integers that represent the coordinates of the n significant locations. The first significant location is the starting point, where your food supply must be purchased; the last significant location is the destination; and the intervening significant locations (if any) are oases. You need not visit any oasis unless you find it helpful in reaching your destination, and you need not visit the oases in any particular order.

The input is terminated by a pair of zeroes.

**Output** For each trial, print the trial number followed by an integer that represents the number of units of food needed for your journey. Use the format shown in the example. If you cannot make it to the destination under the given conditions, print the trial number followed by the word "Impossible."

Place a blank line after the output of each test case.

## Example

#### Input

4 100

10 - 20

-10 5

 $30 \ 15$ 

 $15 \ 35$   $2 \ 100$ 

0.0

100 100

0.0

#### Output

Trial 1: 136 units of food Trial 2: Impossible

**Solution:** First we make question simpler. So we suppose it is impossible to leave food on oases or starting location and possibly return and collect them.

We can model this problem to an undirected graph. The vertices are the starting location, oases and the destination. There is an

edge between u and v, if the amount of required food and water doesn't exceed C.

 $-f_{u,v}$ : The amount of required food from u to v  $-a_{u,v}$ : The amount of required water from u to vWe define weight function w:

$$w(u,v) = \begin{cases} f_{u,v} & f_{u,v} + a_{u,v} \le C \\ \infty & f_{u,v} + a_{u,v} > C \end{cases}$$

Unlike food, we can pick up water in every oases. So we need to order all required food in the starting location. Because we cannot leave food anywhere in the desert, the final path should be simple. Otherwise we have at least one cycle. If we remove that cycle we obtain an equivalent path which required less food. Suppose path p which connects the starting location to the target is an optimal path. We define  $a_m = \max_{(u,v) \in p} (a(u,v))$ . We called

p a valid path if  $\sum_{(u,v)\in p} f(u,v) + a_m \leq C$ . The required food for p must be minimum among all valid paths from the starting location to the destination. We can solve this problem with a greedy algorithm similar to Dijkstra's shortest path. u.d is the amount of required food from the starting location to u. We define a new attribute  $u.a_m$  which we described it before. s is the starting location and t is the target location. The running time of algorithm is like Dijkstra's shortest path which can be  $O(E \log V)$ .

```
1: function Desert(G, w, C, s, t)
       INITIALIZE-SINGLE-SOURCE(G, s)
2:
       S = \emptyset
3:
       Q = G.V
4:
       while Q \neq \emptyset do
5:
          u = \text{EXTRACT-MIN}(G)
6:
          S = S \cup \{u\}
7:
          for all v \in G.Adj[u] do
8:
              RELAX(u, v, w, C)
9:
          end for
10:
       end while
11:
12:
       if t.d < \infty then
13:
          return t.d
       else
14:
           "IMPOSSIBLE"
15:
       end if
16:
17: end function
```

```
1: function RELAX(u, v, w, C)
      m = \max(w(u, v), u.a_m)
2:
      food = u.d + w(u, v)
3:
      if (food + m) \le C \land food < v.d then
4:
5:
         v.d = food
6:
         v.a_m = m
7:
         v.\pi = u
      end if
8:
9: end function
```

```
\begin{array}{l} \textbf{function} \ \text{INITIALIZE-SINGLE-SOURCE}(\mathbf{G},\,\mathbf{s}) \\ \textbf{for all} \ v \in G.V \ \textbf{do} \\ v.d = \infty \\ v.\pi = NIL \\ v.a_m = -\infty \\ \textbf{end for} \\ s.d = 0 \\ \textbf{end function} \end{array}
```

### 2. Single-destination shortest-path problem:

(a) You are given flight schedules between a set of n cities. For each pair of cities (i,j) between which there is a direct flight, you are given the pair  $(d_{ij}, a_{ij})$ , the departure and arrival time of the flight from city i to city j. Assume that there is at most one flight from city i to city j per day. Suppose you start at city A and want to reach city B. You have an important meeting in city B that you need to attend, and you need to reach city B latest by time t. Give an algorithm that outputs a possible sequence of flights you could take starting from city A as late as possible and reaching city B before time t, with at least one hour layover between any two consecutive flights.

Solution: We don't know which flight in A we should choose. We can solve the problem if we consder flights in B. Given the graph G, we need to change that to graph G' such that G'.V = G.V and  $G'.E = \{(u,v): (v,u) \in V.E\}$ . Hence if  $(i,j) \in G'.E$ , there is a flight from j to i in which the departure time is  $d_{ji}$  and arrival time is  $a_{ji}$ . In B we only need to consider all flights  $C = \{(B,u) \in G'.E: a_{uB} \leq t\}$  and choose the edge with latest departure time  $(\max_{(B,u)\in C}(d_{uB}))$ . Because if we arrive at u, flight (u,B) has the latest departure time and it doesn't make any sense to go from u to B through other intermediate vertices. So we add edge (u,B) as an optimal answer between u and B (in G' we should say edge (B,u)). This algorithm is similar to Prim's minimum spanning tree. We can call it "Single-

destination latest-departure problem". We calculate the best possible sequence of flights from u to B. Eventually we calculate an optimal path from A to B. By "best" we mean the departure time of the first flight is as late as possible and the arrival time is at most t and there is a layover of at least one hour between two consecutive flights. u.d store the latest possible departure from u to B. We add a dummy flight from B to an unknown place with departure t+1 to discard all those flights to B with arrival time greater than t. Q contains all those vertices which we don't know yet an optimal flight sequence from them to B. On the other hand, S contains all those vertices which we found out an optimal flight sequence from them to B.

```
1: function SCHEDULING(G, A, B, d, a, t)
       G' = REVERSE-GRAPH-EDGES(G)
       INITIALIZE-SINGLE-SOURCE(G', s)
 3:
       B.d = t + 1
                                 \triangleright To make sure we arrive at B no more than t
 4:
       S = \emptyset
 5:
       Q = G'.V
 6:
       while Q \neq \emptyset do
 7:
          u = \text{EXTRACT-MAX}(G')
 8:
          S = S \cup \{u\}
 9:
          for all v \in G'. Adj[u] do
10:
              RELAX(u, v, d, a)
11:
          end for
12:
       end while
13:
       if A.d > -\infty then
14:
          u = A
15:
          while u \neq B do
16:
              PRINT(u, u.\pi)
17:
              u = u.\pi
18:
19:
          end while
       else
20:
          PRINT("IMPOSSIBLE")
21:
       end if
22:
23: end function
```

```
1: function RELAX(u, v, d, a)
2: if (a_{vu} + 1) \le u.d \land d_{vu} > v.d then
3: v.d = d_{vu}
4: v.\pi = u
5: end if
6: end function
```

```
\begin{array}{l} \textbf{function} \ \text{INITIALIZE-SINGLE-SOURCE}(G,\,s) \\ \textbf{for all} \ v \in G.V \ \textbf{do} \\ v.d = -\infty \\ v.\pi = NIL \\ \textbf{end for} \\ \textbf{end function} \end{array}
```

- 3. Single-pair shortest-path problem:
- 4. All-pair shortest-path problem:

## 5.1 Dijkstra's algorithm

#### 5.1.1 Exercises

**Exercise 6** Suppose the path  $p = (v_0, v_1, \ldots, v_k)$  in which  $v_0 = u$  and  $v_k = v$  is one of the paths between u and v. Since the probabilities are independent we want to find  $\max(\prod_{i=0}^{k-1} r(v_i, v_{i+1}))$ . We can reduce the problem to a shortest-path one by changing the weight function w(u, v) = r(u, v) to  $w'(u, v) = -\log r(u, v)$ .

$$0 \le r(u, v) \le 1$$
$$\log 0 \le \log r(u, v) \le \log 1$$

If we define  $\log 0 = -\infty$ , then  $-\infty \le \log r(u, v) \le 0$  which is equivalent to  $0 \le -\log r(u, v) \le \infty$ .

$$\max(\prod_{i=0}^{k-1} r(v_i, v_{i+1}))$$

$$\equiv \max(\sum_{i=0}^{k-1} \log r(v_i, v_{i+1}))$$

$$\equiv \min(\sum_{i=0}^{k-1} -\log r(v_i, v_{i+1}))$$

which is exactly the shortest path problem. Since w'(u,v) is non-negative, we can use Dijkstra algorithm which its run-time can be  $O(E \log V)$ . We only need to change RELAX function.

```
1: function w'(\mathbf{u}, \mathbf{v}, \mathbf{w})

2: if w(u, v) == 0 then

3: return \infty

4: else

5: return -\log w(u, v)

6: end if

7: end function
```

```
1: function RELAX(u, v, w)

2: if v.d > u.d + w'(u, v, w) then

3: v.d = u.d + w'(u, v, w)

4: v.\pi = u

5: end if

6: end function
```