Matching mechanisms for kidney transplantations

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1 Two Simple Approaches

Question 1

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Algorithm 1: Direct Donation
   Input: A number n of patient and, for all patients i \in \mathbb{N}, their set of
            compatible donors K_i \subseteq [1, n].
   Output: A set of pairs donor-patient pairs c and a waiting list w.
 1 initialization
       c = \emptyset;
       w = [];
 4 begin
       for i = 1 to n do
 5
           if i \in K_i then
 6
            c = c \cup \{(i, i)\};
 7
           else
 8
               w.add(i)
       return c, w
10
```

The implementation in python is pretty straightforward. The python function takes as parameter an integer n and an integer set array of size n. The use of python set allows to test the if condition in constant times. The only difference with pseudo-code is the use of a list rather than a set for the variable c. This choice was made to avoid the unnecessary heaviness of the set data structure, but does not change the complexity.

Question 2

Our matching algorithm must prioritize the patients with the highest priority. It must also ensure that all patients will leave with a donor that suited them better than their paired donor and the waiting list, otherwise some patients may not participate in the exchange program.

Our greedy algorithm therefore consists of browsing the patients in descending order of priority and matching them to their preferred donor whose paired patient does not prefer the waiting list or his donor to the donor of the priority patient.

Algorithm 2: Greedy Matching

Input: A number n of patient, a strict priority list of the patients U and, for all patients $i \in \mathbb{N}$, their set of compatible donors $K_i \subseteq \llbracket 1, n \rrbracket$ and their strict preference relation P_i over $K_i \cup \{k_i, w\}$. The priority list is an integer list that starts with the index of the highest priority patient and goes to the index of the lowest priority patient.

```
Output: A matching M.
```

```
1 initialization
      M = \emptyset;
 2
      is matched = Boolean[1..n];
 3
      sorted K = IntegerList[1..n];
 4
      for i = 1 to n do
 5
 6
         is matched[i] = False;
         sorted K[i] = sort(K_i, P_i);
                                                            // \mathcal{O}(n\log(n))
 8 begin
      /* Iterate over all patient by decreasing priority
                                                                         */
      for not U.is empty() do
 9
          u = U.pop();
10
          if not is matched[u] then
11
             while not sorted K[u].is empty() do
12
                 v = sorted K[u].pop();
13
                 if not is matched[v] then
14
                     /* If t_v prefer k_u over k_v and w
                                                                         */
                    if t_v P_v t_u and t_v P_v w then
15
                        M = M \cup \{((k_u, t_u), (k_v, t_v))\};
16
                        is matched[u] = True;
17
                        is matched[v] = True;
18
19
      return M
```

We start by initializing an $is_matched$ array to keep track of which pair has already been matched and an integer list array $sorted_K$ that contains, for each patient of index i, the indexes of the preferred donors in descending order of preference. The sort function returns a list of the elements of K_i sorted in descending order according to the preference relation P_i .

2 Efficient stategy-proof exchange mechanism

Question 3

Let (k_1, t_1) be a pair of donor-patient and assume there is no cycle. Consider the chain starting from (k_1, t_1) . Since there is a finite number n of donor-patient pairs and there is no cycle, the chain cannot be infinite. Finally, since the only way for a chain to stop is that w belong to the chain, (k_1, t_1) is indeed a tail of a w-chain.

Thus either there exists a cycle, or each pair is the tail of some w-chain.

Question 4

Question 5

Let's run the Cycles and Chains Matching Algorithm with rule A on this example. Graphs associated with every round are shown in figure 1.

The initial graph contains a cycle $C_1 = (k_3, t_3, k_2, t_2, k_{11}, t_{11})$. The first round of the algorithm therefore consists in detecting this cycle, allocating the kidneys to the patients and then eliminating the cycle.

After removing the first circle, a new circle $C_2 = (k_5, t_5, k_7, t_7, k_6, t_6)$ is formed. In the second round, this cycle is therefore deleted.

At the start of the third round, there is no more circle. The graph contains 2 w-chains of length 6: $W_1 = (k_8, t_8, k_4, t_4, k_9, t_9)$ and $W_2 = (k_{10}, t_{10}, k_1, t_1, k_9, t_9)$. There are no longer w-chains and since the highest priority patient t_1 is only present in W_2 , W_2 is chosen. The kidneys are allocated and the chain remains in the graph for the next round but patients pointing to other kidneys than k_10 in the chain are redirected.

After the redirect, a new circle $C_3 = (k_4, t_4, k_8, t_8)$ has formed. The fourth round therefore consists in the elimination of this cycle.

There is no new circle and the patient t_{12} now points to k_{10} forming a chain of maximum length $W_3 = (k_{12}, t_{12}, k_{10}, t_{10}, k_1, t_1, k_9, t_9)$. Round 5 assigns kidneys k_{10} to patient t_{12} and there are no more patients with no kidneys. The algorithm stops.

The final matching is:

$$\begin{pmatrix} t_1 & t_2 & t_3 & t_4 & t_5 & t_6 & t_7 & t_8 & t_9 & t_{10} & t_{11} & t_{12} \\ k_9 & k_{11} & k_2 & k_8 & k_7 & k_5 & k_6 & k_4 & w & k_1 & k_3 & k_{10} \end{pmatrix}$$

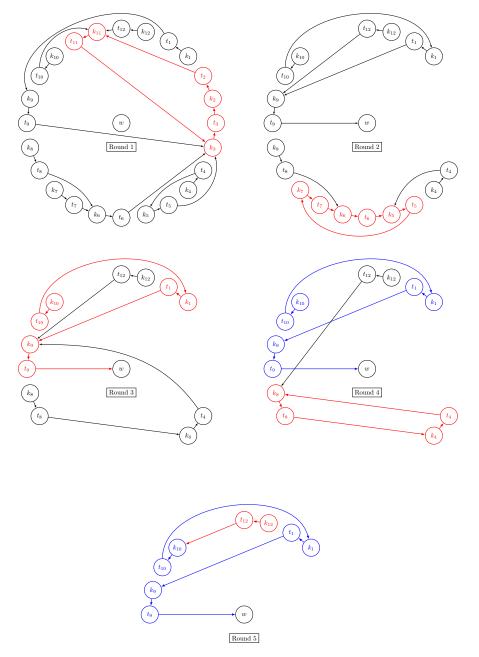


Figure 1: State of the graph at every round.

Question 6

If the selection rule keep in the procedure any selected w-chain at a non-terminal round, then every unasigned kidney is available at every round of the procedure. Therefore, at every round of the algorithm, a patient can only be assigned to his best choice among remaining kidneys. The only way to give him a better kidney is to take it back from a patient whose assignement occurs in a previous round. Since this patient had his best choice at the time, he will strictly disprefer the new matching. Thus, the exchange mechanism is efficient.

Question 7

If we swap k_8 and k_9 in the order of preference of the twelfth patient, then in round 3 the longest w-chain becomes $(k_{12}, t_{12}, k_8, t_8, k_4, t_4, k_9, t_9)$ and t_{12} is assigned to k_8 , which he strictly prefers to k_{10} , that he would get by being honest about his preferences. Therefore the exchange mechanism with chain selection rule A is not strategy-proof.

Question 8

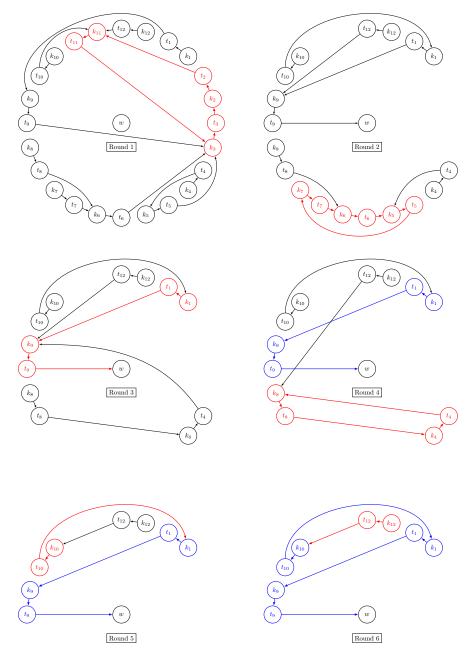


Figure 2: State of the graph at every round.