

Article

MINC-NRL: An Information-Based Approach for Community Detection

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Abstract: Complex networks usually consist of dense-connected cliques, which are defined as communities. A community structure is a reflection of the local characteristics existing in the network topology, this makes community detection become an important research field to reveal the internal structural characteristics of networks. In this article, an information-based community detection approach MINC-NRL is proposed, which can be applied to both overlapping and non-overlapping community detection. MINC-NRL introduces network representation learning (NRL) to represent the target network as vectors, then generates a community evolution process based on these vectors to reduce the search space, and finally, finds the best community partition in this process using mutual information between network and communities (MINC). Experiments on real-world and synthetic data sets verifies the effectiveness of the approach in community detection, both on non-overlapping and overlapping tasks.

Keywords: community detection; mutual information; network representation learning



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1. Introduction

Complex networks are a kind of graph-structured data which abstractly represents the real-world systems and structures. Nodes and edges of a network respectively represent elements and relationships of a system to show its topological characteristics. It has been shown that there are clusters of nodes that can be regarded as an independent whole according to some precisely defined and quantifiable attributes generally exist in many networks. Typically, the density of edges between nodes in a cluster. A group of nodes can be defined as a community if it has a higher density of internal edges than the average edge density of the whole network. Detecting communities in complex networks plays a vital role in understanding the structure and functions of the entire network system and can help us analyze and predict the interactions between the elements of it. Many researchers have focused on methods that can efficiently detect community structure in complex networks.

Traditional community detection algorithms often use modularity [1] to measure the strength and weakness of a community partition of a network; then, by using some process, such as the top-down split GN algorithm [2] and LFM algorithm [3], the maximum or extreme value of community evaluation index is reached. In this way, the best community partition of a network is found. Apart from modularity, some methods define and use other community evaluation index to detect communities more accurately, e.g., the IE [4] model, which aims to reveal the characteristics of the complex networks in an information theoretic view, by defining and calculating the information entropy of communities. Another information-theoretic index for evaluating partitions is the mutual information between network and communities (or MINC for short), proposed by L.C. Reidy et al. By defining structural information and community information between node pairs, the MINC approach specializes the general mutual information formula to calculating the mutual information between network structures and partitions in unweighted and undirected networks.

In recent years, with the development of representation learning algorithm in natural language processing (NLP) and other fields, researchers have begun to work on the relationship between network representation learning and community detection, such as using community annotations to improve network representation algorithms [5,6] and using network representation algorithms to improve the accuracy of community detection [6].

After induction and summary of these studies, we found that a typical community detection method often consists of two parts: The first is the community evaluation index, which measures the strength and weakness of a community structure; the second is the community transformer, which constantly change the community structure and tries to find a partition that reach some peak values of the evaluation index, that is, the solutions of community detection.

In the ideal case, the community evaluation index is able to evaluate the goodness of the community structure precisely, and the community transformer can efficiently search to give a set of partitions which cover the optimal partition. Unfortunately, for the former, since there is no accurate definition of the merits of community structure, researchers can only design community evaluation indices by means of edge density (e.g., modularity) or information-theoretic representation of node clusters (e.g., information entropy), so as to fit the real-world cases as well as possible. For the latter, community transformers have to make a trade-off between efficiency and the accuracy. Most of the methods have no way to use the global information of the network while transforming the community structure. For instance, while propagating labels in label propagation algorithm, to which a node propagates a label depends only on its local structural environment.

Aiming at the problems above, a new approach called MINC-NRL is proposed. The main idea is by using network representation learning (NRL) algorithms such as the Deepwalk method [7], the network is represented as vectors, which reflects the global information of the network. A community evolution process is then generated based on these vectors using hierarchically clustering to reduce the search space. Finally, by the usage of mutual information between network and communities (MINC), the best state in this community evolution process is found, as the final result of community detection.

The main contributions of this paper can be summarized as follows:

- Extend the original MINC evaluation index to make it possible to evaluate overlapping communities in unweighted and undirected networks;
- A community detection approach MINC-NRL is proposed based on network representation learning and MINC;
- Experiments are conducted on real-world data sets and synthetic data sets to verify the effect of the MINC-NRL approach.

The rest of this paper is organized as follows: Section 2 gives a short review of the community detection algorithms. Section 3 discusses how to detect communities using the MINC evaluation index and expand its definition to fit overlapping community detection tasks. In Section 4, the MINC-NRL method is introduced and explained in detail. Section 5 implements the method and verifies its accuracy by experiments. Section 6 is the summary and prospect of our research.

2. Related Work

Detecting community structure on different varieties of networks proves to be a hard task [8]. Various types of approaches for community detection have been proposed including modularity optimization, label propagation, spectral clustering, dynamic analysis, clique percolation, etc. [9].

Since Girvan and Newman pioneered the community evaluation index called modularity [1], modularity optimization is widely applied and studied for its general applicability to network topology which consist only of nodes and edges. GN, FN, and Louvain are typical modularity optimization algorithms. The main motivation of these algorithms is to find a maximum value of modularity on a given network through constantly transforming the community partition. For instance, GN initializes the network as whole community

and breaks it into smaller ones by deleting the edge with the greatest edge-betweenness step by step and tracking the maximum modularity value in the process; FN uses a greedy technique to merge small communities to large ones, following the direction that raises the value of modularity. Modularity provides a link-based criterion to evaluate the goodness of partitions of a network, but it is showed that the modularity values of ground-truth partition on a large number of data sets are usually not the maximum. In addition, modularity has a problem called resolution limit, which means, in some cases, modularity optimization algorithms unreasonably divide a community to smaller ones or combine communities into a large one [10]. Aiming at such problems, researchers have proposed other community evaluation indices as objective functions. By labeling the communities and nodes using two-level encoding, Rosvall et al. have proposed a metric which uses the average length of the description codes generated by random walkers to measure a partition [11]. The accuracy of this evaluation index is time-dependent. In order to obtain a more accurate evaluation, more iterations and more time are needed. R. Lambiotte et al. have proposed a stability index to find and judge which partition has a high degree of stability during the random walking through a network and use stability as the basis for community detection [12]. Stability is defined as a time-dependent function, and when the time parameter t changes, the stability index also changes, hence it is hard to determine the value of t to obtain the best result. L. C. Reidy proposed an information-theoretic index to evaluate the mutual information between network and communities (hereafter referred as MINC) [13]. The MINC approach specializes the general mutual information formula to calculating the mutual information between network structures and partitions, which offers a new thought on detecting community structure within an information-theoretic framework. In our approach, the idea of MINC as a community evaluation index is adopted and extended to figure out the best partition at different resolutions.

Community evaluation indices provide a quantitative manner to evaluate a given partition on a specific network. However, due to the huge number of ways to partition a network, it is impossible to find the maximum value of the evaluation indices by exhausting all possible partitions. Thus, most of community detection algorithms uses community transformers to search local optimal solutions of the evaluation indices by gradually changing the ownership of the nodes. These community transform techniques mainly include greedy techniques, simulated annealing, extremal optimization, spectral optimization, etc. [8]. For instance, local expansion methods such as LFM [3] and GCE [14] are typical greedy techniques for local optimization. The algorithms follow the idea that starting from core nodes as the initial communities, and then greedy include neighbor nodes that are likely to be in the same community until a local maximum of the evaluation index called fitness is reached, and then continue to search other communities one by one in the rest of the network. Such type of methods is often used for the detection of overlapping communities, but the local optimums as the final solutions usually have significant difference with the ground-truths.

Aiming at improving the accuracy of the results and reducing the convergence time while searching optimal partitions, recent research is trying to add some pre-processing steps before community transforming to collect more structural information from the network. For example, the community detection method based on positive/negative connections [15] runs a random walking process in the network and performs statistical analysis on the random-walking sequence. Then, the relationships of the nodes are evaluated as positive/negative for further detection. With a similar idea, the EdMot algorithm [16] uses a motif-based hypergraph of the target network to enhance the edge, and applies other state-of-the-art algorithms to partition the network. The DEMON [17] methods builds an EgoMinusEgo network from the original network, by combining the ego network extraction and the graph-vertex difference operation. In addition, based on an ego network, SONIC-MAN [18] use moderator nodes to integrate the local structural information in distributed online social networks. Such pre-processing steps are verified to be effective in improving the community detection performance of existing approaches.

In recent years, with the development of representation learning algorithms in areas such as natural language processing, researchers have adopted the technique on deep learning to community detection. The idea of these approaches, in general, is to compress the high-dimensional structural information of the network (e.g., an adjacency matrix) into a set of low-dimensional vectors. Such vectors are defined as a low-dimensional representation of the network, which brings two main advantages: (1) Distance and density of nodes can be easily defined based on their representations; (2) The compression of structural information greatly reduces the time and space costs on detecting communities. For example, the ComE algorithm constructs the “Community Detection–Community Representation–Node Representation” closed-loop framework [6] to optimize both the node embedding and community embedding. The model improves the accuracy of community detection on multiple real-world datasets, together with better results in node classification and graph visualization. Based on a two-level representation learning strategies, MemeRep [19] adopts a genetic framework to optimize the representation which preserves the topology structure of the network. It is shown that the algorithm is effective on community detection for that it can make full use of the modularity density to preserve communities. Different from improving the quality of representation learning, our research pays more attention on getting community partitions with high scalability and accuracy, and preserve the structural information of the network at the same time.

3. Mutual Information between Network and Communities

3.1. Original Definition

It is well known that the key part of community detection approach is to design an effective community evaluation index. A well-defined community evaluation index should satisfy the following: The better the given partition is, the higher its value is. Thus, when an optimal partition is given, the index reaches its maximum value. The mutual information between network and communities (or MINC for short) is a community evaluation index defined in a view of informatic theory. Mutual information is a measure of the mutual dependencies of two random variables, i.e., the amount of information that one random variable contains about the other. If the two variables refer to networks and communities, respectively, we can obtain a definition of the mutual information between network X and community partition C as follows:

$$MINC(X, C) = H(X) - H(X|C) \quad (1)$$

Note that X and C are variables related to the structure of the network and community ownership of nodes, which will be specifically defined in the following section. $H(X)$ denotes the entropy of network X , which can be defined as the number of bits on average to describe the network. $H(X|C)$ denotes the conditional entropy of X and C , which quantifies the amount of information needed to describe X given that the value of the known variable C . Based on the perspective of information, the following assertions are made:

- (1) $H(X)$ is determined only by the connectivity of nodes of network.
- (2) For a certain network, the better a given partition of it is, the less the amount of information is needed to describe the network. This results in a smaller value of $H(X|C)$ and a larger value of $MINC(X, C)$.

Therefore, $MINC$ can be used as a community evaluation index to quantify the merits of a partition.

The value of $H(X)$ and $H(X|C)$ can be calculated by:

$$\begin{aligned} H(X) &= - \sum_{x \in X} p(x) \log p(x) \\ &= -[p(X = 1) \log p(X = 1)] - [p(X = 0) \log p(X = 0)] \end{aligned} \quad (2)$$

$$\begin{aligned}
H(X|C) = & -[p(C=1) \cdot p(X=1|C=1) \log p(X=1|C=1)] \\
& -[p(C=1) \cdot p(X=0|C=1) \log p(X=0|C=1)] \\
& -[p(C=0) \cdot p(X=1|C=0) \log p(X=1|C=0)] \\
& -[p(C=0) \cdot p(X=0|C=0) \log p(X=0|C=0)]
\end{aligned} \tag{3}$$

The above equations are derived from the general definition of the mutual information. As for how to calculate the probability distributions (e.g., $p(C=1)$), further definition is needed, and here comes the core idea of the MINC approach.

First, we consider the random variables X and C . X is a variable related to the structure of the network, which can be defined at the micro level of each pair of nodes in the network:

$$X = \begin{cases} 1 & \text{if the nodes are adjacent} \\ 0 & \text{if the nodes are not adjacent} \end{cases}$$

In the same way, the variable for community partition C of the network can be defined as:

$$C = \begin{cases} 1 & \text{if the nodes are in the same community} \\ 0 & \text{if the nodes are not in the same community} \end{cases}$$

Then by considering the connectivity and community ownership of nodes, the probability distributions can be defined as:

$$p(X=1) = \frac{m}{t} \tag{4}$$

$$p(X=0) = 1 - \frac{m}{t} \tag{5}$$

$$p(C=1) = \frac{\pi}{t} \tag{6}$$

$$p(C=0) = 1 - \frac{\pi}{t} \tag{7}$$

$$p(X=1|C=1) = \frac{\theta}{\pi} \tag{8}$$

$$p(X=0|C=1) = 1 - \frac{\theta}{\pi} \tag{9}$$

$$p(X=1|C=0) = \frac{m-\theta}{t-\pi} \tag{10}$$

$$p(X=0|C=0) = 1 - \frac{m-\theta}{t-\pi} \tag{11}$$

In the above equations, t denotes the number of possible node pairs, which is constantly $\binom{n}{2}$ for a particular network with n nodes. m is the number of adjacent node pairs, which is equal to the number of edges. π is the number of node pairs in same community. θ is the number of adjacent node pairs in the same community.

By using the above definitions, the mutual information between a particular network and one of its partitions can be calculated. Based on the assertions that a better partition of a network will have a higher MINC value, MINC can be used as an index to evaluate the community structure and guide the community detection progress.

3.2. Extending the Definition of MINC for Overlapping Communities

The original definition of MINC is not compatible with overlapping communities. This is due to the fact that it divides the edges of a network into two types: intra-community edges and inter-community edge to calculate the probability distributions by counting the number of them. For instance, $p(X=1|C=1) = \frac{\theta}{\pi}$ denotes the probability that a pair of randomly selected nodes in the same community have an edge between them in network X . The value of this probability is equal to the number of intra-community edges divided by the combinatorial number of selecting a pair of nodes in the same community.

However, in the case of overlapping communities, an edge can be ambiguous, which can hardly be defined as being inside or outside the community. As shown in Figure 1, edge a is an internal edge in community A , and edge b is an inter-community edge between communities A and B . However, edge c can be considered either as an internal edge of community A or as an edge between communities A and B because the nodes at both ends of such an edge can either belong to the same community A or to two different communities A and B . This makes it difficult to count the number of edges and calculate the probability distributions in Equations (4)–(11).

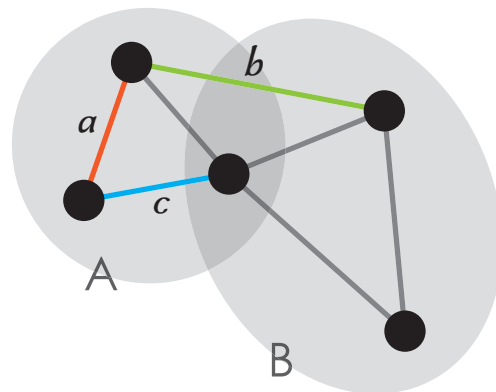


Figure 1. Ambiguous edges between two communities.

This problem can be solved by appending definitions for the ambiguous edges in overlapping communities. A possible solution is that for such an ambiguous edge, it is counted as 0.5 intra-community edges and 0.5 inter-community edges. However, such a solution is not reasonable in some more complicated cases. Consider edges m and n in Figure 2:

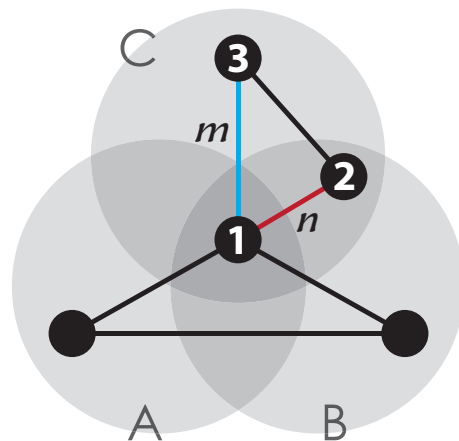


Figure 2. Ambiguous edges among three communities.

For edge m :

- In one case, it can be regarded as an internal edge (in community C);
- In two cases, it can be regarded as an inter-community edge (between communities A and C and between communities B and C).

For edge n :

- In two cases, it can be regarded as an internal edge (in community B and in community C);
- In three cases, it can be regarded as an inter-community edge (between communities A and B , between communities B and C , and between communities A and C).

The two edges m and n are in obviously different conditions, but if using the solution above, they will both be counted as 0.5 intra-community edges and 0.5 inter-community edges. Therefore, a more rigorous method is proposed to count the number of edges within and between communities, when there is overlap between communities. The number of edges in the same community is defined as:

$$\theta = \sum_{l \in M} p(l_a, l_b) \quad (12)$$

where l_a and l_b denote the ends of edge l , and M denotes the set of all the edges of the network. $p(r, s)$ is defined as:

$$p(r, s) = \frac{\sum_{R_i \in R} \sum_{S_j \in S} \delta_{R_i S_j}}{|R| \cdot |S|} \quad (13)$$

where R and S denote the community sets to which node r and s belong, respectively. R_i and S_j are the i -th and j -th communities in R and S , respectively. $\delta_{R_i S_j}$ is the Kronecker delta, which equals to 1 if $R_i = S_j$ and 0 otherwise. $|R|$ and $|S|$ are the numbers of communities in R and S , respectively.

For the number of node pairs π in same community, it can be calculated in the same way by replacing $l \in M$ with $l \in L$, where L denotes the edge set formed by selecting any pair of two nodes in the network:

$$\pi = \sum_{l \in L} p(l_a, l_b) \quad (14)$$

After appending the above definitions for counting the ambiguous edges in overlapping communities, the MINC index is able to be applied to overlapping community partitions.

4. Community Detection Based on MINC

4.1. Community Evolution Process

Apart from MINC as a community evaluation index, a community transformer is still needed to constantly transform the community structure, find the peak value of the evaluation index and finally carry out a solution of community detection. Here, we introduce the community evolution process, which will play a role as a community transformer in the MINC-NRL approach.

Definition 1. Community evolution process

A community evolution process P is an ordered set consist of a series of state, defined as:

$$P = \{P_0, P_1, P_2, \dots, P_N\}$$

where each state P_n is essentially a community partition of the same network.

Definition 2. Community evolution process of bottom-up cohesion

If adjacent states P_n and P_{n+1} in community evolution process P satisfy the following:

$$\forall Y_j \in P_{n+1}, \exists X_i (X_i \in P_n \rightarrow X_i \subseteq Y_j)$$

such process can be named as a community evolution process of bottom-up cohesion.

As a special kind of community evolution process, the community evolution process of bottom-up cohesion is used to represent the evolution process of constant merging of communities over time, as shown in Figure 3.

For a given network, the main idea of the approach is to generate a community evolution process as a community transformer, which simulates a real-world community evolution. That means in any state of such a process, nodes with closer distance or more similar structural environment will have a greater probability to be in the same community, and it brings three advantages comparing with other typical community transformer:

- (1) Each state of the generated process is itself a high-quality partition of the network, and there would be no unreasonable case where a node tries to pull adjacent nodes into its community without discrimination.
- (2) The number of states of the generated process is less, which reduces the calculation time to evaluate the merits of each state.
- (3) For such community evolution process, a stable and unique result can be obtained by finding the maximum of community evaluation index, or by filtering out the states with larger evaluation index, result partitions with different resolution can be output as well.

To generate a community evolution process for given network, network representation learning (NRL) and hierarchical clustering are used in the approach, which will be illustrated in detail in the next subsection.

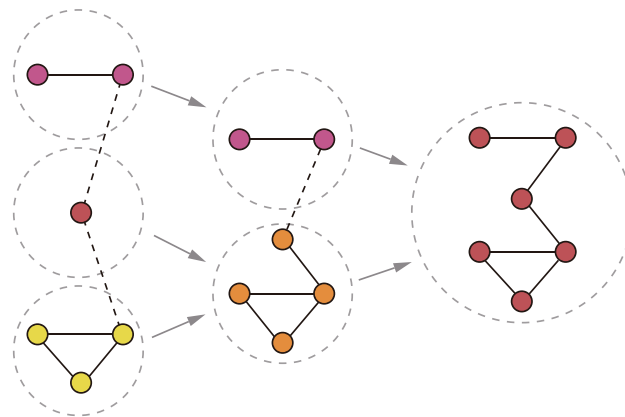


Figure 3. Community evolution process of bottom-up cohesion.

4.2. The Generation of the Community Evolution Process

4.2.1. Network Representation Learning

The generation of the community evolution process in the approach follows the idea that it simulates a real-world community evolution. To achieve this, the first step is sampling the network to vectorize the nodes, which preserves and reveals the structural information of the network. Such process is also known as network representation learning.

Multiple methods can be used to learn representations of nodes in network, e.g., neural-network method including Deepwalk [7], Node2Vec [20], BoostNE [21], Graph-Wave [22], and spectral method such as GLEE [23]. By comparing the results of these methods on our experimental datasets, Deepwalk is finally selected in our approach for its balanced and stable result on undirected real-world networks. The details of the comparative experiments are illustrated in Section 5.3.

4.2.2. Non-Overlapping Hierarchical Clustering

After obtaining the vector representation of the nodes, these vectors are then clustered using hierarchical clustering to form a community evolution process. For non-overlapping community detection, an agglomerative hierarchical clustering (AHC) is used to construct a bottom-up cohesion community evolution process.

The distance between two vectors is measured using the Euclidean distance, while that between two clusters is measured using Ward's method [24]. By clustering the vectors, the distance between each pair of vectors will be smaller when they represent a pair of nodes with closer distances or more similar environments in the network. When we alter the cluster distance threshold from small to large, the nodes will be first completely separated, and then the clusters merge in pairs based on the similarity between them, and eventually, the entire network is merged into one large cluster. Since these vector clusters are the representation of communities in the network, we have obtained a bottom-up cohesion community evolution process P in this way.

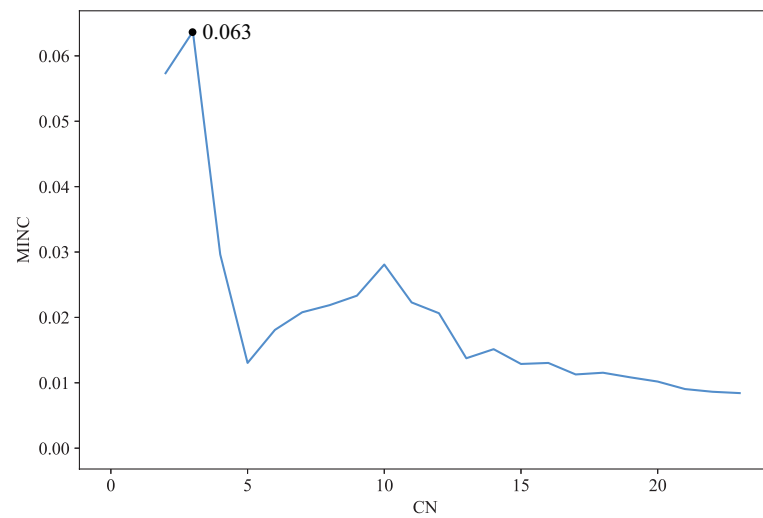


Figure 5. MINC value of community evolution process generated from Polbooks.

The pseudo-codes in Algorithm 1 gives a brief description of MINC-NRL.

Algorithm 1: MINC-NRL

Input: Network $G(E, V)$
Output: Communities detected in network $G(E, V)$

```

1 sequences  $\leftarrow$  List( $R \times N$ );
2 for  $r \leftarrow 0$  to  $R$  do
3   for  $n \leftarrow 0$  to  $N$  do
4     seq  $\leftarrow$  List();
5     currentNode  $\leftarrow n$ ;
6     seq.append(currentNode);
7     for  $l \leftarrow 0$  to  $L$  do
8       currentNode  $\leftarrow$  randomSelect(currentNode.neighbors);
9       seq.append(currentNode);
10    end
11    sequences.append(seq);
12  end
13 end
14 model  $\leftarrow$  Word2Vec(sequences, size  $\leftarrow$  neu_size, window  $\leftarrow$  window_size);
15 vectors  $\leftarrow$  model.get_embeddings();
16 if overlapping = True then
17   dendrogram  $\leftarrow$  OHC(vectors);
18 else
19   dendrogram  $\leftarrow$  AHC(vectors);
20 end
21 list_partition  $\leftarrow$  dendrogram.levels ;
22 list_MINC  $\leftarrow$  List( $N$ ) ;
23 for  $i \leftarrow 2$  to  $N$  do
24   list_MINC[i]  $\leftarrow$  list_partition[i] ;
25 end
26 opt_i  $\leftarrow$  index_of(max(list_MINC));
27 return list_partition[opt_i];

```

4.4. Time Complexity Analysis

The time complexity of this algorithm can be calculated in three parts:

- (1) **Network Representation Learning** Firstly, random walks are performed in the network. Starting from each node, the random walker needs to go L steps in the network, and such process will iterate R times. Hence the time complexity for random-walk is $O(nLR)$, where n is the number of nodes of the network. Then the sequences will be input into the Word2vec model. The model first uses a window of size W sliding through these sequences to count the co-occurrence frequency between nodes, which needs a time complexity of $O(nLRW)$. Meanwhile, the representation vectors for the nodes will be updated using a stochastic gradient descent process of E epochs in $O(E \log(n))$ time, with the acceleration of the hierarchical softmax. To sum up, the overall time complexity of the network representation learning part is $O(nLR) + O(nLRWE \log(n))$.
- (2) **Hierarchical Clustering** With the acceleration by a nearest neighbor chain, the agglomerative hierarchical clustering (AHC) needs a time complexity of $O(n^2)$. For the overlapping hierarchical clustering (OHC), in each iteration, or each level of the dendrogram, each cluster should traverse all nodes to decide which one is the closest, this leads to $O(Cn)$ time, where C is the number of clusters for each level. The number of levels of the dendrogram created by OHC is proportional to the number of nodes n . Thus, the total time complexity for OHC is $O(Cn^2)$.
- (3) **Find the Peak Value of MINC** For a partition, the main time cost for MINC calculation comes from obtaining θ and π , which need a traverse of all edges for their belonging community. To achieve this, a map is first built to store which community each node belongs to, with a time complexity of $O(n)$. Hence, $O(n + m)$ is needed, where m is the number of edges of the network. Since the dendrogram created by hierarchical clustering has $O(n)$ levels, and each level will be regarded as a partition for MINC calculation, the overall time complexity for this part is $O(n(n + m))$.

In conclusion, the time complexity for the approach is $O(nLR) + O(nLRWE \log(n)) + O(n^2) + O(n(n + m))$ for non-overlapping community detection and $O(nLR) + O(nLRWE \log(n)) + O(Cn^2) + O(n(n + m))$ for overlapping community detection. During the actual running of the approach, random-walk length L , iteration R , window size W , and update epochs E are much smaller than the number of nodes n and the number of edges m , the main time cost comes from the clustering part and MINC calculation part. For non-overlapping community detection, the total time complexity is $O(n^2) + O(n(n + m)) = O(n(n + m))$; for overlapping community detection, it is $O(Cn^2) + O(n(n + m))$. It should be noted that the number of clusters C for each level in OHC depends on the merging criterion λ in OHC algorithm. When the λ increases, clusters are more likely to merge, which decreases C . When setting $\lambda = 0.1$, the OHC algorithm tends to have a time cost of about $O(n^{2.45})$ [25]. Hence, the $O(Cn^2)$ time for OHC becomes the major bottleneck in computational efficiency of the approach in overlapping community detection tasks.

5. Experiments

In this section, experiments are conducted to confirm the effectiveness of MINC-NRL.

5.1. Preparation of the Experiment

To verify the accuracy of MINC-NRL, experiments are conducted on both non-overlapping and overlapping community detection tasks. For non-overlapping community detection, we include 4 non-overlapping community detection algorithms as baselines and use the number of communities (CN) and normalized mutual information (NMI) to verify the accuracy of MINC-NRL. For overlapping community detection, 3 overlapping community detection algorithms are included, and 4 accuracy evaluation indices are used, including CN, Overlapping Modularity (Q_{ov}), Extended Modularity (EQ), and Average Conductance (AC).

The data sets used in the experiment are all undirected and unweighted networks with ground-truth partition labels. Among them, the Karate Club [28], Dolphins [29], Football [30], and Polbooks [26] are real-world networks, and LFR500, LFR2000, LFR10000_a, and LFR10000_b are synthetic networks generated with scale-free features according to parameters [31]. Tables 1 and 2 list the main properties of data sets and the hardware and

system information of the experiments, respectively, where k is the average degree and μ is the mixed parameter of LFR networks.

Table 1. Main properties of data sets.

<i>Data Set</i>	<i>Node#</i>	<i>CN</i>	<i>k</i>	μ
Karate Club	34	2	4.6	-
Dolphins	62	2	5.1	-
Football	115	12	10.7	-
Polbooks	105	3	8.4	-
LFR500	500	3	15	0.3
LFR2000	2000	4	30	0.3
LFR10000_a	10,000	4	15	0.3
LFR10000_b	10,000	87	30	0.3

Table 2. Hardware and system information.

CPU	Intel(R) Core(TM) i5-9600K
Cores	8
Frequency	2.4 GHz
Memory	16 GB
Operating system	CentOS 7

5.2. Benchmarks

For non-overlapping community detection, since the data sets are partition-labeled, the number of communities (CN) and normalized mutual information (NMI) [32] are used to measure the accuracy of community detection. As a general rule, it is better that the CN of community detection result is exactly the same as labeled, or at least similar with labeled. NMI can be used to evaluate the difference between the result partition and labeled partition based on an information theory framework. The NMI value between partition A and B can be formulated as:

$$NMI(A, B) = \frac{-2 \sum_{i=1}^{C_A} \sum_{j=1}^{C_B} C_{ij} \cdot \log\left(\frac{C_{ij} \cdot N}{C_{i \cdot} \cdot C_{\cdot j}}\right)}{\sum_{i=1}^{C_A} C_{i \cdot} \cdot \log\left(\frac{C_{i \cdot}}{N}\right) + \sum_{j=1}^{C_B} C_{\cdot j} \cdot \log\left(\frac{C_{\cdot j}}{N}\right)} \quad (15)$$

where N is the total number of nodes, C is a confusion matrix, whose element C_{ij} denotes the number of nodes belonging to Community i in Partition A and also belonging to Community j in Partition B . C_A and C_B denote the number of communities of Partition A and B , respectively. $C_{i \cdot}$ and $C_{\cdot j}$ denote the sum of all elements of a row or column in Matrix C . The value of NMI is between 0 and 1, and it becomes larger when the two partitions are more similar. If the two partitions are exactly the same, the NMI value reaches 1. The NMI index is used in our experiments to compare the result partitions with the labeled partitions. A higher NMI value stands for a better result.

For overlapping community detection, we use overlapping modularity (Q_{ov}) [33], extended modularity (EQ) [34], and average conductance (AC) to evaluate the structural merits of the result partitions, instead of comparing the result partition with the labeled networks. Although we did not use the partition labels directly, we still compared the result CN with the labeled CN because the number of communities still has reference value on overlapping community networks.

The extended modularity (EQ) is a community quality index which extends the definition of Newman's modularity to overlapping community structures. The definition of EQ is shown as follows:

$$EQ = \frac{1}{2m} \sum_{c \in C} \sum_{i,j \in c} \frac{1}{O_i O_j} \left[A_{ij} - \frac{k_i k_j}{2m} \right] \quad (16)$$

where m is the total number of edges in the network, and c is one of the communities. i and j are nodes belonging to Community c ; k_i and k_j are their respective degrees. A_{ij} is the element of the adjacency matrix which follows the condition that when node i and j are linked, the value of A_{ij} is 1, otherwise 0. O_i and O_j are the numbers of communities to which node i and j belong, respectively, and give EQ the ability to deal with the case that a node belongs to more than one community. A high value of EQ indicates a significant overlapping community structure for a particular network.

The overlapping modularity index Q_{ov} is another quality function proposed by V. Nicosia et al. [33] to extend modularity to the more general case of overlapping communities, defined as:

$$Q_{ov} = \frac{1}{m} \sum_{c \in C} \sum_{i,j \in V} \left[\beta_{l(i,j),c} A_{ij} - \frac{\beta_{l(i,j)}^{out} k_i^{out} \beta_{l(i,j)}^{in} k_j^{in}}{m} \right] \quad (17)$$

Same as that in EQ , m is the total number of edges in the network. A_{ij} are the elements of the adjacency matrix. i, j are nodes of the network. k_i^{out} is the out-degree of Node i , while k_j^{in} is the in-degree of Node j . Instead of the number of communities to which a node belongs to, Q_{ov} uses the belonging coefficients $\beta_{l(i,j)}^{in}$ and $\beta_{l(i,j)}^{out}$ to calculate the weight for each link $l(i, j)$ existing between node i and j . Similar to EQ , a higher value of Q_{ov} indicates a stronger community structure.

Another index used in our experiment to evaluate the quality of overlapping community detection result is the average conductance (AC). Conductance is a local measure for the goodness of a node cluster in the network [35]. For a cluster c , conductance is defined as:

$$f(c) = \frac{m_c^{out}}{2m_c^{out} + m_c^{in}} \quad (18)$$

where m_c^{in} is the number of internal edges of Cluster c , m_c^{out} is the number of edges that links the cluster to other parts of the network. To evaluate a partition of the network, average conductance (AC) is used, which is defined as:

$$\Phi(C) = \text{avg}_{c \in C} f(c) \quad (19)$$

where C is a community partition of a network. The equation averages the conductance of all communities of the partition.

5.3. Preliminary Experiments: Comparison on Different Network Representation Methods

As a pre-step to hierarchical clustering, the network representation method can significantly affect the community evolution process generated. As mentioned in Section 4.2, comparative experiments are conducted to evaluate the impact of the network representation methods on the final results and find which of the network representation method is the most appropriate for our framework. The first comparison is made by replacing the network representation process of MINC-NRL (Step (1) and (2) in Section 4.3) by different network representation methods, including Deepwalk [7], Node2Vec [20], Walklets [36], RandNE [37], BoostNE [21], GLEE [23], NetMF [38], GraRep [22], NMFADMM [39], and verified with non-overlapping community detection tasks on real-world datasets.

The following table shows the NMI of using different network representation methods in MINC-NRL. The bold numbers emphasize the best experimental results within each data set.

As illustrated in Table 3, random-walk-based methods perform better than the other algorithms, including Deepwalk, Node2vec, and Walklets, and Deepwalk reaches the highest NMI on most of the datasets. Figure 6 shows the representation of Football network using Deepwalk, which is reduced to 2D by a principal component analysis (PCA). The results indicate that Deepwalk is more appropriate for MINC-NRL.

Table 3. NMI of using different network representation methods in MINC-NRL

	Karate	Dolphins	Football	Polbooks	Avg.
MINC-Deepwalk	1.000	1.000	0.924	0.589	0.878
MINC-Node2Vec	1.000	0.657	0.681	0.521	0.715
MINC-Walklets	0.557	0.659	0.702	0.574	0.623
MINC-Role2Vec	0.447	0.268	0.402	0.306	0.356
MINC-RandNE	0.523	0.333	0.764	0.576	0.549
MINC-BoostNE	0.448	0.676	0.781	0.598	0.626
MINC-GLEE	0.523	0.333	0.764	0.576	0.549
MINC-NetMF	0.350	0.297	0.728	0.593	0.492
MINC-GraRep	0.334	0.889	0.673	0.568	0.616
MINC-NMFADMM	0.390	0.271	0.418	0.220	0.325

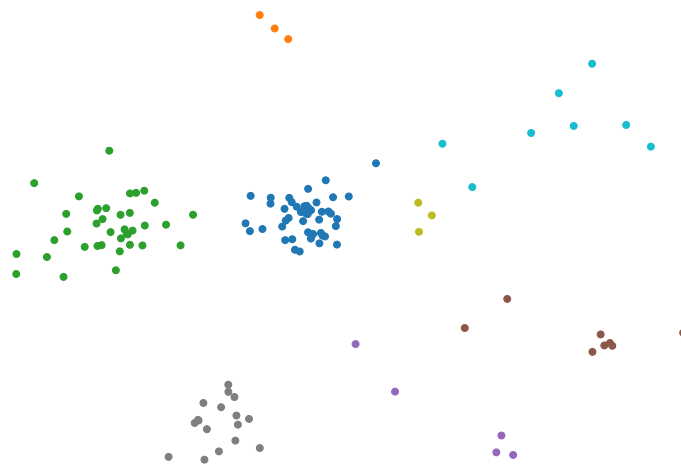


Figure 6. Network representation of Football using Deepwalk.

5.4. Comparison Results with the Other Algorithms

5.4.1. Non-Overlapping Community Detection

The following Tables 4 and 5 show the CN and NMI of running the MINC-NRL algorithm on the data sets, compared with non-overlapping community detection algorithms GN [2], FN [40], Louvain [41], and EdMot [16]. Note that the NMI values quantify the differences between the result partitions with the ground-truth partitions of the data sets.

Table 4. CN of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_a
MINC-NRL	2	2	12	3	3	4	4
GN	5	5	10	5	-	-	-
FN	2	3	7	3	3	8	-
Louvain	2	2	10	3	3	5	4
EdMot	4	5	12	5	3	5	5

Table 5. NMI of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_a	Avg.
MINC-NRL	1.000	1.000	0.924	0.589	0.978	1.000	0.850	0.906
GN	0.580	0.554	0.879	0.558	-	-	-	0.643
FN	0.914	0.606	0.762	0.534	0.969	0.635	-	0.706
Louvain	0.837	0.753	0.885	0.554	0.972	0.755	0.741	0.785
EdMot	0.587	0.511	0.851	0.504	0.980	0.823	0.799	0.722

The results show that MINC-NRL performs well on both real-world networks and LFR synthetic networks. The numbers of communities of the networks are correctly figured out by MINC-NRL. It also obtains the highest average NMIs on most of the data sets compared with the other algorithms.

5.4.2. Overlapping Community Detection

The following Tables 6–9 show the CN, Q_{ov} , AC, and EQ values, respectively, of the results of MINC-NRL algorithm on the data sets, compared with overlapping community detection algorithms ASLPaw [42], DEMON [17], and Ego-splitting [43]. Note that smaller values are better when evaluated by AC.

Table 6. CN of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_b
MINC-NRL	2	2	11	3	3	4	87
ASLPaw	2	3	6	5	1	2	87
DEMON	2	4	8	5	48	344	492
Ego-splitting	3	4	5	3	7	228	43,193

Table 7. Q_{ov} of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_b	Avg.
MINC-NRL	0.753	0.751	0.835	0.697	0.659	0.668	0.700	0.723
ASLPaw	0.739	0.780	0.818	0.741	-	0.460	0.699	0.605
DEMON	0.441	0.417	0.219	0.304	0.003	0.001	0.038	0.203
Ego-splitting	0.641	0.669	0.782	0.550	0.287	0.109	-	0.506

Table 8. EQ of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_b	Avg.
MINC-NRL	0.162	0.170	0.176	0.164	0.133	0.140	0.173	0.160
ASLPaw	0.158	0.165	0.173	0.162	-	0.135	0.173	0.138
DEMON	0.139	0.084	0.073	0.086	0.008	0.004	0.037	0.062
Ego-splitting	0.137	0.145	0.166	0.121	0.0641	0.041	-	0.112

Table 9. AC of MINC-NRL compared with other algorithms.

	Karate	Dolphins	Football	Polbooks	LFR500	LFR2000	LFR10000_b	Avg.
MINC-NRL	0.132	0.071	0.168	0.313	0.303	0.300	0.300	0.227
ASLPaw	0.152	0.175	0.305	0.255	-	0.299	0.307	0.249
DEMON	0.314	0.142	0.123	0.312	0.676	0.845	0.358	0.396
Ego-splitting	0.289	0.271	0.208	0.371	0.706	0.906	-	0.459

As illustrated in the above tables, the algorithm correctly figured out the number of communities on almost all networks. Although it does not get the best result on every data sets, but the average accuracy is the best compared with the other algorithms.

6. Conclusions

In this paper, an information-based approach for community detection MINC-NRL is proposed. The approach adopts network representation learning techniques to obtain the vectorial representation of each node of a network. Then, a community evolution process is generated with these vectors, through an agglomerative hierarchical clustering or overlapping hierarchical clustering, according to a non-overlapping/overlapping community detection task. Finally, the MINC index is used to figure out the optimum partition in the community evolution process. The experimental results show the effectiveness of MINC-NRL on both real-world and synthetic networks.

Due to the definition of MINC, a limitation of the current approach is that it can only be applied in unweighted and undirected networks. In the future, we will try to further extend the definition of MINC to address this limitation. The approach achieves a high degree of accuracy on 10,000 nodes LFR synthetic networks, but it is still difficult to be applied to larger-scale networks. This is mainly due to the time complexity bottleneck of the clustering algorithms. We will continue our research on improving the accuracy and time efficiency of the clustering part and try to extend the algorithm to detect communities on larger-scale networks.

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Abbreviations

The following abbreviations are used in this manuscript:

MINC	Mutual Information between Network and Community
NRL	Network Representation Learning
AHC	Agglomerative Hierarchical Clustering
OHC	Overlapping Hierarchical Clustering

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