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#### Greedily improving our own closeness centrality in a network

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The closeness centrality is a well-known measure of importance of a vertex within a given complex network. Having high closeness centrality can have positive impact on the vertex itself: hence, in this paper we consider the optimisation problem of determining how much a vertex can increase its centrality by creating a limited amount of new edges incident to it. We will consider both the undirected and the directed graph case. In both cases, we first prove that the optimisation problem does not admit a polynomial-time approximation scheme (unless P = NP), and we then propose a greedy approximation algorithm (with an almost tight approximation ratio), whose performance is then tested on synthetic graphs and real-world networks.

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#### 1. INTRODUCTION

Looking for the most important vertices within a given complex network has always been one of the main goals in the field of real-world network analysis. Different measures of importance have been introduced in the literature, and several of them are related to the notion of "centrality" of a vertex. This latter notion, in turn, has been explicitly formalized in different ways: one of the most popular is the closeness centrality measure (see, for example, [Boldi and Vigna 2014]). This measure somehow evaluates the efficiency of a vertex while spreading information to all other vertices in its connected component: more formally, the closeness centrality of u is equal to the sum of the reciprocal of the distances to u from all other vertices. Computing closeness centrality, however, is too time expensive, since it requires to run a breadth first search for each vertex, which is clearly infeasible for networks with millions of vertices and edges (which is the "normal" size of many interesting real-world networks). For this reason, several randomized and/or approximation algorithms have been proposed for the computation of this centrality measure [Cohen et al. 2014].

In this paper, instead, we consider a different problem related to the closeness centrality, that is, the problem of identifying which "strategy" a vertex should adopt in order to increase its own centrality value. Indeed, increasing its own ranking in terms of centrality,

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can have positive consequences for the vertex. For example, in the field of author citation networks closeness centrality seems to be significantly correlated with citation counts (as it has been already observed in the case of collaboration networks) [Yan and Ding 2009]. We then consider the optimization problem of efficiently determining, for a given vertex u, the set of k edges incident to u that, when added to the original graph, allows u to increase as much as possible its closeness centrality and its ranking according to this measure. We will analyze both the undirected and the directed graph case. We first prove that this problem is hard to be approximated within an approximation factor greater than  $1 - \frac{1}{15e}$ in the undirected case (respectively,  $1 - \frac{1}{3e}$  in the directed case), and we then show that a greedy approach yields a  $(1 - \frac{1}{e})$ -approximation algorithm in both undirected and directed cases. Successively, we present several experiments that we have performed (i) in order to evaluate how good is the approximation factor in the case of relatively small randomly generated graphs, (ii) in order to apply the greedy approach to real-world collaboration, citation and transportation networks, and (iii) in order to evaluate the actual improvement in information spread. As a result of the first set of experiments, we have that the greedy algorithm seems to perform much better than the theoretical results, since it often computes an optimal solution and, in any case, it achieves an approximation factor significantly larger than the theoretical one. By applying the greedy algorithm to real-world networks, instead, we observe that by adding very few edges a vertex can drastically increase its centrality measure and, hence, its ranking. We note that, after adding a limited number of edges, the number of informed nodes in the network highly increases.

The problem of adding edges to a graph in order to modify some general properties has been widely studied. To the best of our knowledge, the problems that aim at optimizing some property by adding a limited number of edges are: minimizing the average shortestpath distance between all pair of nodes [Meyerson and Tagiku 2009; Papagelis et al. 2011; Parotsidis et al. 2015], minimizing the average number of hops in shortest paths of weighted graphs [Bauer et al. 2012], maximizing the leading eigenvalue of the adjacency matrix [Saha et al. 2015; Tong et al. 2012], minimizing the diameter [Bilò et al. 2012; Frati et al. 2015], maximizing or minimizing the number of triangles [Dehghani et al. 2015; Li and Yu 2015], minimizing the eccentricity [Perumal et al. 2013], and minimizing the characteristic path length [Papagelis 2015].

The problem analyzed in this paper differs from above mentioned ones as it focus on improving the centrality of a predefined vertex. As far as we know, our problem has never been attacked before, even though similar problems have been studied for other centrality measures, i.e. page-rank [Avrachenkov and Litvak 2006; Olsen and Viglas 2014], eccentricity [Demaine and Zadimoghaddam 2010], average distance [Meyerson and Tagiku 2009], some measures related to the number of paths passing through a given node [Ishakian et al. 2012], and betweenness centrality [Crescenzi et al. 2015; D'Angelo et al. 2015]. Hence, we had no other algorithms to compare with. However, we also consider other potential alternative algorithms and show that the greedy algorithm significantly outperforms them, whenever k > 1.

#### 1.1. Motivating applications

In this section we motivate our study by showing two applications in which improving the centrality of a specific node by adding edges incident to it can give benefits to the node itself or to the whole network.

1.1.1. Increasing the spreading of information. Intuitively closeness centrality evaluates the efficiency of a vertex while spreading information to all other vertices in its connected component. We show that solving our problem for a set of given vertices has positive consequences for the spreading of information through the network. To this aim, we consider the *Linear Threshold Model* which is a widely studied model in network analysis to represent the

spread of information [Kempe et al. 2015]. In this model, we can distinguish between *active* nodes (which spread the information) and inactive ones. The idea is that a node becomes active if a large part of its neighbors are active. In detail, each node u has a threshold a chosen uniformly at random in the interval [0, 1]. The threshold represents the fraction of neighbors of u that must become active in order for u to become active to let the information diffusion process start, this nodes are called *seeds*. In subsequent steps of the process a node becomes active if the fraction of its active neighbors is greater than its threshold.

In our experimental study, we show that adding a small number of edges incident to some randomly-chosen seeds highly increases the spreading of information in terms of number of nodes that become active. Note that this represents an improvement of the whole network in terms of the efficiency of propagating information. We performed such experiments on both undirected and directed networks.

1.1.2. Link recommendation. The link recommendation task consists in suggesting potential connections to social network users with the aim of increasing their social circle. Link recommendations improve the user experience and at the same time help to increase the connectivity inside the network and speed-up the network growth. Most of the existing link recommendation methods focus on estimating the likelihood that a link is adopted by users and recommend links that are likely to be established [Backstrom and Leskovec 2011; Liben-Nowell and Kleinberg 2003; Popescul and Ungar 2003; Yin et al. 2010].

Recently, a new approach has been proposed whose aim is to recommend a set of links that, when added to the network, increases the centrality of a user in a network. In particular, suggesting links that minimize the expected average distance of a node accurately predicts the links that will actually appear in the graph [Parotsidis et al. 2016]. An important step in this approach is to determine the set of links that, when added to the network, maximizes the specific centrality measure considered.<sup>1</sup>

Of particular interest in this context are the collaboration networks in which nodes represent users and links represent collaboration between users (e.g. authors collaborating in the same papers or actors that acted in the same movie). The link recommendation problem in such a case consists in suggesting possible persons to whom request for future collaboration. In our experiments we show that we are able to compute a set of nodes that highly increases the closeness centrality in very large collaboration networks such as those induced by the DBLP and IMDB databases [Ley ; IMDB ].

#### 2. THE UNDIRECTED GRAPH CASE

In this section we will focus on undirected graphs. After giving all necessary definitions and preliminary results, we will introduce the optimisation problem that will be considered, we will prove a non-approximability result, and we will then describe an approximation algorithm. Finally, we will present the experiments that we have performed in order to validate this algorithm and to apply it to two quite big collaboration networks.

#### 2.1. The maximum closeness improvement problem

Let G = (V, E) be an undirected graph, where V denotes the set of nodes, and E denotes the set of edges  $\{u, v\}$  with  $u, v \in V$ . For each node  $u, N_u$  denotes the set of neighbors of u, i.e.  $N_u = \{v \mid \{u, v\} \in E\}$ . Given two vertices u and v, we denote by  $d_{uv}$  the distance from u to v in G, that is, the number of edges in a shortest path from u to v (if there is no path from u to v, we then set  $d_{uv} = \infty$ ). For each node u, the closeness centrality (also

<sup>&</sup>lt;sup>1</sup>Note that the centrality measure used in [Parotsidis et al. 2016] is the inverse of the arithmetic mean of the distances to a node, while in this paper we consider the harmonic mean of the distances to a node.

called harmonic centrality [Boldi and Vigna 2014]) of u is defined as follows

$$c_u = \sum_{\substack{v \in V \\ d_{uv} < \infty}} \frac{1}{d_{uv}}.$$

Given a set S of edges not in E, we denote by G(S) the graph augmented by adding the edges in S to G, i.e.  $G(S) = (V, E \cup S)$ . For a parameter x of G, we denote by x(S) the same parameter computed in the augmented graph G(S) (for example, the distance from u to v in G(S) is denoted as  $d_{uv}(S)$ ).

The closeness centrality of a vertex clearly depends on the graph structure: if we augment a graph by adding a set of edges S, then the centrality of a vertex might change. Generally speaking, adding edges incident to some vertex u can only increase the centrality of u. Given a graph G = (V, E), a vertex  $u \in V$ , and an integer k, the Maximum Closeness Improvement (in short, MCI) problem consists in finding a set S of edges incident to u not in E (that is,  $S \subseteq \{\{u, v\} : v \in V \setminus N_u\}$ ) such that  $|S| \leq k$  and  $c_u(S)$  is maximum.

#### 2.2. The non-approximability result

In this section, in order to derive our approximation hardness result for the MCI problem, we will make use of the Minimum Dominating Set (in short, MDS) problem, which is defined as follows: given an undirected graph G = (V, E), find a *dominating set* of minimum cardinality, that is, a subset D of V such that  $V = D \cup \bigcup_{u \in D} N_u$ . It is known that, for any r with 0 < r < 1, it cannot exist a  $(r \ln |V|)$ -approximation algorithm for the MDS problem, unless P = NP [Dinur and Steurer 2014]. We will now use this result in order to show that the MCI problem does not admit a polynomial-time approximation scheme. To this aim, we will design an algorithm A' that, given an undirected graph G = (V, E) and given the size k of the optimal dominating set of G, by using an approximation algorithm A for the MCI problem will return a dominating set of G whose approximation ratio is at most  $(r \ln |V|)$ . Clearly, we do not know the value of k, but we know that this value must be at least 1 and at most |V|: hence, we run algorithm A' for each possible value of k, and return the smallest dominating set found. Algorithm A' will run the approximation algorithm A for the MCI problem multiple times. Each time A will find k nodes  $u \in V$ which are the "new" neighbours of the node whose centrality has to be increased: we then add these nodes to the dominating set and create a smaller instance of the MCI problem (which will contain, among the others, all the nodes in V not yet dominated). We continue until all nodes in V are dominated.

THEOREM 2.1. For each  $\gamma > 1 - \frac{1}{15e}$ , there is no  $\gamma$ -approximation algorithm for the MCI problem, unless P = NP.

PROOF. We will show that a  $\gamma$ -approximation algorithm A for the MCI problem, with  $\gamma > 1 - \frac{1}{15e}$ , would imply a  $(r \ln n)$ -approximation algorithm A' for the MDS problem, thus proving the theorem. In particular, the algorithm A' is specified in Fig. 1, where k denotes a "guess" of the size of an optimal solution for MDS with input the graph G. In the following,  $\omega$  will denote the number of times the **while** loop is executed. Since, at each iteration of the loop, we include in the dominating set at most k nodes, at the end of the execution of algorithm A' the set D includes at most  $k \cdot \omega$  nodes. Hence, if k is the correct guess of the value of the optimal solution for the MDS instance, then D is a  $\omega$ -approximate solution for the MDS problem (as we have already noticed, we don't know the correct value of k, but algorithm A' can be executed for any possible value of k, that is, for each  $k \in [|V|]$ ).

The first instruction of the **while** loop of algorithm A' computes a transformed graph G' (to be used as part of the new instance for MCI) starting from the current graph  $G = (V, E_V)$ , which is the subgraph of the original graph induced by the set  $\{u_1, \ldots, u_n\}$ ,

Algorithm: A'**Input** : an undirected graph G = (V, E) and an integer k **Output**: a dominating set D

1  $D := \emptyset$ :

while  $V \neq \emptyset$  do  $\mathbf{2}$ 

Compute graph G' starting from G (see Fig. 2); 3

- S := A(G', z, k);4
- $D' := \{u : \{z, u\} \in S\}$ 5
- 6
- $$\begin{split} D &:= D \cup D'; \\ V &:= V D' \bigcup_{u \in D'} N_u; \end{split}$$
  7
- G := subgraph of  $\widetilde{G}$  induced by V; 8

9 return D;

Fig. 1. The approximation algorithm for the MDS problem, given a  $\gamma$ -approximation algorithm A for the MCI problem and a "guess" k for the optimal value of MDS.



Fig. 2. The reduction used in Theorem 2.1. The dashed edges denote those added in a solution to MCI.

where n = |V|, of still not dominated nodes. This computation is done as follows (see Fig. 2). We add a new node z and two new nodes  $x_i$  and  $y_i$ , for each i with  $1 \le i \le n$ . Moreover, we add to  $E_V$  the edges  $\{z, y_i\}, \{x_i, y_i\}, and \{x_i, u_i\}, for each i with <math>1 \le i \le n$ . As it is shown in the second line of the **while** loop, z is the node whose centrality  $c_z$  has to be increased by adding at most k edges: that is, the MCI instance is formed by G', z, and k. Observe that any solution for this instance that contains an edge  $\{x_i, z\}$  can be modified, without decreasing its measure, by substituting this edge with  $\{u_i, z\}$ : hence, we can assume that the solution S computed at the second line of the while loop of algorithm A' contains only edges connecting z to nodes in V (which are shown by dashed lines in Fig. 2).

First of all, note that, since k is (a guess of) the measure of an optimal solution  $D^*$  for MDS with input G, we have that the measure  $c^*(G', z, k)$  of an optimal solution  $S^*$  for MCI with input G' satisfies the following inequality:

$$c^*(G', z, k) \ge k + \frac{1}{2}(n-k) + \frac{3}{2}n = \frac{1}{2}k + 2n.$$

This is due to the fact that, by connecting z to all the k nodes in  $D^*$ , in the worst case we have that k nodes in G are at distance 1, n-k nodes in G are at distance 2 (since  $D^*$  is a dominating set), the n nodes  $y_i$  are at distance 1, and the n nodes  $x_i$  are at distance 2 from z.

Given the solution S computed by the approximation algorithm A for MCI, let a and bdenote the number of nodes in G at distance 2 and 3, respectively, from z in G'(S). Since all nodes in G' are at distance at most 3 from z, we have that n = k + a + b (we can assume,

without loss of generality, that  $n \ge k$ ): hence, a = n - b - k. Since A is a  $\gamma$ -approximation algorithm for MCI, we have that  $c_z(S) \ge \gamma c^*(G', z, k)$ . That is,

$$k + \frac{1}{2}a + \frac{1}{3}b + \frac{3}{2}n \ge \gamma\left(\frac{1}{2}k + 2n\right).$$

From this inequality, it follows that

$$a \ge \gamma(k+4n) - 3n - 2k - \frac{2}{3}b.$$

By using the fact that a = n - b - k, we have that

$$n-b-k \ge \gamma(k+4n) - 3n - 2k - \frac{2}{3}b.$$

That is,

$$b \le 12(1-\gamma)n + 3(1-\gamma)k.$$

Since  $k \leq n$ , we then have that

$$b \le 15n(1-\gamma).$$

Assuming  $\gamma > 1 - \frac{1}{15e} > \frac{14}{15}$  (which implies  $15(1 - \gamma) < 1$ ), then after one iteration of the **while** loop of algorithm A', the number of nodes in G decreases by a factor  $15(1 - \gamma)$ . Hence, after  $\omega - 1$  iterations, the number n of nodes in the graph G is at most a fraction  $[15(1 - \gamma)]^{\omega - 1}$  of the number N of nodes in the original graph. Since we can stop as soon as n < k, we need to find the maximum value of  $\omega$  such that  $k \le N[15(1 - \gamma)]^{\omega - 1}$ . By solving this inequality and by recalling that  $15(1 - \gamma) < 1$ , we obtain

$$\omega - 1 \le \log_{15(1-\gamma)} \frac{k}{N} \le \log_{15(1-\gamma)} \frac{1}{N} = \frac{\ln(N)}{\ln \frac{1}{15(1-\gamma)}}.$$

One more iteration might be necessary to trivially deal with the remaining nodes, which are less than k. Hence, the total number  $\omega$  of iterations is at most  $\frac{\ln(N)}{\ln \frac{1}{15(1-\gamma)}} + 1$ . If  $\gamma > 1 - \frac{1}{15e}$ , we have that  $r' = \frac{1}{\ln \frac{1}{15(1-\gamma)}} < 1$ : as a consequence of the observation at the beginning of the proof, the solution reported by algorithm A' is an  $(r' \ln N + 1)$ -approximate solution. Clearly, for any r with 0 < r' < r < 1, there exists  $N_r$  sufficiently large, such that for any  $N > N_r$ ,  $r' \ln N + 1 \leq r \ln N$ : hence, algorithm A' would be an  $r \ln N$ -approximation algorithm for MDS, and, because of the result of [Dinur and Steurer 2014], P would be equal to NP. Thus, we have that, if  $P \neq NP$ , then  $\gamma$  has to be not greater than  $1 - \frac{1}{15e}$  and the theorem is proved.  $\Box$ 

#### 2.3. The greedy approximation algorithm

Let us consider the following optimisation problem. Given a set X and an integer k, find a subset Y of X of cardinality at most k that maximises the value f(Y), where  $f: 2^X \to \mathbb{N}$  is a specific objective function. If f is monotone submodular, that is, if, for any pair of sets  $S \subseteq T \subseteq X$  and for any element  $e \in X \setminus T$ ,  $f(S \cup \{e\}) - f(S) \ge f(T \cup \{e\}) - f(T)$ , then the following greedy algorithm approximates the above problem within a factor  $1 - \frac{1}{e}$  [Nemhauser et al. 1978]: start with the empty set, and repeatedly add an element that gives the maximal marginal gain. In this section, we exploit this result by showing that  $c_u$  is monotone and submodular with respect to the possible set of edges incident to u. Hence, the greedy algorithm reported in Fig. 3 provides a  $(1 - \frac{1}{e})$ -approximation. Note that the computational complexity of such algorithm is  $O(k \cdot n \cdot g(n, m + k))$ , where g(n, m + k) is the complexity of computing  $c_u$  in a graph with n nodes and m + k edges.

Algorithm: GREEDYIMPROVEMENT **Input** : an undirected graph G = (V, E); a vertex  $u \in V$ ; and an integer  $k \in \mathbb{N}$ **Output**: set of edges  $S \subseteq \{\{u, v\} \mid v \in V \setminus N_u\}$  such that  $|S| \leq k$ 1  $S := \emptyset;$ **2** for i = 1, 2, ..., k do foreach  $v \in V \setminus N_u(S)$  do 3 Compute  $c_u(S \cup \{u, v\})$ ;  $\mathbf{4}$  $v_{\max} := \arg\max\{c_u(S \cup \{u, v\}) \mid v \in V \setminus N_u(S)\};$  $S := S \cup \{\{u, v_{\max}\}\};$ 5 6 7 return S: Fig. 3. The greedy algorithm for undirected graphs.

THEOREM 2.2. For each vertex u, function  $c_{\mu}$  is monotone and submodular with respect to any feasible solution for MCI.

PROOF. To simplify the notation, in the following we will assume that  $\frac{1}{\infty} = 0$ . To show that  $c_u$  is monotone increasing, it is enough to observe that, for each solution S to MCI, for each edge  $\{u, v\} \notin E \cup S$ , and for each node  $x \in V \setminus \{u\}, d_{ux}(S \cup \{\{u, v\}\}) \leq d_{ux}(S)$ (since adding an edge cannot increase the distance between two nodes) and, therefore,  $\frac{1}{d_{ux}(S \cup \{\{u,v\}\})} \ge \frac{1}{d_{ux}(S)}$ . We now show that, for each pair S and T of solutions for MCI such that  $S \subseteq T$ , and for each edge  $\{u, v\} \notin T \cup E$ ,

$$c_u(S \cup \{\{u, v\}\}) - c_u(S) \ge c_u(T \cup \{\{u, v\}\}) - c_u(T)$$

To this aim, we prove that each term of  $c_u$  is submodular, that is, that, for each vertex  $x \in V \setminus \{u\},\$ 

$$\frac{1}{d_{ux}(S \cup \{\{u, v\}\})} - \frac{1}{d_{ux}(S)} \ge \frac{1}{d_{ux}(T \cup \{\{u, v\}\})} - \frac{1}{d_{ux}(T)}.$$
(1)

Let us consider the shortest paths from u to x in  $G(T \cup \{\{u, v\}\})$ , and let us distinguish the following two cases.

- (1) The first edge of a shortest path from u to x in  $G(T \cup \{\{u, v\}\})$  is  $\{u, v\}$  or be-(1) The first edge of a shortest path from u to x in G(T ∪ {{u,v}}) is {u,v} or belongs to S ∪ E. In this case, such a path is a shortest path also in G(S ∪ {{u,v}}), as it cannot contain edges in T \ S (since these edges are all incident to u). Then, d<sub>ux</sub>(S ∪ {{u,v}}) = d<sub>ux</sub>(T ∪ {{u,v}}) and 1/(d<sub>ux</sub>(S ∪ {{u,v}})) = 1/(d<sub>ux</sub>(T ∪ {{u,v}})). Moreover, d<sub>ux</sub>(S) ≥ d<sub>ux</sub>(T) (since S ⊆ T) and, therefore, -1/(d<sub>ux</sub>(S) ≥ -1/(d<sub>ux</sub>(T)).
   (2) The first edge of all shortest paths from u to x in G(T ∪ {{u,v}}) belongs to T \ S. In this case, d<sub>ux</sub>(T) = d<sub>ux</sub>(T ∪ {{u,v}}) and, therefore, 1/(d<sub>ux</sub>(T ∪ {{u,v}})) = -1/(d<sub>ux</sub>(T)) = 0. As 1/(d<sub>ux</sub>(S)) is monotone increasing, then 1/(d<sub>ux</sub>(S ∪ {{u,v}})) - 1/(d<sub>ux</sub>(S)) ≥ 0.

In both cases, we have that the inequality (1) is satisfied and, hence, the theorem follows.  $\Box$ 

COROLLARY 2.3. The MCI problem is approximable within a factor  $\left(1-\frac{1}{e}\right)$ .

As it can be seen, there is quite a significant gap between the non-approximability result proved in Theorem 2.1 (that is, the upper bound equal to  $1 - \frac{1}{15e} \approx 0.98$ ), and the approximability result of the above corollary (that is, the lower bound  $(1 - \frac{1}{e}) \approx 0.63$ ). One of the main goals of the next experimental session is to analyse the "real" performance, in terms of solution quality, of the greedy algorithm on relatively small real-world and synthetic graphs.

Table I. Comparison between the GREEDYIMPROVEMENT algorithm and the optimum in random graphs. The first three columns reports the type and size of the graphs; the fourth column reports the minimum measured approximation ratio.

Network	n =  V	m =  E	Min Approx.
PA	100	130	0.9939
PA	500	650	0.9921
ER	100	200	0.9828
ER	100	500	0.9938
ER	100	1000	0.9970
ER	500	5000	0.9971
ER	500	12500	0.9991
ER	500	25000	1
CM	100	200	0.9946
CM	500	1000	0.9995
WS	100	500	0.9798
WS	100	600	0.9798
WS	100	800	0.9856
WS	100	1200	0.9946

Table II. Comparison between the GREEDYIMPROVEMENT algorithm and the optimum in real world graphs. The first three columns reports the name and size of the graphs; the fourth column reports the minimum measured approximation ratio.

Network	n =  V	m =  E	Min Approx.
s838_ st	512	819	0.9862
jazz	198	2742	0.9968
coli1	328	456	0.9947
celegans_metabolic	346	1493	0.9981

#### 2.4. The experimental study: part I

In this section we analyse the greedy algorithm from an experimental point of view. First, we compare the solution of the greedy algorithm with the optimal solution computed by using an integer program formulation of the MCI problem, in order to assess its real performance in terms of solution quality. Then, we compare the greedy algorithm with several alternative baselines. Finally, we study how the spreading of information increases as a consequence of the augmentation of the graph due to our algorithm.

All our experiments have been performed on a computer equipped with two Intel Xeon E5-2643 CPUs, each with 6 cores clocked at 3.4GHz and 128GB of main memory, and our programs have been implemented in C++ (gcc compiler v4.8.2 with optimization level O3).

2.4.1. Evaluating the solution quality. In this section we evaluate the quality of the solution produced by the greedy algorithm by measuring its approximation ratio on several, relatively small, randomly generated networks and on four real-world networks. In particular, we considered four random graph generating models, that is, undirected Preferential Attachment (in short, PA) [Barabasi and Albert 1999], Erdős-Rényi (in short, ER) [Erdős and Rényi 1959], Configuration Model (in short, CM) [Molloy and Reed 1995; Bender and Canfield 1978], and Watts-Strogatz model (in short WS) [Watts and Strogatz 1998]. The size of the generated graphs is reported in Table I. For each combination (n, m), we generated five random undirected graphs. Moreover, we considered the four real-world graphs, whose size is reported in Table II. The first graph is the collaboration network between Jazz musicians that have played together in a band, and it has been obtained from the Konect database [Kunegis 2013], while the last three graphs have been downloaded from the Uri AlonLab [Uri AlonLab ] database: in particular, s838\_st is an electronic network, while the other two graphs are biological networks.

For both random and real-world graphs we focused our attention on twenty vertices u, which have been chosen on the basis of their original closeness ranking. In particular, we have divided the list of vertices, sorted by their original ranking, in four intervals, and chosen five random vertices uniformly at random in each interval: we denote by  $u_{X\%}$  the average value of the vertices in the interval of the top Xth percentile. The value of k ranged from 1 to 10. In the experiments, we measured the ratio between the value of the solution found by the greedy algorithm and the optimal value computed by using the integer program formulation of the MCI problem, defined as follows.

The decision variables  $x_v$  and  $y_{sv}$  specify a solution S of the MCI problem as follows. For any  $v \in V \setminus N_u$ ,

$$x_v = \begin{cases} 1 & \text{if } \{u, v\} \in S, \\ 0 & \text{otherwise,} \end{cases}$$

and, for each  $s \in V \setminus \{u\}$  and  $v \in V \setminus N_u$ ,

 $y_{sv} = \begin{cases} 1 & \text{if a shortest path from } s \text{ to } u \text{ in } G(\{u, v\}) \text{ passes through edge } \{u, v\}, \\ 0 & \text{otherwise.} \end{cases}$ 

The first constraint of the integer program ensures that each node s can be covered by at most one edge  $\{u, v\}$  and, hence, that the distance from s to u is counted only once in the objective function, while the second constraint ensures that if  $y_{sv} = 1$ , then  $x_v = 1$ and, hence, that the shortest path from s to u passing through  $\{u, v\}$  is considered only if  $\{u, v\} \in S$ . Finally, note that in the objective function, the value of  $\frac{1}{d_{su}(\{u,v\})}$  and  $\frac{1}{d_{su}}$  can be preprocessed, and that the term  $\sum_{s \in V \setminus \{u\}} \frac{1}{d_{su}}$  is a constant.

We solved the above integer program by using the GLPK solver [glp]. The results are reported in Table I and in Table II where we show the minimum (i.e. worst-case) approximation ratio obtained by the greedy algorithm. The experiments clearly show that the experimental approximation ratio is by far better than the theoretical one proven in the previous section. In fact, in the worst case the ratio is 0.9798.

In Fig. 4, instead, we plot the average closeness centrality and ranking of vertices u as a function of k in a small real-world network, namely the  $s838\_st$  electrical network. We observe that the charts on the top, where the values are computed using the GREEDY-IMPROVEMENT algorithm, and the charts on the bottom, in which we used the optimal algorithm, are almost identical. Indeed, the approximation ratio in the worst case is 0.9862: that is, the GREEDYIMPROVEMENT algorithm performs very well in practice.

Finally, we tested our algorithm on several artificial instances generated by the Erdős-Rényi and the Watts-Strogatz models. In the former model we can choose appropriate values of the graph density, while in the latter one we can choose the clustering coefficient. It turned out that the performance of our algorithm are not influenced by these two factors.



Fig. 4. Closeness centrality and ranking of vertices in the four intervals u as a function of k in the network s838\_st. Comparison between the GREEDYIMPROVEMENT algorithm and the optimal one.

Indeed, the approximation ratio ranges in [0.9798, 1] and improves a little when the density is very high (i.e.  $m > 0.5n^2$ ).

2.4.2. The comparison with alternative baselines. In this section we compare our algorithm with the following algorithms:

- (1) The algorithm that connects u to a set of k nodes extracted uniformly at random (RANDOM).
- (2) The algorithm that connects u to a set of k nodes having the highest degree (DEGREE).
- (3) The algorithm that connects u to a set of k nodes having the highest harmonic centrality (TOP-K).
- (4) The algorithm that connects u to a set of k nodes that have the highest fractional value when solving the linear relaxation of the integer program (ROUNDING).
- (5) The algorithm that connects u to a set of k nodes computed with an approximation algorithm for the k-median with penalties problem given in [Meyerson and Tagiku 2009] (K-MEDIAN).

The first two algorithms are easy to describe and implement efficiently. In what follows we give more details on the implementation of the last three algorithms.

The TOP-K algorithm. The classical algorithm to find the k nodes having the highest value of centrality, consists, for each node v, in determining all the distances to v by running a Breadth First Search (BFS) and computing  $c_v$ . With such an approach computing the k nodes having the highest value of centrality requires  $O(n \cdot (n + m))$ . In Fig. 6 we give

an algorithm that reduces the computation time by using a branch-and-bound technique that prunes the unnecessary BFS by comparing the intermediate results of centrality with a properly defined upper bound.

Algorithm: PrunedBFS **Input** : An undirected graph G(S); a node v, a double  $min_c$ **Output**:  $c_v$ 1  $Q := \emptyset;$ **2** visited :=  $\emptyset$ ; **3**  $d_{uv}(S \cup \{u, v\}) := 0;$ 4 foreach  $x \in N_v(S)$  do  $d_{ux}(S \cup \{u, v\}) := 1;$  $\mathbf{5}$  $V_x := V_x + 1;$ 6 **7** visited :=  $\{u, v\} \cup N_u(S);$ **8**  $c_v := 0;$ 9 foreach  $x \in N_v(S)$  do | Q.push(x); $\mathbf{10}$ 11 while  $\neg Q.empty()$  do x := Q.pop();12 $V_x := V_x - 1;$  $c_v := \frac{1}{d_{ux}(S)};$  $\mathbf{13}$  $\mathbf{14}$ 15foreach  $y \in N_x(S)$  do  $\begin{array}{l} \text{if } (y \notin visited) \land (d_{uy}(S) > d_{ux}(S \cup \{u, v\} + 1)) \text{ then} \\ \mid d_{uy}(S \cup \{u, v\}) := d_{ux}(S \cup \{u, v\}) + 1; \end{array}$ 16  $\mathbf{17}$ Q.push(y); $\mathbf{18}$  $visited := visited \cup \{y\};$ 19  $V_x := V_x + 1$  $\mathbf{20}$  $UB_v := c_v + |V_x| \cdot \frac{1}{d_{vx}} + (|V| - |visited| - |V_x|) \cdot \frac{1}{d_{vx} + 1};$ if  $UB_v \le min_c$  then  $\mathbf{21}$  $\mathbf{22}$ return 23 24 return  $c_v$ 

Fig. 5. Algorithm PrunedBFS.

**Algorithm**: TOP-K **Input** : an undirected graph G = (V, E); and an integer  $k \in \mathbb{N}$ **Output**: set of nodes  $S \subseteq V$  such that |S| = k

1 S: Min priority queue; 2 Sort V according to according to the node degree; 3 Compute the centrality of the first k nodes in V and compute their centrality; 4 for i = k, k + 1, ..., |V| do 5  $min_c := S.getMin();$ 6  $c_j := PrunedBFS(j, min_c);$ 7 if  $c_j > min_c$  then 8 S.pop();9  $S.push(j, c_j);$ 10 return S

Fig. 6. Algorithm TOP-K.

Let  $C_k$  be the set of k nodes having the highest centrality. We represent  $C_k$  with a minheap in order to find the minimum in constant time. First of all, TOP-K algorithm inserts the k nodes with highest degree in  $C_k$  and computes their centrality. Then, it computes the centrality of other nodes v by performing a BFS starting at each v. The algorithm uses the minimum value of centrality in  $C_k$  as a lower bound and prunes the BFS from v when such lower bound is greater then an upper bound (to be defined later) on  $c_v$ . Such upper bound is computed every time a node is extracted from the BFS queue. If the BFS is completed without any pruning, it removes the minimum from  $C_k$  and it inserts the node v in it.

The upper bound estimates the value of the centrality of a node v. The main idea is that, at each BFS step, when we extract a node x at distance  $d_{vx}$  from v, we can maintain the exact number of nodes that are at distance  $d_{vx}$  and that are not visited yet. Moreover, we can upper bound the distance to any other node. When x is extracted from the queue, let  $V_x$  be the set of nodes at distance  $d_{vx}$  from the source v that are not visited be the set of nodes currently visited during the BFS, and Currc be the value of the centrality at the current step, that is  $Currc = \sum_{y \in visited} \frac{1}{d_{vy}}$ . Then, we have that  $|V_x|$  nodes are at distance  $d_{vx}$  from v, while the remaining  $|V| - |visited| - |V_x|$  nodes are at distance at most  $d_{vx} + 1$  from v. Hence the upper bound is defined as:

$$UB_{v} = Currc + |V_{x}| \cdot \frac{1}{d_{vx}} + (|V| - |visited| - |V_{x}|) \cdot \frac{1}{d_{vx} + 1}.$$

The ROUNDING algorithm. This approach consists in adding the edges which are obtained by rounding to one k variables of the optimal solution to the linear relaxation of the integer program for the MCI problem given in Section 2.4.1. In particular, we connect u to the nodes corresponding to the k highest values in an optimal fractional solution.

The K-MEDIAN algorithm. This approach consists in connecting u with the k nodes which are solution of the k-median with penalties problem [Meyerson and Tagiku 2009]. In this problem, we are given a set of cities and a set of potential facility locations. Each city has a demand that needs to be served by a facility. Each city also has a penalty cost, which we pay if we refuse to serve the city. If we choose to serve a city, we must pay the distance between the city and its assigned facility for each unit demand. Our job is to find a set of kfacilities to open, a set of cities to be served, and an assignment of cities to open facilities such that our total cost is minimized. We implemented the approximation algorithm given in [Meyerson and Tagiku 2009] that is based on local search.

Analysis of the results. In order to compare the solution obtained by the GREEDYIM-PROVEMENT algorithm with that obtained by using the other aforementioned approaches, we run all the algorithms on several real-world networks reported in Tables II and III. We first observe that the rounding and the k-median algorithms cannot be executed on networks having more than few hundred of nodes in reasonable computational time. Therefore, in what follows we first compare GREEDYIMPROVEMENT with DEGREE, RANDOM, and TOP-K on network ca-HepPh, which is a well known collaboration network obtained from the SNAP database [Leskovec and Krevl 2014], and then we compare GREEDYIMPROVEMENT with ROUNDING and K-MEDIAN on the network jazz. The results for the other networks are similar (but for the networks in Table III and the ROUNDING and K-MEDIAN algorithms for which we have no results due to the high computational time).

In Fig. 7 we plot the closeness centrality and the ranking of vertex u as a function of k on network ca-HepPh. We observe that any vertex becomes central by adding just few edges. In Fig. 8, we compare the ranking obtained with the solution given by our algorithm with that obtained with the solution given by the other approaches on the same network.

Table III. Real-world graphs used in the comparison between the GREEDYIMPROVEMENT algorithm and the other baselines. The columns reports the type and size of the graphs.

${f Network}$	n =  V	m =  E
advogato	5272	45903
ca-AstroPh	17903	196972
ca-CondMat	21363	91286
ca-HepPh	11204	117619
ca-HepTh	8638	24806
dip20090126	19928	41202
Newman-Cond_mat_95-99	22015	58578
PGPgiant.compo	10680	24316



Fig. 7. Closeness centrality and ranking of vertex u as a function of k in network ca-HepPh.

In particular, we show the *average relative ranking position* that is:

$$\frac{r_u(S_B) - r_u(S_{GR})}{r_u(S_{GR})}$$

where  $S_{GR}$  and  $S_B$  are the solutions given by our algorithm and one baseline algorithm, respectively, and  $r_u(S)$  denotes the closeness ranking of node u in G(S). The average relative ranking position represents the gain of our algorithm over any other baseline in terms of ranking position. Each curve represents the average relative ranking position in a given interval and the values are expressed in percentage. We observe that the greedy algorithm significantly outperforms RANDOM, DEGREE, and TOP-K, whenever k > 1.

In Fig. 9, we compare, the ranking of node u in the solution given by our algorithm with that given by the ROUNDING and K-MEDIAN. We confirm that our algorithm is by far better than the other approaches. In some cases, ROUNDING gives better solutions in terms of objective function value, however, such cases correspond to the instances in which the fractional solution is integral and therefore is optimal for the problem. Note that computing such a solution requires a long computational time and that we cannot apply such an approach for instances having more than few hundred nodes.

2.4.3. The analysis of information spreading. In this section we analyze how adding a limited number of edges incident to some randomly-chosen seeds highly increases the number of nodes that become active in the threshold model.

In the experiments we have choose a number of seeds that is 2%, 4%, 6%, 8% and 10% of the number of nodes of the graph. The seeds are chosen uniformly at random. We run different experiment where the threshold a is uniform and equal to 0.2, 0.3 and 0.4. We



Fig. 8. Average relative ranking position between the Ranking of the solution obtained by the GREEDYIM-PROVEMENT algorithm and the different baselines in network ca-HepPh.



Fig. 9. Average relative ranking position between the Ranking of the solution obtained by the GREEDYIM-PROVEMENT algorithm and the different baselines in network jazz.

measured the number of nodes that become active at the end of the process in the graphs of Tables II and III.

In Fig. 10 we plot the percentage of active nodes as a function of k for the coli1 undirected network. The value for k = 0 is the percentage of active nodes in the original graph. The plots clearly show that the number of active nodes highly increases even with few edges addition and that the percentage of active nodes tends to 100%. The results for the other networks in Tables II and III are similar.



Fig. 10. Percentage of active nodes in coli1 network as a function of k. Parameter a denotes the threshold and the percentage of seeds is equal to 2%, 4%, 6%, 8%, 10%, respectively.

#### 2.5. Improving the greedy algorithm running time

In this section we show how to improve the running time of GREEDYIMPROVEMENT. This algorithm requires  $O(k \cdot n \cdot g(n, m + k))$  computational time, where g(n, m + k) is the complexity of computing  $c_u$  in a graph with n nodes and m+k edges. The classical algorithm to compute  $c_u$  consists in determining all the distances to u by running a BFS starting from u. Therefore, with such an approach GREEDYIMPROVEMENT requires  $O(k \cdot n \cdot (n + m + k))$  in the worst case. In this section we provide a dynamic algorithm to reduce the time required to compute  $c_u$ .<sup>2</sup> Furthermore, we show how to exploit the submodularity

<sup>&</sup>lt;sup>2</sup>Note that the idea of incrementally updating the closeness centrality as been already explored in the literature [Kas et al. 2013; Sariyüce et al. 2013]. However, in this paper we consider the harmonic mean to compute the closeness centrality instead of the arithmetic mean that is used in other papers. The motivation is that the harmonic mean has been showed to be more robust in the case of undirected disconnected

of  $c_u$  in order to reduce the running time of iterations  $i \ge 2$  of the **for** loop at line 2 of GREEDYIMPROVEMENT.

Let us assume that we add an edge  $\{u, v\} \notin E \cup S$  to graph G(S). The dynamic algorithm aims at computing only the distances between u and any other node that change as a consequence of the addition of edge  $\{u, v\}$  (i.e. nodes w such that  $d_{uw}(S) \neq d_{uw}(S \cup \{u, v\})$ ) and keep the old distances to any other node in the graph. The algorithm is based on the following observation: if we add an edge  $\{u, v\}$  to G(S), then  $d_{uw}(S) \neq d_{uw}(S \cup \{u, v\})$ , for some  $w \in V$ , only if the shortest path between u and w in  $G(S \cup \{u, v\})$  contains edge  $\{u, v\}$ . Therefore, we can determine the nodes that change their distance to u by finding all the shortest paths passing through edge  $\{u, v\}$  in  $G(S \cup \{u, v\})$ . To this aim, the dynamic algorithm executes a BFS starting from node v and prunes the search as soon as a node that does not change its distance to u is extracted from the queue. We report the dynamic algorithm *DynamicBFS* in Figure 11. In detail, Procedure *DynamicBFS* returns the value  $\Delta Clo$  which corresponds to the increment to  $c_{\mu}(S)$  which is obtained by adding edge  $\{u, v\}$ . To compute  $\Delta Clo$ , the algorithm computes the distances between u and any node y such that  $d_{uy}(S) \neq d_{uy}(S \cup \{u, v\})$ . First, it computes the distances of u and its neighbors (lines 3–5) and the initial increment  $\Delta Cl_0$  that is equal to the difference between the reciprocal of the new distance and that of the old distance (line 7). Then, it pushes in queue Q the neighbors of u (lines 8 and 9) and performs the BFS starting from v (lines 10–17). For each extracted node, it updates  $\Delta Clo$  by subtracting the reciprocal of the old distance and adding the new one (line 12). After that, it enqueues a neighbor yof the extracted node x only if the old distance  $d_{uy}(S)$  is greater than the length of the path made of the shortest path from u to x in  $G(S \cup \{u, v\})$  and the edge  $\{x, y\}$  (that is  $d_{ux}(S \cup \{u, v\}) + 1$ , see the test at line 14). Note that this condition is satisfied only if the shortest path between u and y passes through edge  $\{u, v\}$ . The procedure repeats this process until the queue is empty.

We give an example of execution of Algorithm 11 in Figure 12.

In order to analyze the computational complexity of Algorithm 11, let us define as  $\gamma_{uv}(S)$  as the set of nodes that change their distance to u as a consequence of the addition of edge  $\{u, v\}$  to G(S), that is

$$\gamma_{uv}(S) = \{ w \in V \mid d_{xu}(S) \neq d_{xu}(S \cup \{\{u, v\}\}) \}.$$

Moreover, let  $\Gamma_{uv}(S)$  be the number of edges incident to nodes in  $\gamma_{uv}(S)$ , that is  $\Gamma_{uv}(S) = \sum_{w \in \gamma_{uv}(S)} |N(w)|$ . Parameters  $|\gamma_{uv}(S)|$  and  $\Gamma_{uv}(S)$  measure the minimal number of nodes and edges, respectively, that must be visited in order to update all the distance to u after the addition of edge  $\{u, v\}$ . Note that  $\Gamma_{uv}(S) = O(m + n)$  in the worst case, however it is much smaller than m in many practical cases as shown in the next section. In Figure 12, the nodes in  $\gamma_{uv}(S)$  are represented in gray, while the number of double edges is  $\Gamma_{uv}(S)$ . The next theorem gives the computational complexity of Algorithm 11 as a function of  $O(\Gamma_{uv}(S))$ .

THEOREM 2.4. Algorithm 11 requires  $O(\Gamma_{uv}(S))$  time.

PROOF. Lines 1–9 require  $O(N_v(S)) = O(\Gamma_{uv}(S))$  time. In the loop at lines 10–17, variable visited ensures that each node is inserted into Q at most once. Therefore, the overall time requirement of such loop is equal to the sum of  $N_x(S)$ , for all the nodes x that are inserted into Q. Hence, to prove the statement, we show that all the nodes inserted into Q belong to  $\gamma_{uv}(S)$ . We first show that, for each  $x \in \gamma_{uv}(S)$  all the distances  $d_{xu}(S \cup \{\{u,v\}\})$  between u and x in  $G(S \cup \{\{u,v\}\})$  are correctly computed by Algorithm 11. By contradiction, suppose that the distance between some node in  $\gamma_{uv}(S)$  and u is not

networks or directed not-strongly connected networks [Boldi and Vigna 2014]. Therefore, we cannot directly use the algorithms in the literature and we devise a new dynamic algorithm.

Algorithm: DynamicBFS **Input** : An undirected graph G(S); edge  $\{u, v\}$ ; distances  $d_{ux}(S)$ , for each  $x \in V$ **Output:**  $\Delta Clo$ , the increment to  $c_u$  obtained when adding edge  $\{u, v\}$  to G(S)1  $Q := \emptyset;$ **2** visited :=  $\emptyset$ ; **3**  $d_{uv}(S \cup \{u, v\}) := 1;$ 4 foreach  $x \in N_v(S)$  do 6 visited :=  $\{u, v\} \cup N_u(S);$ 7  $\Delta Clo := 1 - \frac{1}{d_{uv}(S)};$ 8 foreach  $x \in N_v(S)$  do | Q.push(x);9 10 while  $\neg Q.empty()$  do x := Q.pop();11 
$$\begin{split} \Delta Clo &:= \Delta Clo + \frac{1}{d_{ux}(S \cup \{u,v\})(x)} - \frac{1}{d_{ux}(S)};\\ \text{for each } y \in N_x(S) \text{ do} \end{split}$$
12 13 if  $(y \notin visited) \land (d_{uy}(S) > d_{ux}(S \cup \{u, v\}) + 1)$  then 14  $d_{uy}(S \cup \{u, v\}) := d_{ux}(S \cup \{u, v\}) + 1;$ 15 Q.push(y);16  $visited := visited \cup \{y\};$ 17 18 return  $\Delta Clo$ 

Fig. 11. Algorithm DynamicBFS.

correctly computed and consider a node  $y \in \gamma_{uv}(S)$  having minimal distance to u among such nodes. At the last iteration when y is inserted into Q, there exists a node  $x \in N(y)$ such that  $d_{uy}(S) > d_{ux}(S \cup \{u, v\}) + 1$ . It follows that  $d_{uy}(S \cup \{u, v\}) = d_{ux}(S \cup \{u, v\}) + 1$ (see the test at line 14). Since the distance between y and u is minimal among those that are not correctly computed by the algorithm, then  $d_{ux}(S \cup \{u, v\})$  is correct. It follows that the distance between y and u is correctly computed at line 15, a contradiction. By contradiction, suppose that some node not in  $\gamma_{uv}(S)$  is inserted into Q and consider a node  $y \notin \gamma_{uv}(S)$  having minimal distance to u among such nodes. Since y has minimal distance to u among the nodes not in  $\gamma_{uv}(S)$  inserted into Q, then the node x for which the condition at line 14 is satisfied when y is inserted into Q must belong to  $\gamma_{uv}(S)$ . By the previous arguments,  $d_{ux}(S \cup \{u, v\}) + 1 < d_{uy}(S)$ , a contradiction to the fact that y does not belong to  $\gamma_{uv}(S)$ .  $\Box$ 

The new dynamic algorithm can now be obtained by the GREEDYIMPROVEMENT algorithm shown in Figure 3, by doing the following modifications.

- Before line 2, we compute  $c_u$  in G.
- At line 4, we incrementally compute  $c_u(S \cup \{u, v\})$  by making use of algorithm DynamicBFS instead of a full BFS.

Note that, for each  $v \in V$ ,  $\Gamma_{uv}(S)$  is maximized when  $S = \emptyset$ , then the algorithm requires an overall  $O(k \cdot n\Gamma)$  computational time, where  $\Gamma = \max_{v \in V} \{\Gamma_{uv}(\emptyset)\}$ .

We now show how to exploit the definition of submodularity to reduce the running time of iterations  $i \geq 2$  of the **for** loop at line 2 of GREEDYIMPROVEMENT. Let  $\Delta c_u(S \cup \{\{u, v\}\})$ be the increment to the centrality of node u after adding the edge  $\{u, v\}$  to graph G(S). Since  $c_u$  is submodular, then  $\Delta c_u(S \cup \{\{u, v\}\})$  is monotonic non-increasing. It follows that  $\Delta c_u(S \cup \{\{u, v\}\})$  is upper bounded by  $\Delta c_u(S' \cup \{\{u, v\}\})$ , where  $S' \subseteq S$ . We exploit this



(a) Graph G, the dashed edge is the newly added edge  $\{u, v\}$ , gray nodes and double edges are visited by Algorithm 11.

				r		$d (f_{M}, y_{1})$
Iter.	Node extracted from $Q$	$\Delta Clo$	Q	$\begin{array}{c} x \\ \hline u \\ a \\ b \end{array}$	$\begin{array}{c} u_{xu} \\ 0 \\ 1 \\ 1 \end{array}$	$\begin{array}{c} 0 \\ 1 \\ 1 \end{array}$
0	v	2/3	(c, f, g)	c U	$\frac{1}{2}$	2
1		$\frac{2}{3}$	(f,g)	$\overset{\circ}{d}$	2	2
2		$\frac{5}{0}$	(g, n, a)	v	3	1
4	h	7/6	(d, i, j)	f	3	2
5	d	7'/6	(i,j)	$g_{h}$	4	2
6	i	7/6	(j)	1 i	3	3
$\frac{7}{1}$		7/6	<u> </u>	j	4	4
(b) Itera	tions of the algo	orithm: th	e second col-	$\frac{1}{(c)}$	listone	os boforo and af

(b) Iterations of the algorithm: the second column is the node extracted from Q, the last two columns represent the status of  $\Delta Cl_0$  and Qat the end of the iteration. Iteration 0 corresponds to lines 1–9 of Algorithm 11.

(c) Distances before and after the edge addition.

Fig. 12. Example of execution of Algorithm 11.

observation in algorithm DYNAMICGREEDYIMPROVEMENT given in Figure 13. First, we compute  $c_u$  and initialize  $\Delta c_u$  (lines 1–3). For each iteration i of the **for** loop at line 6, we use variable LB (line 7) to maintain the maximum improvement to closeness found so far, that is LB is a lower bound to the improvement that will be found at the end of iteration i. If at iteration  $i \geq 2$ , for some node  $v \in V \setminus N_u(S)$ , we have that  $LB \geq \Delta c_u(\{S' \cup \{\{u, v\}\}\})$  (line 9), where S' is the value of S at iteration i - 1, then edge  $\{u, v\}$  cannot increase the value of  $c_u$  more than the maximum found so far. Therefore, in this case we prune the search. Otherwise, we compute  $\Delta c_u(S \cup \{\{u, v\}\})$  and check whether it improves LB or not (line 11). In the affirmative case, we update LB (line 12).

We can improve the performance of DYNAMICGREEDYIMPROVEMENT by means of two further heuristics. First, we sort the nodes of  $N_v(S)$ , for each  $v \in V$ , in non-increasing order of distance from u and we stop the **for** loop of line 13 of algorithm DynamicBFS when a node y such that  $d_{uy}(S) \leq d_{ux}(S \cup \{u, v\}) + 1$  is extracted. In fact, for any other node adjacent to x with a distance to u greater than  $d_{uy}(S)$  the condition at line 14 is not satisfied. Then, we can easily parallelize algorithm DYNAMICGREEDYIMPROVEMENT over p processors since  $V \setminus (N_v(S))$  can be divided into sets of  $\left\lfloor \frac{|V \setminus (N_v(S))|}{p} \right\rfloor$  nodes and the **for** 

Algorithm: DynamicGreedyImprovement **Input** : An Undirected graph G = (V, E); a vertex  $v \in V$ ; and an integer  $k \in \mathbb{N}$ **Output**: Set of edges  $S \subseteq \{\{u, v\} \mid u \in V \setminus N_v\}$  such that  $|S| \leq k$ 1 Compute  $c_u$  by using full BFS; 2 foreach  $v \in V \setminus N_u$  do  $| \Delta c_u(\{\{u,v\}\}) := 0;$ 3 **4**  $S := \emptyset$ ; **5**  $S' := \emptyset;$ 6 for i = 1, 2, ..., k do LB := 0;7 foreach  $v \in V \setminus N_u(S)$  do 8 if  $(i=1) \lor (LB < \Delta c_u(S' \cup \{\{u,v\}\}))$  then 9  $\Delta c_u(S \cup \{\{u, v\}\}) := DynamicBFS(G(S), \{u, v\}, \{d_{ux}(S)\}_{x \in V});$ 10 if  $\Delta c_u(S \cup \{\{u, v\}\}) > LB$  then 11  $LB := \Delta c_u (S \cup \{\{u, v\}\});$  $\mathbf{12}$ max := v;13 S' := S; $\mathbf{14}$ 15 $S := S \cup \{\{u, max\}\};\$ Compute distances  $d_{ux}(S)$ , for each  $x \in V$ ;  $\mathbf{16}$ 17 return S

Fig. 13. Algorithm DYNAMICGREEDYIMPROVEMENT.

loop at line 8 of algorithm DYNAMICGREEDYIMPROVEMENT can be executed in parallel for each set. In this case, LB is given by the maximum over each subset.

#### 2.6. The experimental study: part II

We also conducted a second type of experiment by measuring the improvement in the value of closeness of u and in the closeness ranking of u within the network. In particular, we studied two large real-world networks obtained from the DBLP [Ley ] and IMDB database [IMDB]. In such networks, the nodes are authors or actors and there is an edge connecting vertex x and vertex y if the author, or actor, corresponding to vertex x collaborated with y for writing a paper or for acting in the same movie. For each graph, we used twenty vertices as u but, differently from the experiments on random graphs, these vertices have been chosen on the basis of their degree ranking: in particular, we divided the list of vertices sorted by their ranking in 4 parts and chosen randomly five vertices for each interval. The value of k ranges from 1 to 10.

The analysis of these two large networks has been possible only by using the DYNAMIC-GREEDYIMPROVEMENT algorithm, since this algorithm visits only few edges of the graph (as explained in the previous section): in particular, for all the iterations i = 1, 2, ..., k the algorithm visits only 0.09% of the edges. The results for the DBLP network (n = 1305445, m = 6108727) are plotted in Fig. 14, while the results for the IMDB network (n = 1797446, m = 72880156) are similar and are shown in Fig. 15. In the chart on the left we plot the closeness centrality of vertex u as a function of k. We observe that any vertex improves its closeness value by adding just few edges. In the right chart we plot the execution time of the algorithm DYNAMICGREEDYIMPROVEMENT. We notice that the computational effort is high for k = 1 but then it is almost constant for k > 1: this is due to the submodularity property.

In order to test the scalability of the parallelized version of DYNAMICGREEDYIMPROVE-MENT algorithm, we run the same set of experiments with different numbers of cores and we measured the execution time and the speedup i.e. the ratio between the execution time



Fig. 14. (Left) Performance of DYNAMICGREEDYIMPROVEMENT algorithm on network DBLP. (Right) Execution time of DYNAMICGREEDYIMPROVEMENT algorithm.



Fig. 15. (Left) Performance of DYNAMICGREEDYIMPROVEMENT algorithm on network IMDB. (Right) Execution time of DYNAMICGREEDYIMPROVEMENT algorithm.

Table IV. Execution time and speedup of  $\rm DynamicGREEdyIMPROVEMENT$  algorithm on DBLP and IMDB networks with different number of cores.

	Avg.	Avg.	Avg.	Avg.	Avg.
	Execution Time	Speedup	Speedup	Speedup	Speedup
Network	(1  core)	(2  cores)	(4  cores)	(6  cores)	(8  cores)
DBLP	17671 s	1.91	3.01	4.01	4.63
IMDB	$10710 \ s$	1.57	2.10	3.12	3.27

with 1 core and the execution time with p cores for p = 2, 4, 6, 8. The results are reported in Table IV. We notice that the parallel algorithm shows a good scalability in terms of execution time. Note that the small increase in the case of 8 cores is due to the fact that in our machine, each CPU has 6 physical cores and hence in the case of 8 cores the computation is performed by two different processors.

#### 3. THE DIRECTED GRAPH CASE

In this section we will focus on directed graphs. After giving all necessary definitions and preliminary results, we will introduce the optimisation problem that will be considered, we will prove a non-approximability result, and we will then describe an almost optimal approximation algorithm. Finally, we will present the experiments that we have performed in order to validate this algorithm and to apply it to a quite big citation network and a web graph.

#### 3.1. The maximum directed closeness improvement problem

Let G = (V, A) be a directed graph, where V denotes the set of nodes, and A denotes the set of arcs (u, v) with u and v in V (note that  $(u, v) \in A$  does not imply that  $(v, u) \in A$ ). For each node u,  $N_u$  denotes the set of in-neighbours of u, i.e.  $N_u = \{v \mid (v, u) \in A\}$ . Given two vertices u and v,  $d_{vu}$  is defined as in the undirected graph case. Given a set S of arcs not in A, we denote by G(S) the graph augmented by adding the arcs in S to G, i.e.  $G(S) = (V, A \cup S)$ . Once again, for a parameter x of G, we denote by x(S) the same parameter in graph G(S). For each node u, the closeness centrality of u is defined as follows:

$$c_u = \sum_{\substack{v \in V \setminus \{u\}\\d_{vu} < \infty}} \frac{1}{d_{vu}}.$$

Given a directed graph G = (V, A), a vertex  $u \in V$ , and an integer k, the Maximum Directed Closeness Improvement (in short, MDCI) problem consists in finding a set S of arcs entering u not in A (that is,  $S \subseteq \{(v, u) : v \in V \setminus N_u\}$ ) such that  $|S| \leq k$  and  $c_u(S)$  is maximum.

We observe that the following results hold also for the related problem in which the edges to be added to the graph are outgoing from u and the closeness centrality considers distances  $d_{uv}$  instead of  $d_{vu}$ .

#### 3.2. The non-approximability result

In this section, in order to derive our approximation hardness result for the MDCI problem, we will make use of the *Maximum Set Coverage* (in short, MSC) problem, which is defined as follows: given a set X, a collection  $\mathcal{F} = \{S_1, S_2, \ldots S_{|\mathcal{F}|}\}$  of subsets of X, and an integer k, find a sub-collection  $\mathcal{F}' \subseteq \mathcal{F}$  such that  $|\mathcal{F}'| \leq k$  and  $s(\mathcal{F}') = |\bigcup_{S_i \in \mathcal{F}'} S_i|$  is maximised. It is known that the MSC problem cannot be approximated within a factor greater than  $1 - \frac{1}{e}$ , unless P = NP [Feige 1998]. We will now use this result in order to show that the MDCI problem does not admit a polynomial-time approximation scheme.

THEOREM 3.1. The MDCI problem cannot be approximated within a factor greater than  $1 - \frac{1}{3e}$ , unless P = NP.

PROOF. We give an *L*-reduction with parameters *a* and *b* [Papadimitriou and Yannakakis 1991]. In detail, we will give a polynomial-time algorithm that transforms any instance  $I_{\text{MSC}}$  of MSC into an instance  $I_{\text{MDCI}}$  of MDCI and a polynomial-time algorithm that transforms any solution *S* for  $I_{\text{MDCI}}$  into a solution  $\mathcal{F}'$  for  $I_{\text{MSC}}$  such that the following two conditions are satisfied for some values *a* and *b*:

$$OPT(I_{\rm MDCI}) \le aOPT(I_{\rm MSC})$$
 (2)

$$OPT(I_{\rm MSC}) - s(\mathcal{F}') \le b \left( OPT(I_{\rm MDCI}) - c_u(S) \right).$$
(3)

where OPT denotes the optimal value of an instance of an optimization problem. If the above conditions are satisfied and there exists a  $\alpha$ -approximation algorithm for MDCI, then there exists a  $(1 - ab(1 - \alpha))$ -approximation algorithm for MSC [Papadimitriou and Yannakakis 1991]. Since MSC is hard to approximate within a factor greater than  $1 - \frac{1}{e}$ , then  $1 - ab(1 - \alpha) < 1 - \frac{1}{e}$ , unless P = NP. This implies that  $\alpha < 1 - \frac{1}{e}$ .

then  $1 - ab(1 - \alpha) < 1 - \frac{1}{e}$ , unless P = NP. This implies that  $\alpha < 1 - \frac{1}{abe}$ . Given an instance  $I_{\text{MSC}} = (X, \mathcal{F}, k)$  of MSC, we define an instance  $I_{\text{MCI}} = (G, u, k)$  of MDCI as follows (see Fig. 16): G = (V, A), where  $V = \{u\} \cup \{v_{x_i} \mid x_i \in X\} \cup \{v_{S_j} \mid S_j \in \mathcal{F}\}$  and  $A = \{(v_{x_i}, v_{S_j}) \mid x_i \in S_j\}$ .

Without loss of generality, we can assume that any solution S of MDCI contains only arcs  $(v_{S_j}, u)$  for some  $S_j \in \mathcal{F}$ . In fact, if a solution does not satisfy this property, then we can improve it in polynomial time by repeatedly applying the following rule: if S contains an arc  $(v_{x_i}, u)$ , for some  $x_i \in X$ , then exchange such arc with an arc  $(v_{S_j}, u)$  such that



Fig. 16. The reduction used in Theorem 2.1 (in this example,  $x_1 \in S_1$ ,  $x_1 \in S_2$ ,  $x_2 \in S_1$ , and  $x_2 \in S_{|\mathcal{F}|}$ ). The dashed arcs denote those added in a solution.

 $(v_{S_j}, u) \notin S$  (note that such an arc must exist, since otherwise  $|\mathcal{F}| \leq k$  and  $I_{\text{MSC}}$  could be easily solved). The above rule does not decrease the value of  $c_u(S)$ : indeed, if we exchange an arc  $(v_{x_i}, u)$  with an arc  $(v_{S_j}, u)$  such that  $(v_{S_j}, u) \notin S$ , then the closeness centrality of u decreases by either 1 or  $\frac{1}{2}$  (because of the deletion of  $(v_{x_i}, u)$ ) but certainly increases by 1 (because of the insertion of  $(v_{S_i}, u)$ ).

Given a solution S of MDCI, let  $\mathcal{F}'$  be the solution of MSC such that  $S_j \in \mathcal{F}'$  if and only if  $(v_{S_j}, u) \in S$ . We now show that  $c_u(S) = \frac{1}{2}s(\mathcal{F}') + k$ . To this aim, let us note that the distance from a vertex  $v_{x_i}$  to u is equal to 2 if an arc  $(x_{S_j}, u)$  such that  $x_i \in S_j$  belongs to S, and it is  $\infty$  otherwise. Similarly, the distance from a vertex  $v_{S_j}$  to u is equal to 1 if  $(x_{S_j}, u) \in S$ , and it is  $\infty$  otherwise. Moreover, the set of elements  $x_i$  of X such that  $d_{v_{x_i}u}(S) < \infty$  is equal to  $\{x_i \mid x_i \in S_j \land (v_{S_j}, u) \in S\} = \bigcup_{S_i \in \mathcal{F}'} S_j$ . Therefore,

$$\begin{aligned} c_u(S) &= \sum_{\substack{v \in V \setminus \{u\} \\ d_{vu}(S) < \infty}} \frac{1}{d_{vu}(S)} = \sum_{\substack{x_i \in X \\ d_{vx_iu}(S) < \infty}} \frac{1}{d_{vx_iu}(S)} + \sum_{\substack{S_j \in \mathcal{F} \\ d_{vS_ju}(S) < \infty}} \frac{1}{d_{vS_ju}(S)} \\ &= \frac{1}{2} |\{x_i \in X \mid d_{vx_iu}(S) < \infty\}| + |\{S_j \in \mathcal{F} \mid d_{vS_ju}(S) < \infty\}| \\ &= \frac{1}{2} \left| \bigcup_{S_j \in \mathcal{F}'} S_j \right| + |\{S_j \mid (v_{S_j}, u) \in S\}| = \frac{1}{2} s(\mathcal{F}') + k. \end{aligned}$$

It follows that Conditions (2) and (3) are satisfied for  $a = \frac{3}{2}$  and b = 2. Indeed,  $OPT(I_{\text{MDCI}}) = \frac{1}{2}OPT(I_{\text{MSC}}) + k \leq \frac{3}{2}OPT(I_{\text{MSC}})$ , where the inequality is due to the fact that  $OPT(I_{\text{MSC}}) \geq k$ , since otherwise the greedy algorithm would find an optimal solution for  $I_{\text{MSC}}$ . Moreover,  $OPT(I_{\text{MSC}}) - s(\mathcal{F}') = 2(OPT(I_{\text{MDCI}}) - k) - 2(c_u(S) - k) = 2(OPT(I_{\text{MDCI}}) - c_u(S))$ . The theorem follows by plugging the values of a and b into  $\alpha < 1 - \frac{1}{abe}$ .  $\Box$ 

#### 3.3. The greedy approximation algorithm

As in the case of undirected graphs, we can show that, for each vertex u, the function  $c_u$  is monotone and submodular with respect to any feasible solution for MDCI. Indeed, the proof of Theorem 2.2 can be easily adapted to the directed graph case, and the following result holds.

COROLLARY 3.2. The algorithm shown in Fig. 17 is a  $(1-\frac{1}{e})$ -approximation algorithm for the MDCI problem.

Also in this case, there is a gap between the non-approximability result proved in Theorem 3.1 (that is, the upper bound equal to  $1 - \frac{1}{3e} \approx 0.88$ ), and the approximability result of the above corollary (that is, the lower bound  $(1 - \frac{1}{e}) \approx 0.63$ ). One of the main goals of the 

 Algorithm: DIRECTEDGREEDYIMPROVEMENT

 Input : a directed graph G = (V, A), a vertex  $u \in V$ , and an integer  $k \in \mathbb{N}$  

 Output: set of arcs  $S \subseteq \{(v, u) \mid v \in V \setminus N_u\}$  such that  $|S| \leq k$  

 1  $S := \emptyset$ ;

 2 for  $i = 1, 2, \dots, k$  do

 3 [ foreach  $v \in V \setminus N_u(S)$  do

 4 [ Compute  $c_u(S \cup \{(v, u)\})$ ]

 5 [ S := S \cup \{(v\_{\max}, u)\};

 7 return S;

 Fig. 17. The greedy algorithm for directed graphs.

next experimental session is to analyse the "real" performance, in terms of solution quality, of the greedy algorithm on relatively small real-world and synthetic graphs.

#### 3.4. Experimental results

As in the undirected graph case in this section we analyse the greedy algorithm from an experimental point of view. First, we compare the solution of the greedy algorithm with the optimal solution in order to assess its real performance in terms of solution quality. Then, we compare the greedy algorithm with the other approaches used in the undirected cases, but K-MEDIAN because it cannot be applied in the directed case. We adapted the DYNAMICGREEDYIMPROVEMENT algorithm used for the undirected case. In particular, we run the (pruned) BFSs on the transpose graph of G to reduce the time required to compute  $c_u$  and to improve the computational complexity of GREEDYIMPROVEMENT. Moreover, we show the results of our experiments on a real-world graph measuring the improvement in the value of closeness of u within the network. Finally, we analyse how the information spread increases.

We measured the approximation ratio of the greedy algorithm on five types of randomly generated directed networks, namely directed Preferential Attachment (in short, PA) [Bollobás et al. 2003], Erdős-Rényi (in short, ER) [Erdős and Rényi 1959], Copying (in short, COPY) [Kumar et al. 2000], Compressible Web (in short, COMP) [Chierichetti et al. 2009] and Forest Fire (in short, FF) [Leskovec et al. 2007]. The size of the graphs is reported in Table V. For each combination (n, m), we generated five random directed graphs and used twenty vertices as u. These vertices have been chosen on the basis of their original closeness ranking as in the undirected case.

The results of the experiments are reported in Table V, where we show, similarly to the undirected graph case, the minimum ratio obtained. The experiments show that in the worst case the ratio is 0.9668.

For the comparison with the other approaches we used real-world citation networks obtained from the Arnetminer database [arn ] (see Table VI for details). In the Arnetminer's networks, there is a vertex for each author and an arc from vertex x to vertex y if the author corresponding to vertex x cited in his paper one paper written by the author corresponding to y. We parsed the Arnetminer database in order to select a sub-network induced by the authors that published at least a paper in one of the main conferences or journals. As in the previous experiment, for each graph, we used twenty vertices as u. The value of k ranges from 1 to 10.

The results for the citation network Information Security are plotted in Fig. 18. In the two charts we plot the closeness centrality and the ranking of vertex u as a function of k. We observe that any vertex becomes central by adding just few arcs. For example a

Table V. Comparison between the DIRECTEDGREEDYIMPROVE-MENT algorithm and the optimum. The first three columns reports the type and size of the graphs; the fourth column reports the approximation ratio.

Network	n =  V	m =  E	Min Approx. Ratio
PA	100	130	0.9816
PA	500	650	0.9956
PA	1000	1300	1
ER	100	200	0.9668
ER	100	500	0.9744
ER	100	1000	0.9780
ER	500	5000	0.9890
ER	500	12500	0.9819
ER	500	25000	0.9994
COMP	100	200	0.9968
COMP	100	500	0.9764
COMP	100	1000	1
COMP	500	5000	0.9848
COMP	500	12500	1
COMP	500	25000	1
COPY	100	200	0.9911
COPY	100	500	0.9753
COPY	100	1000	0.9820
COPY	500	5000	0.9825
COPY	500	12500	0.9726
COPY	500	25000	0.9690
FF	100	200	0.9911
FF	200	400	0.9714
FF	500	1000	0.9892

Table VI. Collaboration networks obtained from Arnetminer database.

Network	n =  V	m =  E
Software Engineering	3141	14787
Information Security	1067	4253
Computer Graphics Multimedia	8336	41925
Theoretical Computer Science	4172	14272
Artificial Intelligence	27617	268460
High-Performance Computing	4869	35036
Computer Networks	9420	53003
Interdisciplinary Studies	577	1504



Fig. 18. Performance of the DIRECTEDGREEDYIMPROVEMENT algorithm on network Information Security.

vertex with the smallest closeness centrality which initially has closeness 0 and is ranked 509, improves its closeness and ranking to 213.32 and 1, respectively, by adding only 7 arcs.



(c) RANDOM

Fig. 19. Average relative ranking position between the Ranking obtained by the DIRECTEDGREEDYIM-PROVEMENT algorithm and the different baselines in the network Information\_security.

In the chart in Fig. 19 we compare the greedy algorithm with the other approaches. We report the comparison of the average relative ranking position reached by the nodes. We do not show the results for ROUNDING as such algorithm is not able to terminate within a reasonable amount of time. The experiments clearly show that the greedy algorithm outperforms the other approaches.

We further used the DYNAMICGREEDYIMPROVEMENT algorithm to analyse a web network [LAW]. The results for the web network uk-2007 (n = 100000, m = 3050615) are plotted in Fig. 20. In the right chart we report the execution time of the algorithm. The DYNAMICGREEDYIMPROVEMENT algorithm is up to  $10^3$  times faster than the basic GREEDY-IMPROVEMENT algorithm and for all the iteration it visits only the 0.18% of the arcs of the graph: using the DYNAMICGREEDYIMPROVEMENT algorithm, it is possible to solve the MCI problem on very large graphs where it is impossible to obtain a solution using the GREEDYIMPROVEMENT algorithm.

Like in the undirected case, we run the same set of experiments with different numbers of cores and we measured the execution time and the speedup i.e. the ratio between the execution time with 1 core and the execution time with p cores for p = 2, 4, 6, 8. The results are reported in Table VII. We notice that the parallel algorithm shows a very good scalability in terms of execution time.

Finally, we measured the increase in information spreading when few edges are added to a small set of randomly-chosen seeds. We performed experiments similar to those shown in the undirected case in the graphs of Table VI. In Fig. 21, we plot the results for the Information\_security directed network. We observe that also in this case the informed nodes percentage increases. The results for the other networks in Table VI are similar.



Fig. 20. (Right) Performance of the DYNAMICGREEDYIMPROVEMENT algorithm on network uk-2007. (Left) Time of execution.

Table VII. Execution time and speedup of DYNAMICGREEDYIMPROVEMENT algorithm on uk-2007 network with different number of cores.

	Avg.	Avg.	Avg.	Avg.	Avg.
	Execution Time	Speedup	Speedup	Speedup	Speedup
Network	(1  core)	(2  cores)	(4  cores)	(6  cores)	(8  cores)
uk-2007	1382 s	1,83	3,70	5,58	7,44

#### 4. CONCLUSION AND FUTURE RESEARCH

We considered the problem of adding k edges in a (directed or undirected) graph in order to maximize the closeness of a predefined vertex. For undirected graphs, we have shown that the problem cannot be approximated within a factor larger than  $1 - \frac{1}{15e}$ , and we proposed a greedy algorithm that guarantees an approximation factor of  $1 - \frac{1}{e}$ . For directed graphs, the problem cannot be approximated within a factor larger than  $1 - \frac{1}{3e}$ , while the greedy algorithm still guarantees an approximation factor of  $1 - \frac{1}{e}$ . We experimentally evaluated such algorithms and showed that they often compute an optimal solution and, in any case, they achieve an approximation factor significantly better than the theoretical one. Moreover, by adding very few edges a vertex can drastically increase its centrality measure and its ranking.

As future works, we plan to extend our work to further centrality measures such as betweenness, to generalize the problem by allowing the addition of edges incident to other nodes of the graph, and to maximize the ranking of a node instead of the centrality value.

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Fig. 21. Percentage of active nodes in Information\_security network as a function of k. Parameter a denotes the threshold and the percentage of seeds is equal to 2%, 4%, 6%, 8%, 10%, respectively.

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