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Detecting overlapping communities in complex networks using non-cooperative games

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ABSTRACT

Detecting communities in complex networks is of paramount importance, and its wide range of real-life applications in various areas has caused a lot of attention to be paid to it, and many efforts have been made to have efficient and accurate algorithms for this purpose. In this paper, we proposed a non-cooperative game theoretic-based algorithm that is able to detect overlapping communities. In this algorithm, nodes are regarded as players, and communities are assumed to be groups of players with similar strategies. Our two-phase algorithm detects communities and the overlapping nodes in separate phases that, while increasing the accuracy, especially in detecting overlapping nodes, brings about higher algorithm speed. Moreover, there is no need for setting parameters regarding the size or number of communities, and the absence of any stochastic process caused this algorithm to be stable. By appropriately adjusting stop criteria, our algorithm can be categorized among those with linear time complexity, making it highly scalable for large networks. Experiments on synthetic and real-world networks demonstrate our algorithm's good performance compared to similar algorithms in terms of detected overlapping nodes, detected communities size distribution, modularity, and normalized mutual information.

Introduction

Complex relationships between components existing in society, technology, biology, economy, and other various fields, in many cases, can be modeled as complex networks by regarding components as nodes and relationships as edges¹. As a consequence, all of tools available for complex networks analysis could be applied to extract valuable information about under investigation system. An important consideration of network structures is the possibility of classifying nodes into groups or communities². Indeed, it is observed that many of real-world networks have community structure³. In a network, community refers to a group of nodes that are densely connected internally and have sparser connection with the rest of the network³. Detecting communities is of great importance since nodes in a community usually have similarities in function, property, and characteristics⁴. For instance, community detection in the network of protein-protein interaction could reveal groups of closely connected proteins that possess an identical function in the body⁵. The discovery of community structure can be constructive in many fields such as drug discovery⁶, precision marketing⁷, brain neural network⁸, online social interaction analysis⁹, and public opinion analysis¹⁰. Network communities typically can be categorized in two types. Disjoint communities with no shared member (also called non-overlapping communities or partitions) and overlapping communities with shared members (also called covers). Examples of overlapping communities are widely seen in real world. Researchers, based on their various research interests or multiple affiliations, can be a member of more than one research group, or a gene can be involved in causing various diseases¹¹. As a result, it is crucial to design community detection algorithms that be able to identify overlapping nodes. In recent years, a variety of approaches have been employed for this purpose. The use of game theory in this context was initialized by Athey and Jha in 2006 to model an organization workers interaction¹² and followed by game theoretic based algorithm proposed by Chen in 2010^{13,14}. A comprehensive discussion of game theoretic-based methods for detecting community structure in networks is provided in a survey done by Jonnalagadda and Kuppusamy¹⁵. However, the number of game theoretic based algorithms proposed in last decay is not very large and most of them are not scalable for large networks¹¹. Community detection algorithms using the game theory are typically based on cooperative or non-cooperative games. Our proposed algorithm is based on the non-cooperative game in which nodes are assumed as rational selfish players who decide to be part of the communities which bring them the most profit. Although, our algorithm is designed to detect overlapping communities, in contrast with similar algorithms, nodes are not allowed to be part of multiple communities before the exact boundaries of communities are determined (phase one), and overlapping nodes is identified in phase two. Such two phases algorithm not only increase accuracy but also, along with the appropriate stop criterion used in current work, speed up convergence. Moreover, in the present work, players have only local interactions, which leads the algorithm to be more effective than some other game theoretic based algorithms in which interaction with all nodes is considered in the utility function^{13,16,17}. The remainder of this paper is organized as follows. In next section, the framework of the proposed algorithm and related definitions are given and it is followed a discussion on the time complexity of the algorithm. Afterward,

the experimental results of our algorithm and its comparison with some other state-of-the-art algorithms are given. Finally, the concluding remark is stated.

Proposed Algorithm

The proposed algorithm consists of two phases. The non-cooperative game is the basis of the first phase leading to non-overlapping community detection, while in the second phase, the overlap of the communities is determined. The game-theoretic framework is based on considering each node as a selfish agent trying to maximize its payoff by choosing different strategies, and each agent's choice can influence the other ones'. Strategy is a term in game theory which in the current context, refers to the communities to which the agent wants to participate. Based on this, each agent's strategy s_i is actually a list of community labels it is a member of and strategy profile of all agents is defined as $S = (s_1, s_2, \dots, s_n)$. As stated, each agent aims to maximize its payoff, which for the agent v_i is represented through a utility function defined as follows.

$$U(S_{-i}, s_i) = \sum_{a_{ij} \neq 1} (1 + sim_{ij}) \frac{|s_i \cap s_j|}{\sqrt{|s_j|}} \quad (1)$$

Where a_{ij} is the adjacency matrix element; s_i is the strategy of agent v_i and S_{-i} is strategy profile of all agents but her; $|s_i \cap s_j|$ is the number of common labels between agent v_i and v_j ; and $|s_j|$ is the number of communities agent v_j belongs to. Unlike some other game-theoretic overlapping community detection algorithms, in phase one agents are not allowed to acquire multiple labels and consequently, expression $\frac{|s_i \cap s_j|}{|s_j|}$ can only have two values of 0 or 1. Also, in this phase agents are only allowed to do switch operation among different community labels. In utility function, sim_{ij} is the similarity between agents v_i and v_j , which can be calculated through different available metrics as follows.

$$Jaccard\ coefficient(JC) : sim_{ij}^{JC} = \frac{|\Gamma_i \cap \Gamma_j|}{|\Gamma_i \cup \Gamma_j|} \quad (2)$$

$$Saltin\ index(SI) : sim_{ij}^{SI} = \frac{|\Gamma_i \cap \Gamma_j|}{\sqrt{|\Gamma_i| |\Gamma_j|}} \quad (3)$$

$$Sorensen\ index(SO) : sim_{ij}^{SO} = \frac{2|\Gamma_i \cap \Gamma_j|}{|\Gamma_i| + |\Gamma_j|} \quad (4)$$

$$Hub\ promoted\ index(HP) : sim_{ij}^{HP} = \frac{|\Gamma_i \cap \Gamma_j|}{\min(|\Gamma_i|, |\Gamma_j|)} \quad (5)$$

$$Hub\ depressed\ index(HD) : sim_{ij}^{HD} = \frac{|\Gamma_i \cap \Gamma_j|}{\max(|\Gamma_i|, |\Gamma_j|)} \quad (6)$$

Where Γ_i and Γ_j denote the neighbors of agent v_i and v_j respectively. The proposed algorithm results do not significantly depend on different similarity metrics except for a few special cases. However, represented results have been obtained using HP similarity, which slightly performs better than other similarity metrics. The algorithm starts with an initial condition in which each agent v_i is assigned to a singleton community c_i . Next, in each iteration, all agents, by order of their degrees, update their strategy by imitating their neighbors with the aim of maximizing payoff. For more clarity, the phase one framework is given in Algorithm 1 in Figure 1.

Lines 4 to 23 repeat until the stop criterion is met and finally agents with the same label belong to the same community. The stop criterion should be defined in a way that satisfies accuracy and efficiency at the same time. In the proposed algorithm, there are some cases in which some agents' strategy fluctuates permanently, and some other agents need too many iterations to reach their stable one. Since a minimal number of agents often fall in such category, defining stop criterion that ignore such agents stability could speed up algorithm without significant loss of accuracy. For this reason, instead of waiting for all agents' strategy to be fixed, the stop criterion is satisfied as soon as the number of agents with a fixed strategy does not

Algorithm 1 phase one

Input : A Network $G = (V, E)$
Output : single strategy profile of all agents S

- 1 : making initial partition : assign different label to each agent resulting singleton partitions.
- 2 : $iter \leftarrow 0$
- 3 : $\Delta \leftarrow 0$
- 4 : **while** $\Delta \leq \Delta_{stop}$ **and** $iter < 2$ **do**
- 5 : $iter \leftarrow iter + 1$
- 6 : $n_{fixed}[iter] \leftarrow n$
- 7 : **for** agent v_i in G **do**
- 8 : $U0 \leftarrow Utility(v_i, s_i)$
- 9 : **for** each label l available in the vertex v_i neighborhood **do**
- 10 : **if** $Utility(v_i, \{l\}) > U0$ **then**
- 11 : $s_{new} \leftarrow \{l\}$
- 12 : **end if**
- 13 : **end for**
- 14 : **if** $s_{new} \neq s_i$ **then**
- 15 : $s_i \leftarrow s_{new}$
- 16 : $n_{fixed}[iter] \leftarrow n_{fixed}[iter] - 1$
- 17 : **end if**
- 18 : **end for**
- 19 : **if** $iter > 2$ **then**
- 20 : $\Delta \leftarrow |n_{fixed}[iter] - n_{fixed}[iter - 1]|$
- 21 : $\Delta_{stop} \leftarrow \alpha \cdot n_{fixed}[iter - 1]$
- 22 : **end if**
- 23 : **end while**

Algorithm 2 phase two

Input : A Network $G = (V, E)$ and non overlapping strategy profile S
Output : overlapping strategy profile \bar{S}

- 1 : **for** agent v_i in G **do**
- 2 : **for** each label l available in the vertex v_i neighborhood **do**
- 3 : $add\ Utility(v_i, \{l\})\ to\ U_list[v_i]$
- 4 : **end for**
- 5 : $max_U[v_i] = \max(U_list[v_i])$
- 6 : **for** each label l available in the vertex v_i neighborhood **do**
- 7 : $add\ Utility(v_i, \{l\}) / max_U[v_i]\ to\ Normalized_U_list[v_i]$
- 8 : **end for**
- 9 : $Thresh_list[v_i] = RMS(Normalized_U_list[v_i])$
- 10 : **end for**
- 11 : **for** agent v_i in G **do**
- 12 : **for** each label l available in the vertex v_i neighborhood **do**
- 13 : **if** $> Thresh_list[v_i]$ **then**
- 14 : **if** $Utility(v_i, \{l\}) / max_U[v_i] > Thresh_list[v_i]$ **then**
- 15 : $add\ l\ to\ \bar{S}[v_i]$
- 16 : **end if**
- 17 : **end for**
- 18 : **end for**

Figure 1. Phase one and Phase two framework algorithms.

increase more than a specific value. This value in each iteration Δ_{stop} is defined as a fraction of fixed agents number n_{fixed} in previous iteration.

$$\Delta_{stop} = \varepsilon \cdot n_{fixed} \quad (7)$$

By adjusting ε value, a balance between accuracy and efficiency can be obtained. Variation of relative phase one execution time (execution time divided by longest execution time) and relative NMI (Obtained NMI divided by the best achievable NMI) obtained for LFR synthetic networks is represented in Figure 2. According to the results, nonzero but small values of ε such as 0.005 and 0.01 can reduce phase one elapsed time while giving acceptable accuracy. The effect of ε value on scalability of algorithm will be discussed more in algorithm complexity section.

Phase two is responsible for finding overlapping nodes. In some non-cooperative game theoretic algorithms a *loss function* is used as a method for controlling multiple membership of agents^{13, 16–18}. In such method multiple membership criterion usually is defined in a way that is similar for all nodes in spite of different condition they may have. Moreover, in some other algorithms like^{19, 20}, the manually defined threshold is responsible for determining multiple memberships of nodes. Nevertheless, in our algorithm, this criterion is defined uniquely for each agent based on payoffs it acquires from membership in different communities. Accordingly, phase two contains two stages. In the first stage in which payoff thresholds is calculated, following operations should be done for each agent:

1. Calculating payoffs that agent acquires by Adopting any of community label available in its neighborhood.
2. Normalizing all obtained payoffs with respect to maximum payoff the agent has obtained.
3. Finding payoff threshold for the agent by calculating root mean square of normalized payoff values obtained for that agent.

In following stage, each agent adds community labels that have payoff above her payoff threshold value to her strategy. The framework of phase two is given in Algorithm 2 in Figure 1.

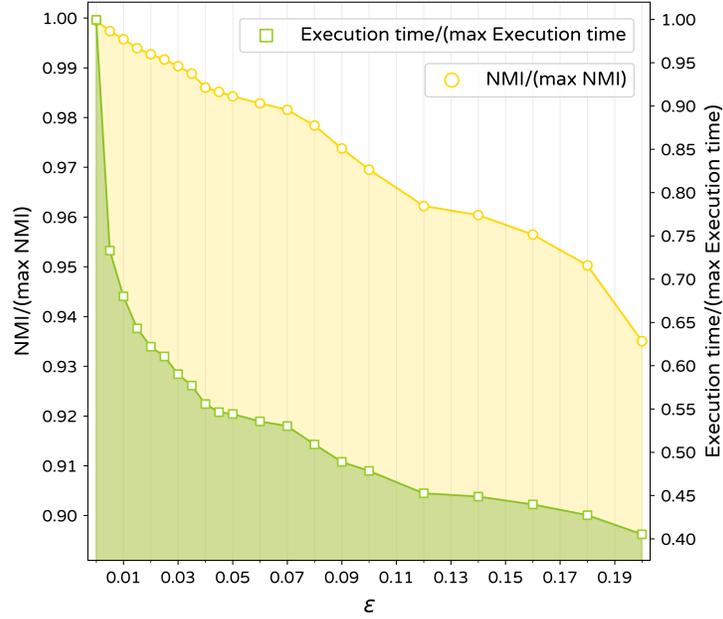


Figure 2. Effect of ϵ value on algorithm accuracy and elapsed time.

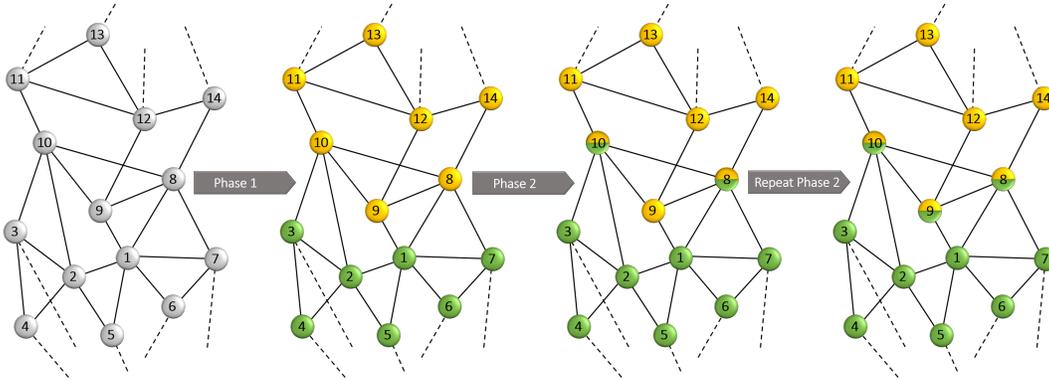


Figure 3. The toy example to illustrate the performance of each phase.

Finally, each agent belongs to all communities which those labels exist in its strategy list. In networks with high degree of overlap, it is very probable for overlapping nodes to be connected with another overlapping ones. In such cases repetition of phase two can help discover overlapping nodes more reliably. For more illustration, a toy model representing community structure before and after applying each phase is shown in Figure 3.

It should be noted that described phase two returns crisp communities with binary membership coefficient of nodes in different communities. Although often it is desirable form, sometimes the fuzzy communities are more suitable for intended use. In such cases, the normalized payoff values of each agent are representative of that agent fuzzy membership coefficients.

Time Complexity of Algorithm

The proposed algorithm consists of three parts. The first one is initialization that requires $O(n)$, where n is total number of nodes. In phase one, the outer loop continues until stop criterion satisfaction. In inner loops, for each agent, the payoff should be calculated for all labels in the neighborhood, which is maximally equal to the number of agent's neighbors. Therefore, phase one requires $O(T.n.K)$ on average, where K is the average degree and T is maximum iteration. In some other algorithms T is defined manually. In the proposed algorithm, although the maximum iteration number is determined dynamically based on stop criterion satisfaction, it does not depend on n or the total number of edges m if the network topology is kept the same and if the ϵ value is selected appropriately. For LFR synthetic networks, the variation of maximum iteration number for three small values of ϵ were calculated as a function of n and m (Figure 4 (a,b)). As it can be seen, especially for small

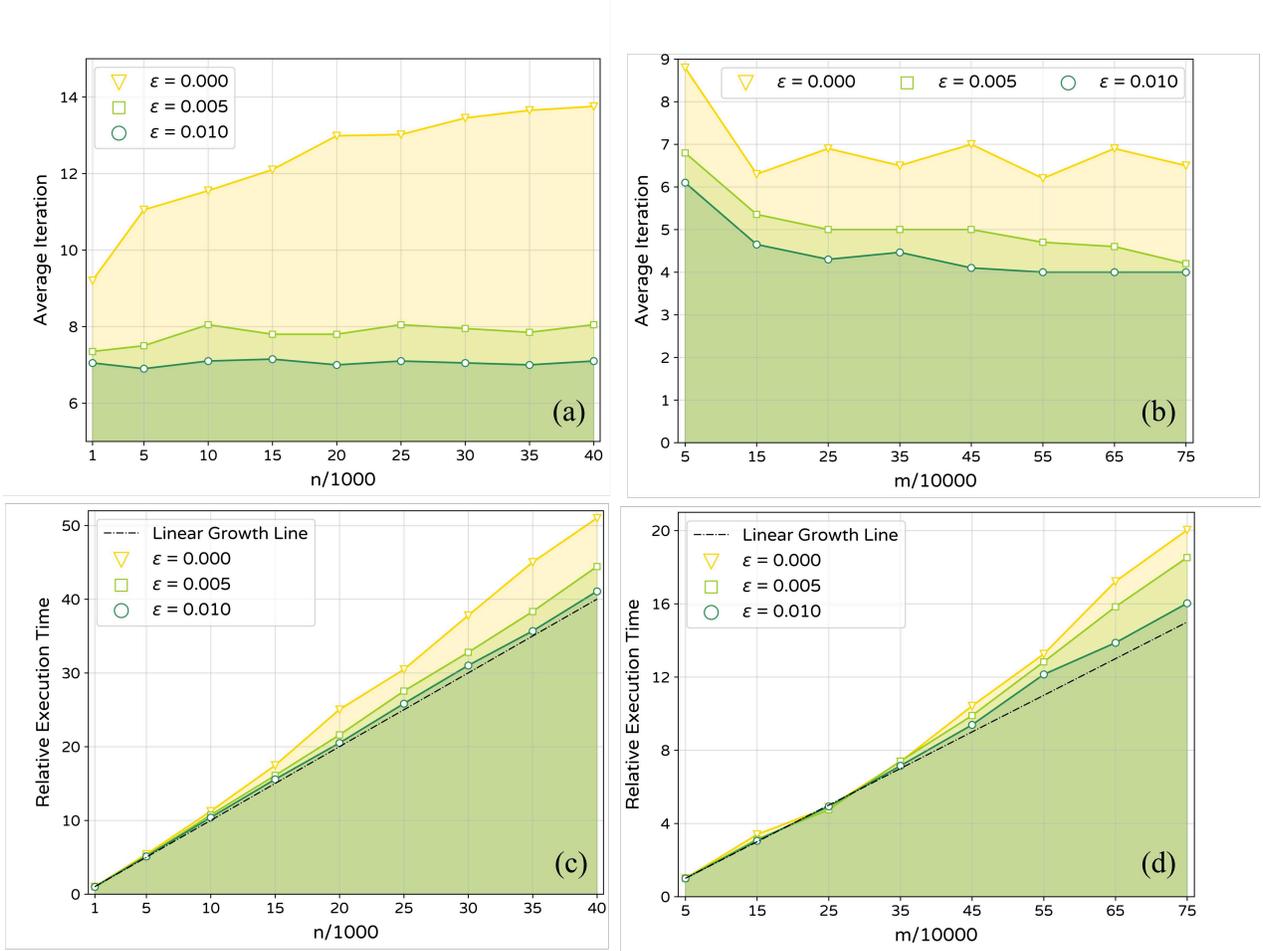


Figure 4. a,b) Phase one average max iteration as a function of n and m . c,d) Entire relative algorithm execution time as a function of n and m . Results of a and c is obtained for LFR networks with $\bar{k} = 10$. Results of b and d is obtained for LFR networks with $n = 5000$.

nonzero values of ϵ the maximum iteration number does not depend considerably on n or m . The phase two have similar calculation structure as inner part of phase one algorithm. considering the second phase repeats for two times, it requires $O(2n)$. Therefore, time complexity of entire algorithm is $O(n)$ in sparse networks and $O(m)$ in arbitrary ones. For a naive implementation of algorithm, Figure 4 (c,d) shows the execution time for LFR synthetic networks. As it can be seen, for ϵ value of 0.01 the execution time is just slightly slower than linear growth.

Experimental Results and Comparison

With the aim of evaluating our proposed algorithm performance, we compare it with some other algorithms named GAME1¹³, GAME2²¹, GAME3²², SLPA¹⁹, OSLOM²³, CPM²⁴, GCE²⁵ and LFM²⁶. GAME1 is based on non-cooperative game theory with time complexity of $O(m^2)$. GAME 2 and GAME3 are based on cooperative game theory with time complexity of $O(n^2)$ and $O(n \cdot \log(n)) + O(n \cdot k_{max})$, respectively (k_{max} is graph maximum degree). Our algorithm results in this section is obtained by set ϵ value to 0.01 and for other algorithms with tunable parameters, the results with the best setting are reported. Moreover, the reported results of other algorithms were obtained from what have reported in the comparative study papers²⁷ or each algorithm original paper.

Evaluation Criteria

There are various metrics in order to evaluate obtained results of algorithms. One of the most common metric for networks with ground truth is normalized mutual information (NMI). An extended version of it which is used in current study and is appropriate for comparison of two overlapping communities is the one proposed by²⁶. The closer value of NMI to 1, the more

similar the detected community structure to ground truth; and the 0 value indicates the least similarity.

When it comes to testing the performance of overlapping community detection algorithms, especially when the ground truth of communities is unknown, the Q_{ov} is a well-known and frequently used metric²⁸. It is an extension of the classical modularity, and the higher value of this means the better-detected communities. For directed networks this metric is defined as follow:

$$Q_{ov} = \frac{1}{m} \sum_{c \in C} \sum_{i,j \in V} \left[\beta_{l(i,j),c} A_{j,j} - \frac{\beta_{l(i,j)}^{out} k_i^{out} \beta_{l(i,j)}^{in} k_j^{in}}{m} \right] \quad (8)$$

By applying minor changes as follows, it can be used for undirected networks:

$$Q_{ov} = \frac{1}{2m} \sum_{c \in C} \sum_{i,j \in V} \left[\beta_{l(i,j),c} A_{j,j} - \frac{\beta'_{l(i,j)} k_i \beta'_{l(i,j)} k_j}{2m} \right] \quad (9)$$

The components of this equation is given by:

$$\beta'_{l(i,j)} = \beta_{l(i,j)}^{out} = \beta_{l(i,j)}^{in} = \frac{\sum_{i \in V} F(\alpha_{i,c}, \alpha_{j,c})}{|V|} \quad (10)$$

$$\beta_{l(i,j)} = F(\alpha_{i,c}, \alpha_{j,c}) \quad (11)$$

$$k_i^{out} = k_i^{in} = k_i \quad (12)$$

$$F(\alpha_{i,c}, \alpha_{j,c}) = \frac{1}{(1 + e^{f(\alpha_{i,c})})} \quad (13)$$

$$f(x) = 2px - p, p \in \mathbb{R} \quad (14)$$

Where $\alpha_{(i,c)}$ is the belonging coefficient of node i to community c and p in $f(x)$ is an arbitrary value that in the current study is set to 30.

Synthetic Networks

	n	\bar{k}	k_{max}	μ	τ_1	τ_2	Min Community Size (minC)	Max Community Size (maxC)	Om	On
LFR1	1000	10	50	0.1	2	1	20	100	2-6	0.1
LFR2	1000	10	50	0.3	2	1	20	100	2-6	0.1
LFR3	5000	10	50	0.3	2	1	20	100	2-6	0.1
LFR4	5000	10	50	0.3	2	1	20	100	2-6	0.5
LFR5	5000	10	50	0.3	2	1	10	50	2-6	0.1
LFR6	5000	10	50	0.3	2	1	10	50	2-6	0.5

Figure 5. LFR synthetic networks used for performance tests.

The most widely used benchmark network that has been used frequently for performance tests of community detection algorithms is synthetic networks called LFR which can be generated by the method proposed by Lancichinetti and Fortunato²⁹.

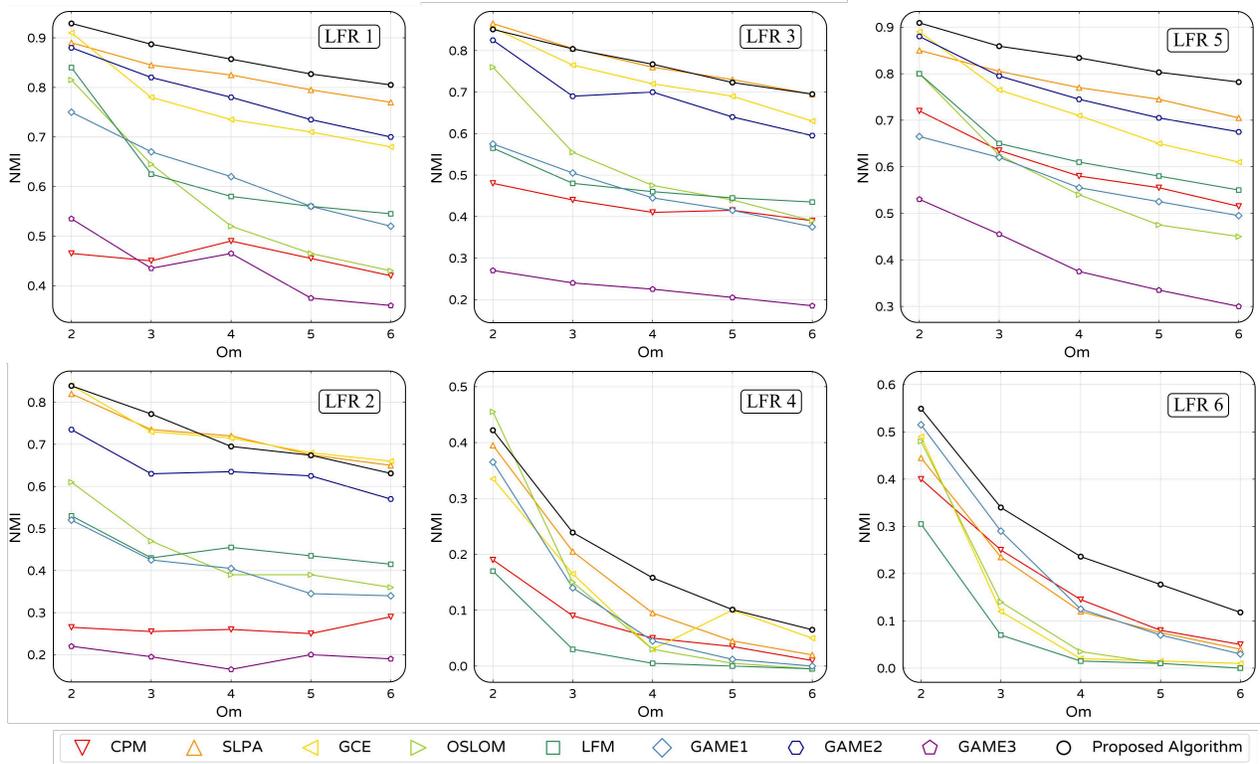


Figure 6. Comparative NMI value for proposed and other algorithms on LFR synthetic networks listed in Figure 5.

In the networks made by this method 10 parameters are adjustable. By setting these parameters we generated 6 group of LFR network for the performance tests as shown in Figure 5. The mixing parameter μ refers to fraction of links through which a node connects to other nodes in other communities; $k_i^{in} = (1 - \mu)k_i$. τ_1 and τ_2 are exponents of power law distribution of node degrees and community sizes respectively. Furthermore, overlapping features of LFR network is controlled by Om (the number of communities to which each overlapping node belongs) and On (the fraction of nodes that belongs to more than one community). It should be noted that for our algorithm performance test on LFR networks, we have reported averaged results of runs over at least 10 instantiations of these networks for each parameter set.

The NMI value for results obtained using our proposed and other algorithms are represented in Figure 6. As expected, by increasing Om the NMI value gradually decrease. However, it is observed that in most cases our algorithm outperforms others, especially in synthetic networks with smaller community sizes and more overlapping nodes.

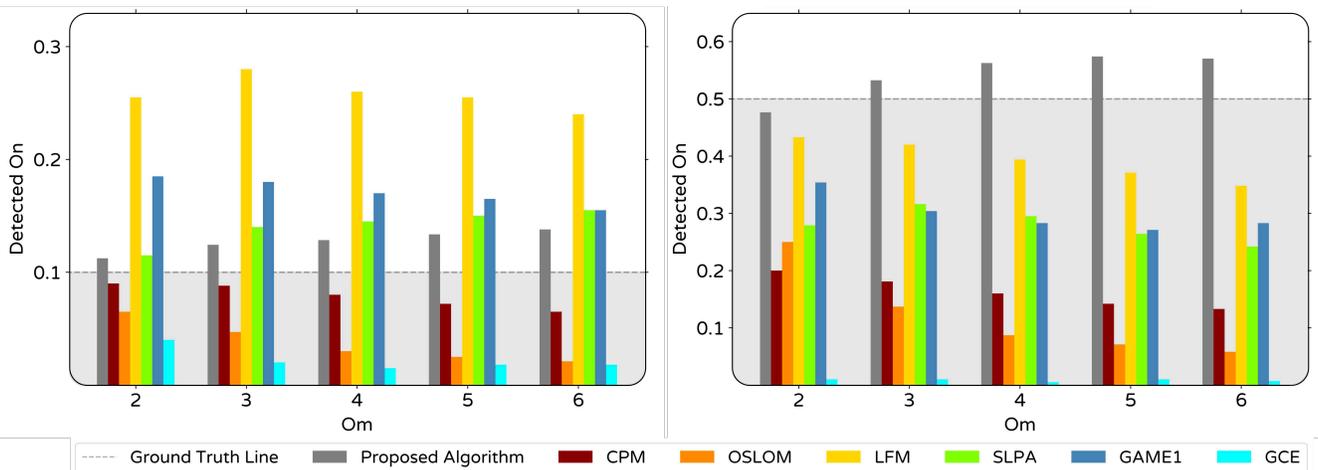


Figure 7. Overlappin nodes fraction detected by proposed and other algorithms in LFR3 and LFR4.

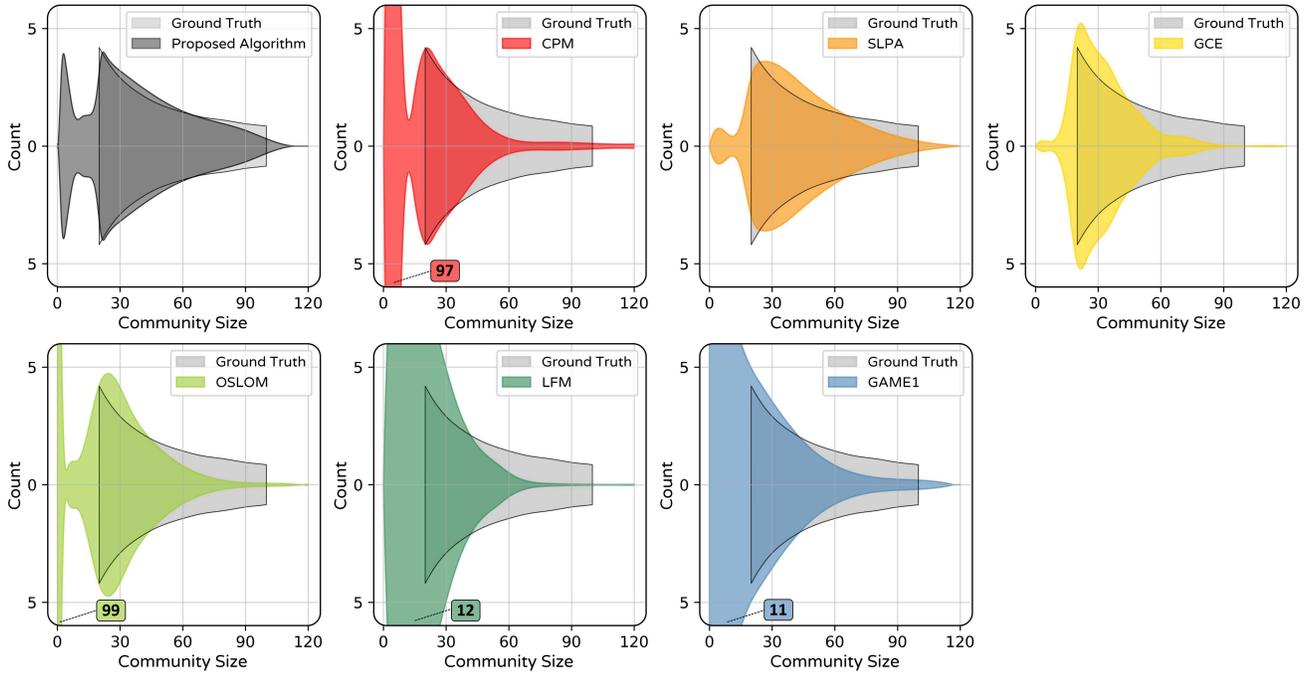


Figure 8. Histogram of detected community sizes for LFR3 (averaged on all Om). In each plot, the height of peaks is written next to them if they locate out of the frame.

when it comes to networks with overlapping communities, evaluation of a community detection algorithm performance must include checking the number of identified overlapping nodes which is one of important parameter determining the algorithm's accuracy. Overlapping nodes play crucial role in real-world social networks considering the fact they usually act as bridges or messengers between communities²⁷. Identified On detected by proposed and other algorithms for two groups of LFRs with ground truth O_n of 0.1 and 0.5 are shown in Figure 7. Overlapping nodes identified by our algorithm increase gradually by increase of Om . This trend is in contrast with other algorithms except SLPA in LFR3 network.

Aiming to finding more comprehensive insight on algorithms performance it would be beneficial to investigate distribution of detected community sizes (CS). For this purpose we used algorithms results on LFR3 averaging on all values Om and 10 instantiations of these networks. In histogram of community sizes which is shown in Figure 8, small fluctuations were omitted by representing fitted curves instead of raw data. for comparison, the ground truth power law distribution is visible in each histogram. Except for ours and SLPA algorithms, other algorithms have remarkable weakness in detecting larger size communities. Besides, some algorithms tend to break communities into smaller parts that cause distribution concentration in the range of small communities which do not exist in real distribution. Although such miss clustering occurs to some extent by our algorithm, it is not as much as some other algorithms such as GAME1, LFM, and especially CPM and OSLOM. Particularly, results demonstrate the relatively better performance of our algorithm in detecting larger communities.

Real Networks

	n	\bar{k}
Karate	34	4.5
Dolphins	62	5.1
Lesmis	77	6.6
Polbooks	105	8.4
Football	115	10.6

Figure 9. real networks in test.

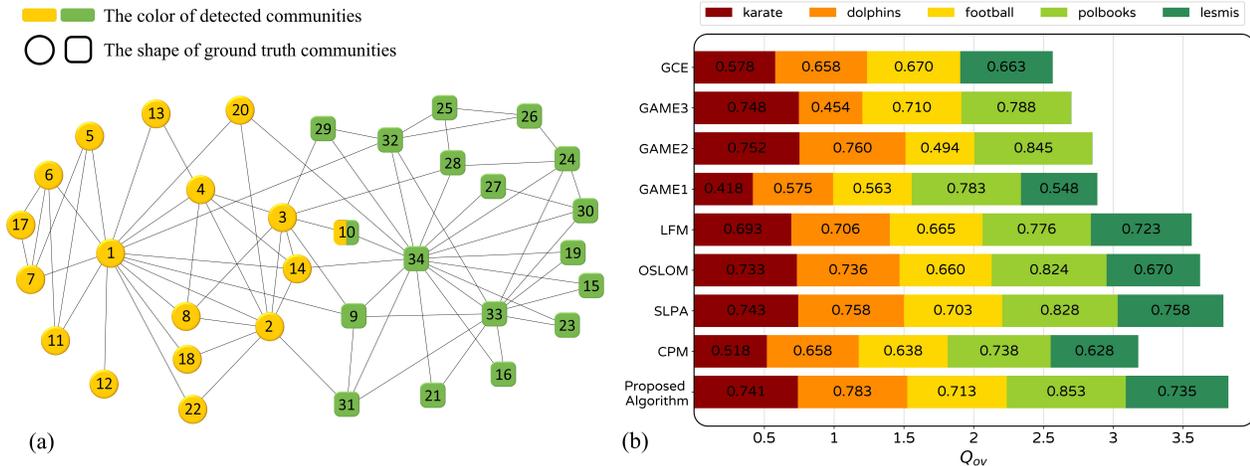


Figure 10. Ground truth and detected community structure of karate network. b) The Q_{ov} value obtained by proposed and other algorithm on five real networks.

In order to further evaluation of the proposed algorithm, we tested its performance on some real-world networks. As an evaluation measure, the overlapping modularity described in section 4.1 is used. Five real networks have been chosen for this test which the description of them can be observed in Figure 9.

Stack bar chart of Q_{ov} for obtained community structure of these five networks by ours and other algorithms is shown in Figure 10. Such illustration makes us able to compare the overall performance of algorithms on all five networks. our algorithm gets Q_{ov} value for Dolphins, Football and Polbooks which is slightly higher than other algorithms. Moreover, sum of Q_{ov} obtained by our algorithm is higher than others.

As an example, the community structure of karate network which is obtained by our algorithm is shown in Figure 10. This network is of traditional importance and was studied by Wayne W. Zachary during three years from 1970 to 1972³⁰. The ground truth of this network that was observed by Zachary, contains two communities representing in Figure 10. As it can be seen, detected community structure is exactly fitted to ground truth if excluding node 10. However, locating node 10 in the overlapping of two communities is sensible considering its equal connection with both.

Conclusion

In this paper, we proposed a novel game theoretic-based Algorithm for community detection in networks. Algorithm performance test on synthetic and real-world networks indicates our algorithm has a relatively better performance compared with similar algorithms presented in the literature. Our proposed algorithm has a time complexity of $O(m)$ making it a good choice for applying on ultra-large networks. Besides, no stochastic factors are influencing the process of community detection, which eliminates the need for multiple executions and averaging of results and causes our algorithm to be categorized among stable ones. In addition, this framework can be straightforwardly applied to weighted networks by making minor changes.

data availability

All data generated or analyzed during this study are included in this published article. The proposed algorithm python code is available in the Supplementary Material.

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Author contributions statement

F. Ferdowsi and K. Aghababaei Samani contributed in algorithm development and result analysis. F. Ferdowsi wrote the code and conducted the experiments, generated the figures, and wrote the manuscript. All authors reviewed and edited the manuscript.

competing interests

The authors declare no competing interests.