

**Last name(s)**

**Name**

**ID**

**Final EDA Exam**

**Length: 3 hours**

**12/01/2017**

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- *The exam has 4 sheets, 8 sides and 4 problems.*
  - *Write your full name and ID on every sheet.*
  - *Write your answers within the reserved space in the exam sheet.*
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**Problem 1**

**(2 pts.)**

Fill the following gaps as precisely as possible:

- (a) (0.25 pts.) A graph with  $n$  vertices has  $O(\text{ } \text{ } )$  edges.
- (b) (0.25 pts.) A connected graph with  $n$  vertices has  $\Omega(\text{ } \text{ } )$  edges.
- (c) (0.25 pts.) A complete graph with  $n$  vertices has  $\Omega(\text{ } \text{ } )$  edges.
- (d) (0.25 pts.) A min-heap with  $n$  vertices has  $\Theta(\text{ } \text{ } )$  leaves.
- (e) (0.25 pts.) A binary search tree with  $n$  vertices has height  $\Omega(\text{ } \text{ } )$ .
- (f) (0.25 pts.) A binary search tree with  $n$  vertices has height  $O(\text{ } \text{ } )$ .
- (g) (0.25 pts.) An AVL tree with  $n$  vertices has height  $\Omega(\text{ } \text{ } )$ .
- (h) (0.25 pts.) An AVL tree with  $n$  vertices has height  $O(\text{ } \text{ } )$ .

*This side would be intentionally blank if it were not for this note.*

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**Problem 2**

**(3 pts.)**

Let  $G = (V, E)$  be a directed graph with weights  $\omega : E \rightarrow \mathbb{R}$ . We want to solve the problem of, given a vertex  $s \in V$ , to compute the distance from  $s$  to all vertices of the graph.

Recall that:

- a *path* is a sequence of vertices that are connected consecutively by arcs; that is to say,  $(u_0, u_1, \dots, u_k)$  such that for all  $1 \leq i \leq k$ , we have  $(u_{i-1}, u_i) \in E$ .
- the *weight* of a path is the sum of the weights of its arcs:

$$\omega(u_0, u_1, \dots, u_k) = \sum_{i=1}^k \omega(u_{i-1}, u_i).$$

- the *distance* of a vertex  $u$  to a vertex  $v$  is the weight of the path (called *minimum path*) with minimum weight among those leaving from  $u$  and arriving at  $v$ , if it exists.
- (a) (0.2 pts.) Let us assume that for any edge  $e \in E$ , we have  $\omega(e) = 1$ ; that is to say, all weights are 1. Which algorithm can we use to solve efficiently the problem of the distances?
- 
- (b) (0.2 pts.) Let us assume that for any edge  $e \in E$ , we have  $\omega(e) \geq 0$ ; that is to say, all weights are non-negative. Which algorithm can we use to solve efficiently the problem of the distances?
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- (c) (0.2 pts.) Which algorithm can we use to solve efficiently the problem of the distances, if some weights can be negative?
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- (d) (0.4 pts.) Which condition over the cycles of the graph must be satisfied so that, for all pairs of vertices  $u, v \in V$ , there exists a minimum path from  $u$  to  $v$ ?
-

- (e) (1 pt.) A function  $\pi : V \rightarrow \mathbb{R}$  is a *potential* of the graph if it satisfies that, for any edge  $(u, v) \in E$ , we have  $\pi(u) - \pi(v) \leq \omega(u, v)$ . Moreover, the *reduced weights*  $\omega_\pi$  are defined as  $\omega_\pi(u, v) = \omega(u, v) - \pi(u) + \pi(v)$  for any  $(u, v) \in E$ .

Prove that if  $c$  is a path from  $u$  to  $v$  then  $\omega_\pi(c) = \omega(c) - \pi(u) + \pi(v)$ .

- (f) (1 pt.) Suppose that the graph has a potential  $\pi$ . Then, it can be proved that for all pairs of vertices  $u, v \in V$  there is a minimum path with weights  $\omega$  from  $u$  to  $v$ . Assuming this fact, explain how to use the potential  $\pi$  to compute the distance from a given vertex  $s$  to all vertices of the graph with an alternative algorithm to that of part (c) when the weights can be negative.

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**Problem 3**

**(2 pts.)**

We define a type *matrix* to represent square matrices of real numbers. Consider the following program:

```
typedef vector<vector<double>> matrix;

matrix aux(const matrix& A, const matrix& B) {
    int n = A.size ();
    matrix C(n, vector<double>(n, 0));
    for (int i = 0; i < n; ++i)
        for (int j = 0; j < n; ++j)
            for (int k = 0; k < n; ++k)
                C[i][j] += A[i][k] * B[k][j];
    return C;
}

matrix mystery(const matrix& M) {
    int n = M.size ();
    matrix P(M), Q(M);
    for (int i = 1; i < n; ++i) {
        P = aux(P, M);
        Q = aux(Q, P);
    }
    return Q;
}
```

- (a) (0.5 pts.) What does the function *matrix mystery*(const *matrix*& *M*) compute in terms of the matrix *M*?

- (b) (0.5 pts.) If *M* is an  $n \times n$  matrix, what is the cost in the worst case of the function *matrix mystery*(const *matrix*& *M*) as a function of *n*? Justify your answer.

- (c) (1 pt.) Implement in C++ a function that computes the same as the function *matrix\_mystery*(**const** *matrix*& *M*) and which takes  $\Theta(n^3 \log n)$  time in the worst case. Also implement the auxiliary functions you use, except for *aux* and the functions of the standard library of C++. Justify that the cost of your function is as required.

**Last name(s)****Name****ID****Problem 4****(3 pts.)**

The problem of HAMILTONIAN GRAPH consists in, given an (undirected) graph, to decide whether it is Hamiltonian, that is to say, whether there exists a cycle that visits all its vertices exactly once. It is well-known that HAMILTONIAN GRAPH is an NP-complete problem.

It is also known that, from the proof that a problem belongs to class NP, one can derive a brute force algorithm that solves it. The following function **bool ham(const vector<vector<int>>& G)** which, given a graph G represented with adjacency lists, returns if G is Hamiltonian, implements this algorithm in the case of HAMILTONIAN GRAPH.

```

1  bool ham_rec(const vector<vector<int>>& G, int k, vector<int>& p) {
2      int n = G.size ();
3      if (k == n) {
4          vector<bool> mkd(n, false);
5          for (int u : p) {
6              if (mkd[u]) return false;
7              mkd[u] = true;
8          }
9          for (int k = 0; k < n; ++k) {
10             int u = p[k];
11             int v;
12             if (k < n-1) v = p[k+1];
13             else v = p[0];
14             if (find(G[u].begin(), G[u].end(), v) == G[u].end()) return false;
15         }
16         return true;
17     }
18     for (int v = 0; v < n; ++v) {
19         p[k] = v;
20         if (ham_rec(G, k+1, p)) return true;
21     }
22     return false;
23 }
24
25 bool ham(const vector<vector<int>>& G) {
26     int n = G.size ();
27     vector<int> p(n);
28     return ham_rec(G, 0, p);
29 }
```

Note: The function of the STL library

*Iterator find ( Iterator first , Iterator last , **int** val );*

returns an iterator to the first element in the range *[first, last)* that compares equal to *val*. If no such element exists, the function returns last.

- (a) (0.5 pts.) Identify the variable in the previous program which represents the witness when the function *ham* returns **true**.

- (b) (0.5 pts.) Identify the code corresponding to the verifier in the previous program. To that end, use the line numbers on the left margin.

- (c) (1 pt.) Fill the following gaps so that the function *ham2* computes the same as the function *ham*, but more efficiently.

```

bool ham2_rec(const vector<vector<int>>& G, int k, int u, vector<int>& next) {
    int n = G.size ();
    if (  == n)
        return find (G[u].begin (), G[u].end (),  )  $\neq$  G[u].end();

    for (int v : G[u])
        if (next[v] ==  ) {
            next[u] =  ;
            if (ham2_rec(G, k+1, v, next)) return true;
            next[u] = -1;
        }
    return false;
}

bool ham2(const vector<vector<int>>& G) {
    int n = G.size ();
    vector<int> next(n, -1);
    return ham2_rec(G,  , 0, next);
}

```

- (d) (0.5 pts.) Suppose that the adjacency lists of the representation of *G* are sorted (for example, increasingly). Explain how to use this to make the function of part (c) more efficient.

- (e) (0.5 pts.) Suppose that *G* is a **disconnected** graph. Explain how to use this to make the function of part (c) more efficient.