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# A local community detection algorithm based on potential community exploration

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Local community detection aims to detect local communities that have expanded from the given node. Because of the convenience of obtaining the local information of the network and nearly linear time complexity, researchers have proposed many local community detection algorithms to discover the community structure of real-world networks and have obtained excellent results. Most existing local community detection algorithms expand from the given node to a community based on an expansion mechanism that can determine the membership of nodes. However, when determining the membership of neighboring nodes of a community, previous algorithms only considered the impact from the current community, but the impact from the potential communities around the node was neglected. As the name implies, a potential community is a community structure hidden in an unexplored network around a node. This paper gives the definition of potential communities of a node for the first time, that is, a series of connected components consisting of the node's neighbors that are in the unexplored network. We propose a three-stage local expansion algorithm, named *LCDPC*, that performs Local Community Detection based on Potential Community exploration. First, we search for a suitable node to replace the given node as the seed by calculating the node importance and the node similarity. Second, we form the initial community by combining the seed and its suitable potential community. Finally, the eligible nodes are selected by comparing the similarities between potential communities and the expanding community and nodes and adding them to the initial community for community expansion. The proposed algorithm is compared with eight state-of-the-art algorithms on both real-world networks and artificial networks, and the experimental results show that the performance of the proposed algorithm is better than that of the comparison algorithms and that the application of potential community exploration can help identify the community structure of networks.

## KEYWORDS

local community detection, seed selection, local expansion, potential community, node similarity

## 1 Introduction

In recent years, there have been many changes in people's lifestyles brought by the emergence of various kinds of complex networks in different domains, such as computer networks, social media networks, biological networks, and power system networks [1]. Research on complex networks, especially the community structure of complex networks, has received much attention in various fields and interdisciplinary subjects [2, 3]. In a real-world network, the node represents an entity and the edge represents the correlation among entities [4]. In

addition, edges in the same community are connected densely, and edges between communities are sparse in contrast [3, 5].

The detection and analysis of community structure are helpful to discover the interaction between people in social networks, the function of proteins in protein networks, and the research fields of scholars in academic cooperation networks. These can help people solve practical problems in society, such as personalized recommendations of products and information in the commercial field, technological breakthroughs in the medical field, and expert mining in the academic field.

Research on community detection aims to detect the community structure in complex networks quickly and accurately. Girvan *et al.* [6] proposed the classic Girvan–Newman (*GN*) algorithm based on the *Betweenness*, which denotes the number of shortest paths between two nodes. The *GN* algorithm is a hierarchical clustering algorithm based on global information. The basic idea of *GN* is to delete edges in the network with the maximum *Betweenness* relative to all source nodes continuously and then recalculate the *Betweenness* of remaining edges in the network relative to all source nodes all edges in the network are deleted.

Subsequently, various excellent community detection algorithms that depend on the global information of the network were proposed [7]. However, accessing the global information of a real-world network is sometimes impossible and in some cases, unnecessary [8]. On the one hand, for large-scale or dynamic real-world networks, it is difficult and time-consuming to obtain global information [9]. On the other hand, in some practical applications, one does not need the global information of the entire network but just needs to obtain local information from a specific node [10]. Different from the algorithms based on global information, the local community detection algorithm is capable of obtaining the local community based on only local information around the target node. Therefore, an increasing number of local community detection algorithms based on local information have emerged over time.

Most existing local community detection algorithms expand from the given node to a community based on an expansion mechanism that can determine the membership of nodes. The expansion mechanism of previous algorithms is generally based on the relationship between the community under expansion and its neighboring nodes. However, the communities existing in the undetected area of the network also impact these neighboring nodes. It is arbitrary and inefficient to determine the membership of neighboring nodes based only on the relationship between the community and its neighboring nodes while ignoring the relationship between the neighboring nodes and these communities in the undetected area of the network. In addition, ignoring the communities in the undetected area of the network means that the lack of topological information about the node leads to a decline in the accuracy of node membership.

Accordingly, this paper introduces the concept of the potential community which represents the potential community structure hidden in the unexplored network, to solve the problem above. We define the potential communities of a node as a set of connected components composed of node neighbors. The connected component is a set of nodes, where there is a path between each of the two nodes. The consideration of the potential community serves to provide the suspicious node with the topological information of its neighboring communities in the undetected area of the network. We propose a three-stage local community detection algorithm named *LCDPC*,

which performs Local Community Detection based on Potential Community exploration. The algorithmic process of *LCDPC* consists of three stages: the seed selection stage, the community initialization stage, and the community expansion stage. First, the seed selection searches for a suitable seed to replace the given node. Second, the community initialization process forms the initial community by combining the seed and its suitable potential community. Finally, the community expansion process adds eligible nodes to the initial community for community expansion. The main contributions of this paper are as follows.

- This paper gives the definition of the potential community. For the first time, the notion of potential community is applied to the process of node identification to increase the accuracy of the local community detection algorithm.
- In addition, we propose a three-stage local expansion algorithm based on potential community exploration, which performs seed selection, community initialization, and community expansion in order.
- Experimental results show that the application of potential community exploration can help identify the community structure of networks.

The rest of this paper is organized as follows. Section 2 reviews the basic theory and the related algorithms. The basic definitions and detailed description of the proposed algorithm are presented in Section 3. Experimental results are shown in Section 4. Section 5 concludes this paper and outlines future work.

## 2 Related works

A typical local community detection algorithm consists of two main processes: the seed selection process and community expansion process. The seed or the seed community generated by the seed selection process is the basis of the algorithm, which directly determines the quality of the result. The community expansion process takes the seed as the initial community and expands the community by optimizing the quality function or node similarity. This section outlines representative methods in terms of seed selection and community expansion as well as node centrality metrics and quality functions.

### 2.1 Seed selection

To obtain high-quality communities, the seed selection method is applied to detect the most appropriate node of the target community that contains the given node as the seed. In recent years, scholars have proposed a variety of local community detection algorithms based on local expansion. Lancichinetti *et al.* [11] developed an algorithm that takes random nodes as the seed. Although this method has a fast running time, it reduces the quality of the resulting community, and the test is unstable. Baumes *et al.* [12] proposed a robust algorithm, named IC (iterative scan), which takes a random edge as the seed. However, this method is time-consuming because it produces a great many duplicate communities when searching for seeds. Lee *et al.* [13] proposed a method named *GCE* (greedy clique expansion), which takes a *k-clique* as the seed. Whang *et al.* [14] proposed a new seeding

strategy based on the same distance kernel and degree. Taking maximal cliques as the seed, *Li et al.* [15] introduced a deep and broad searching method to detect maximal cliques, where different communities are allowed to be merged into a larger subgraph according to some given rules. *Li et al.* [16] considered some eligible nodes as a seed community and expanded the seed community by absorbing adjacent nodes of the seed community based on the absorbing degree function. *Zareie et al.* [17] introduced a hierarchical community detection method, which proposed two indices, the topological location of a node and the closeness to the network graph core, to measure the influence of node and rank them based on these two indices. To solve the seed-dependent problem, *Ding et al.* [18] introduced a core detecting method to replace the seed with the core member of the target community. *Mohammadi et al.* [19] developed a new node ranking strategy based on nodes' global potential values, such as influence and importance, to reveal the core of the community. *Cheng et al.* [20] proposed an algorithm that performs the *TOPSIS* (Technique for Order of Preference by Similarity to Ideal Solution) to rank each node and take the node with the highest score as the seed. *Rezaei et al.* [21] proposed a non-heuristic algorithm *EML* (Extended Machine Learning-based vital node identification), which makes use of the vitality of a part of a network for training a *SVR* model and predicts the vitality of each node based on this trained *SVR*.

## 2.2 Community expansion

The community expansion method takes a seed or a seed community as the initial community and expands it by absorbing the appropriate neighboring nodes iteratively. There are two common ways to determine the fitness of a node: running a greedy optimization process for a quality function [22–26] and spreading the influence of the seed throughout the network [27–32]. The quality function evaluates the quality of the community partition, and the scores generated by it can be used for partition ranking [2]. In a study on 13 quality functions based on 230 large real-world social, collaboration, and information networks, *Yang et al.* distinguished quality functions from (1) only internal community connectivity, (2) only connectivity between internal nodes and external networks; (3) both internal and external community connectivity and (4) modularity [33].

Influence spreading is a method that spreads the seed influence throughout the entire network. Inspired by the epidemic spreading model, *Raghavan et al.* [34] proposed the classical *LPA* (Label Propagation algorithm). *LPA* initializes each node in the network with a unique label and propagates the label throughout the network. The method stops when the node label does not change. *Gregory et al.* [35] proposed an improved *LPA* algorithm, named *COPRA* (Community Overlap Propagation Algorithm), which allows overlapping by multiple labels on each node. *Tang* [36] proposed an algorithm based on speaker-listener label propagation to detect overlapping nodes.

In recent years, some other excellent expansion methods have been proposed. *Tao et al.* [37] proposed a method based on local similarity, which expands the community by connecting the node of a small-scale community with the nodes that have a high degree. *Li et al.* [38] proposed an algorithm to detect overlapping communities by diffusing the local spectral which is simulated by short random walks.

Based on the idea that walkers who follow the same node sets should be assigned to the same community, *Makoto et al.* [39] proposed a retrained random-walk similarity algorithm for community expansion. Taking advantage of *NGC* (Nearest Greater Centrality) nodes, *Luo et al.* [40] proposed a novel community expansion method that adds the node that has the largest fuzzy relation with its *NGC* nodes into the local community. *Bahadori et al.* [41] proposed a probabilistic overlapping community detection method called *PODCD* which addresses dynamic communities efficiently. *PODDCD* scans communities by a probabilistic generative model which is constructed by a multi-objective optimization evolutionary-based method.

To enable readers to clearly understand recent achievements in the field of seed selection and community expansion, we summarize the characteristics, advantages, and disadvantages of the abovementioned methods in the following [Table 1](#).

## 3 Basic definitions and algorithms

### 3.1 Motivation

As mentioned in [Section 2](#), researchers have proposed many local expansion algorithms and made great progress in terms of seed selection and community expansion. However, there are still problems in local expansion algorithms when determining the membership of suspicious nodes. Previous algorithms only consider the relationship between the community and suspicious nodes while ignoring the relationship between the suspicious node and its potential communities.

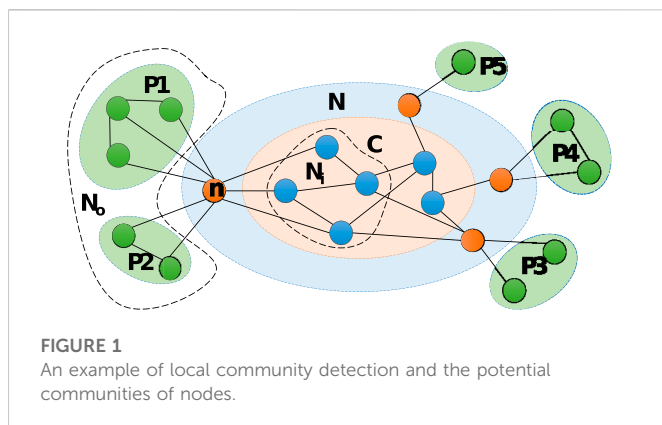
Let us take a brief figure legend of local community detection shown in [Figure 1](#) to illustrate this problem. In [Figure 1](#), subgraph *C* denotes the community under detection. Subgraph *N* denotes the neighboring nodes of *C*. The orange nodes in subgraph *N* denote the suspicious nodes of *C*. Subgraph  $P_{i(i \in \{1-5\})}$  denotes the potential communities of the suspicious nodes. The area  $N_o$  in the dotted line denotes the neighboring nodes outside *C* of suspicious node *n*. The area  $N_i$  in the dotted line denotes the neighboring nodes inside *C* of suspicious node *n*.

Node *n* in [Figure 1](#) is a suspicious node of community *C* that is under detection. The neighboring nodes of *n* can be divided into two parts: subgraph  $N_i$  located in community *C* and subgraph  $N_o$  located in the rest of the network. Most proposed algorithms determine the membership of suspicious node *n* on the basis of the relationship between  $N_i$  and *n*. Some algorithms take the relationship between  $N_o$  and node *n* into consideration, but they regard  $N_o$  as a whole. Actually,  $N_o$  is not an integrated whole, where nodes likely belong to different communities. As shown in [Figure 1](#),  $N_o$  is composed of  $P_1$  and  $P_2$ , which impact *n*. Therefore, treating adjacent nodes as a whole in undetected areas will lead to a reduction in the accuracy of node membership judgment in the process of community detection.

Topological structure refers to describing the entities and their relationships in complex networks by two basic graphic elements: nodes and links. The more topological structure information we have, the more precisely and rapidly we can detect the community structure from the complex network. Therefore, the motivation of this paper is to explore the topological structure information and subdivide the neighborhood of suspicious nodes. Therefore, we introduce the

TABLE 1 Pros and cons of common methods.

Methods	Characteristics	Advantages and disadvantages	Ref
Lancichinetti	Random selecting nodes as seeds	Low time complexity, but high randomness	[11]
Baumes	Random selecting edges as seeds	Low time complexity, but producing duplicate communities	[12]
GCE	Selecting $k$ -clique as seeds	Unable to address diversity of community	[13]
Whang	Seeding based on the same distance kernel	Needing big trainset size	[14]
Li	Taking maximal cliques as seeds	High time complexity	[15]
Mohammadi	Ranking nodes based on nodes' global potential values to reveal the core of a community	Strong adaptability	[19]
Cheng	Performing the TOPSIS to rank each node, and took the node with the highest score as seeds	High time complexity	[20]
Zareie	Ranking nodes with two indices, the topological location of a node and the closeness to the network graph core	Sufficient topological information	[17]
Ding	Searching the core of community as the alternative seed for the given node	Solving seed-dependent problem	[18]
Rezaei	Predicting the vitality of each node based on the trained SVR model	Strong adaptability, but relies on simulating the dynamics	[21]
LPA	Low time complexity	Unstability and high randomness	[34]
COPRA	An improved LPA algorithm allows overlapping by multiple labels on each node	Detecting overlapping communities	[36]
Tao	Expanding community by connecting the node in small scale community with the nodes own high degree	High time complexity	[37]
Li	Detecting the overlapping communities by diffusing the local spectral which is simulated by short random walks	Unstability and unpredictable	[38]
Makoto	Proposing a retrained random-walk similarity algorithm for community expansion	Unstability and unpredictable	[39]
Bahadori	Scanning communities by a probabilistic generative model which is constructed by a multi-objective optimization evolutionary-based method	Identifying dynamic communities efficiently	[41]



abstract concept of potential community into the local community detection algorithm. Detailed topological structure information can be obtained by exploring the potential communities of suspicious nodes to determine the membership of suspicious nodes accurately. As shown in Figure 1,  $P_1$  and  $P_2$  are the potential communities of node  $n$ . In our method, we first calculate the similarity between  $P_1$ ,  $P_2$ , and  $n$ . Then calculate the similarity between  $N_i$  and  $n$ . Finally, we choose the most similar one from  $P_1$ ,  $P_2$ , and  $n$  as the result community. Thus we can obtain a more accurate result on the basis of more topological structure information.

## 3.2 Problem definition

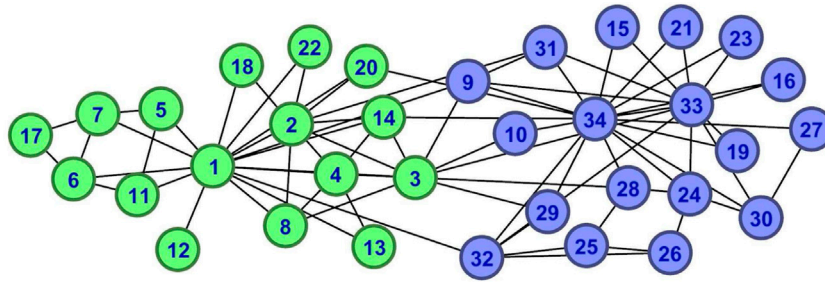
In this paper, we use a graph  $G = (V, E)$ , where  $V$  is the node set and  $E$  is the link set. The graph  $G$  can be represented as an adjacency matrix  $A$ , where  $A_{ij}$  denotes the connection of node  $i$  and node  $j$ , if node  $i$  and node  $j$  are connected,  $A_{ij} = 1$ ; otherwise  $A_{ij} = 0$ . The graph  $G$  consists of communities, where  $C = \{C_1, C_2, \dots, C_i\} (C_1 \cup C_2, \dots, \cup C_i \subseteq V)$ . A community can be represented as a node set  $C = \{v_1, v_2, \dots, v_j\} (C \in \mathbf{C}, v_i \in V)$ . The given node  $v_g$  is the initial node for local community detection, where  $v_g \in V$ . The target community  $C_{target}$  is a community where the given node is located in the real network, where  $C_{target} \in \mathbf{C}$ ,  $v_g \in C_{target}$ . The local community detection aims to detect the most similar community to the target community based on the given node.

## 3.3 Basic definitions

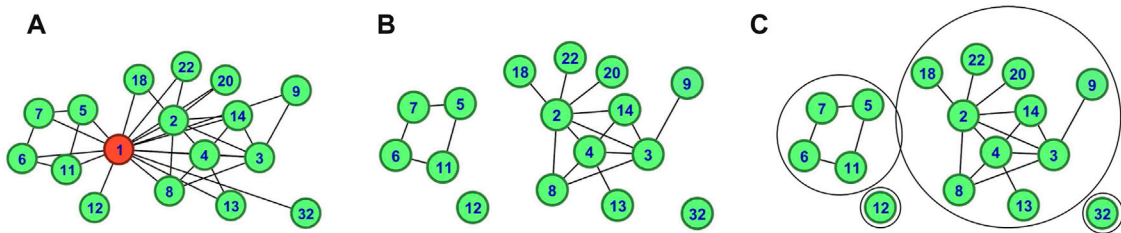
The definitions related to this paper are presented in this subsection.

**Definition 1.** (Neighboring nodes). The neighboring nodes  $N(v)$  of node  $v$  is defined as follows:

$$N(v) = \{u | (v, u) \in E, u \in V\}, v \in V \quad (1)$$



**FIGURE 2**  
The node distribution of the *Karate network*.



**FIGURE 3**  
An example of exploring the potential community of node  $v_1$ . (A) The distribution of node  $v_1$  and its neighbors. (B) Remove  $v_1$  and its links between its neighbors. (C) Potential communities of  $v_1$ .

where node  $v$  and node  $v_i$  are connected by a link.  $E$  is the link set and  $V$  is the node set of network  $G$ .

**Definition 2.** (Neighboring Community). The neighboring community  $\Gamma(v)$  of node  $v$  is defined as follows:

$$\Gamma(v) = N(v) \cup \{v\}, v \in V \tag{2}$$

The neighboring community of a node is the union of the node and its neighboring nodes. Figure 2 displays the nodes distribution of Karate network, and Figure 3 displays a part of Karate network. For instance, all nodes in Figure 3A make up the neighboring community of node  $v_1$ ,  $\Gamma(1) = \{1, 2, 3, 4, 5, 6, 7, 8, 9, 11, 12, 13, 14, 18, 20, 22, 32\}$ . Our algorithm takes the neighboring community of the seed node as the initial community for expansion.

**Definition 3.** (Community Neighbors). The community neighbors  $N(C)$  of community  $C$  is defined as follows:

$$N(C) = \{u|u \notin C, \exists v \in C, (u, v) \in E\}, C \in V \tag{3}$$

The community neighbors are neighboring nodes of community members, which are not in the community. Our algorithm takes the community neighbors of the detected community as suspicious nodes.

**Definition 4.** (Connected component). The connected component is defined as follows:

The connected component is a set of nodes, where there is a path between each pair of nodes. Nodes in the connected component are closely connected.

**Definition 5.** (Potential Community). The potential community of a node is defined as follows:

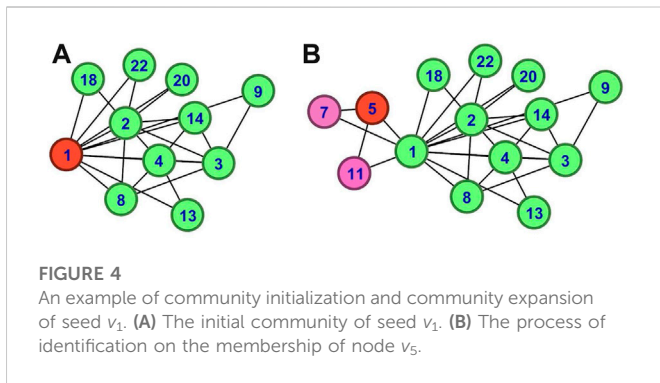
The potential communities of a node are a series of connected components that consist of the node’s neighbors. That is, every pair of nodes in the adjacent node is connected to each other.

We can search for the potential communities of node  $v$  by the following steps.

- Step 1: Set up a subgraph  $G_v$ , that is composed of node neighbors  $N(v)$  and links between them. That is, a tight node set formed by neighbors where each pair of nodes have a path, which is called the potential community.
- Step 2: Starting from one node in  $G_v$ , the proposed algorithm walks along the link and records the nodes encountered. When a path ends, it continues to walk along the previous branch until no branch can be continued. The nodes recorded constitute a connected component (The subnetwork with only one node is also considered to be a connected component.)
- Step 3: The connected component found in step 2 is one potential community of  $v$ , denoted by  $P(v)_i$ . Remove  $P(v)_i$  from  $G_v$ . The proposed algorithm iterates step 2 until there are no nodes left in  $G_v$ .

Note that  $\bigcup_{P(v)_i \in P(v)} P(v)_i = N(v)$ , and  $\forall P(v)_i, P(v)_j \in P(v), i \neq j, P(v)_i \cap P(v)_j = \emptyset$ , where  $P(v)$  represents the union of  $P(v)_i$ .

Figure 3 displays the process of exploring the potential communities of node  $v_1$  in the *Karate Club Network* [42].



**FIGURE 4**  
An example of community initialization and community expansion of seed  $v_1$ . (A) The initial community of seed  $v_1$ . (B) The process of identification on the membership of node  $v_5$ .

We use all neighboring nodes of node  $v_1$  to form a subgraph  $G_{v_1}$ . The connected components determined by the proposed algorithm are  $\{2, 3, 4, 5, 13, 14, 18, 20, 22\}$ ,  $\{5, 6, 7, 11\}$ ,  $\{12\}$  and  $\{32\}$ .

**Definition 6.** (Node Similarity). The similarity between a pair of nodes is defined as follows:

$$NS(v_i, v_j) = \frac{|\Gamma(v_j) \cap \Gamma(v_i)|}{|\Gamma(v_i) \cup \Gamma(v_j)|}, v_i \in V, v_j \in V \quad (4)$$

The Jaccard similarity coefficient [43] is a common measure used to compare the similarity between two finite sets. It is easy to compute and has linear time complexity. Therefore, we use the Jaccard similarity coefficient value of two nodes' neighboring communities to measure the similarity between the two nodes.

**Definition 7.** (Node Community Similarity). The similarity between community  $C$  and node  $v$  is defined as follows:

$$NCS(v, C) = |\Gamma(v) \cap C| \times \sum_{i,j \in (\Gamma(v) \cap C)} A_{ij} (d(v_i) + d(v_j)), v \in V, C \in V \quad (5)$$

where  $|\Gamma(v) \cap C|$  denotes the number of nodes in the intersection of the neighboring community of node  $v$  and community  $C$  and  $d(v)$  denotes the degree of node  $v$ .

We use the internal links in the intersection of the neighboring community of the node and community as the measurement of similarity between the node and community. To distinguish the situation in which there is the same number of links in the intersection, we set the degree of nodes on both sides of the link as the weight of the link. In addition, a larger scale of intersection means a higher similarity between the node and community. Therefore, we add the number of nodes in the intersection to the formula.

**Definition 8.** (Fittest Community). The fittest community  $FC$  to which node  $v$  belongs is defined as follows:

$$FC(v) = \arg \max (NCS(v, C)), C \in (\mathbf{P} \cup C_{detected}), v \in V \quad (6)$$

where  $\mathbf{P}$  is the potential community set of node  $v$ .

The fittest community to which a node belongs is the community that has the greatest similarity to the node among the potential communities of the node and the detected community. In the process of community expansion, nodes with the fittest community being the detected community will be added to the detected

community. In addition, when the detected community and the potential communities have the greatest similarity to the node at the same time, we choose the detected community as the fittest community.

### 3.4 The proposed algorithm

The proposed algorithm named *LCDPC* includes three stages: seed selection, community initialization, and community expansion. In order to give readers a clear description of the proposed algorithm, we show an example on *Karate Club Network* in Figures 3, 4.

Figure 3 shows an example of exploring the potential community of node  $v_1$ . As described in Definition.5, we first get the network composed of the node  $v_1$  and its neighboring nodes in Figure 3A. Second, we removed node  $v_1$  and its links which is shown in Figure 3B. Third, each circle in Figure 3C is a connected component, the potential community of node  $v_1$ .

When node  $v_1$  is the seed, the initial community of seed  $v_1$  is shown in Figure 4A. We can get the connected components of node  $v_1$  is  $\{v_2, v_3, v_4, v_8, v_9, v_{13}, v_{14}, v_{18}, v_{20}, v_{22}\}$  with similarity 3,784,  $\{v_5, v_6, v_7, v_{11}\}$  with similarity 450,  $\{v_{12}\}$  with similarity 30 and  $\{v_{32}\}$  with similarity 40 from Figure 3C. Therefore, we will form the initial community with node  $v_1$  and the connected component that is most similar to node  $v_1$ .

Figure 4B shows the process of identification on the membership of node  $v_5$ . Note that, the value of node degree is in the whole network rather than in the subgraph. First, we get the neighboring nodes of  $v_5$ , where  $\{v_1\}$  is neighbor inside the community and  $\{v_7, v_{11}\}$  are neighbors outside the community. Then, we explore the potential community of  $v_5$ ,  $\{v_7\}$  and  $\{v_{11}\}$ . The similarity between  $\{v_7\}$  and  $v_5$  is  $(3 + 4) * 2 = 14$  is bigger than the similarity between  $\{v_{11}\}$  and  $v_5$  is  $(3 + 3) * 2 = 12$ . So the external similarity is 14 The internal similarity between  $\{v_1\}$  and  $v_5$  is  $(16 + 3) * 2 = 38$  is bigger than 14. Therefore, node  $v_5$  is assigned to the community. If we perform the process without the application of potential community, the external neighboring nodes are considered as a whole  $\{v_7, v_{11}\}$ . The similarity between  $\{v_7, v_{11}\}$  and  $v_5$  is  $((3 + 4) + (3 + 3)) * 2 = 39$  which is bigger than internal similarity 38. Therefore, node  $v_5$  is not a member of the detected community.

There are four pseudo-code of *LCDPC* displaying in Algorithm 1, Algorithm 2, Algorithm 3, and Algorithm 4. The flow chart of each process is shown in Figure 5. The details of each procedure are shown in the following text.

*Potential community exploration.* Line three initializes a community  $P$  to the empty set to store the potential community. Line four initializes a *list* to store the nodes that make up the potential community. Then the first node from *list* is pulled out and  $v_{temp}$  is initialized as this node (Line 6). Lines eight to nine search for node sets that are linked in the common neighbors between the neighbors of  $v'$  and the neighbors of  $v_{temp}$  and store these nodes in  $P$  and *list*, respectively. Lines 5–10 repeat the algorithm until there are no nodes in *list*. Saving a potential community  $P$  obtained above to  $P_s$  (Line 11). Lines 2–12 repeat the algorithm until each node  $v_i$  neighbors  $v$ . Line 13 returns all the potential communities  $P_s$ .

*Seed selection.* The seed selection procedure aims to search for the most suitable node as the seed of the target community where the

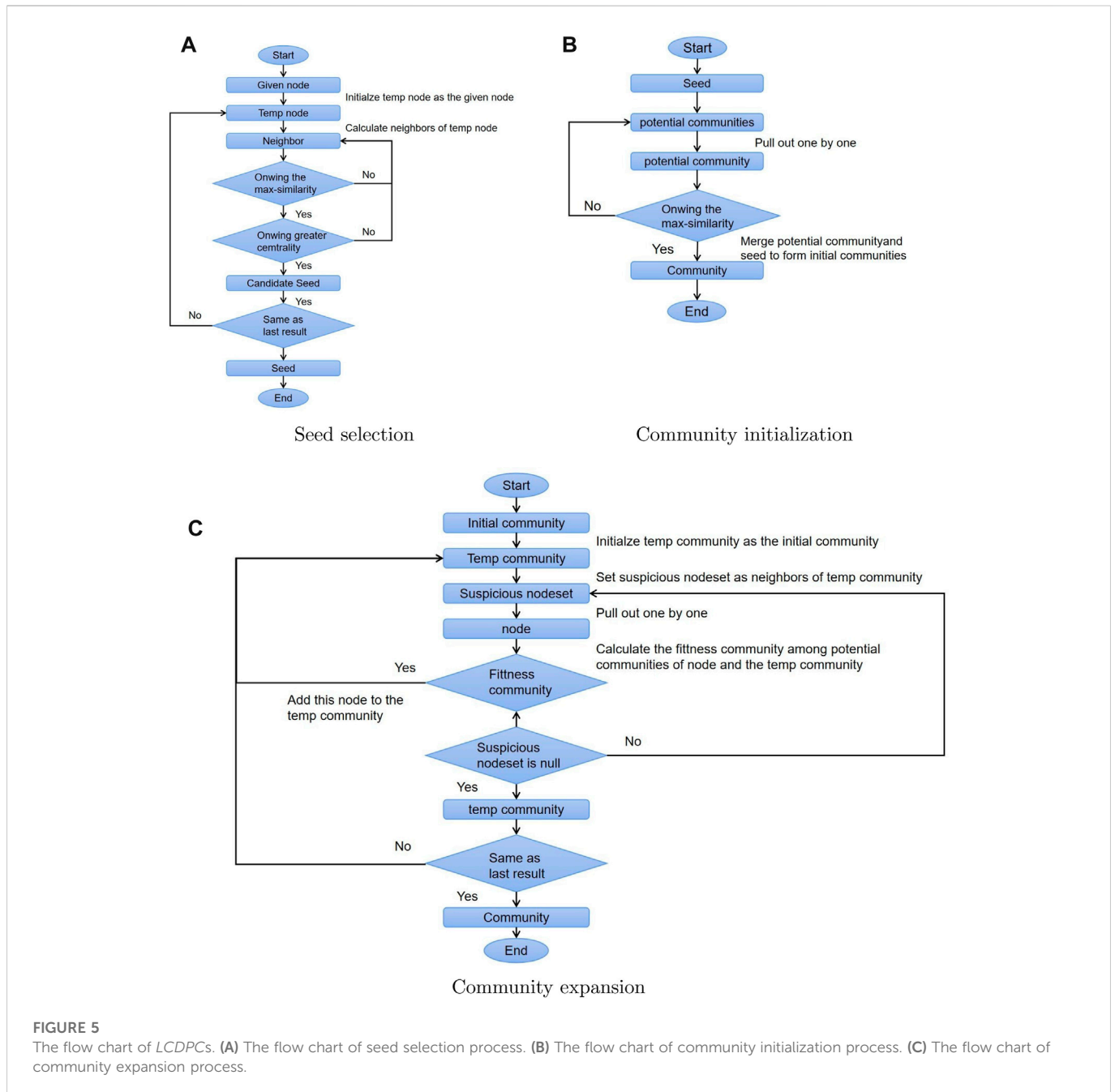


FIGURE 5

The flow chart of LCDPCs. (A) The flow chart of seed selection process. (B) The flow chart of community initialization process. (C) The flow chart of community expansion process.

given node is located. In the process of seed selection, LCDPC searches the node that meets the following two conditions: first, the degree of the node is greater than that of the given node; second, the node similarity between this node and the given node is the highest among neighboring nodes of the given node. In Algorithm 2, Line one initializes a community to the empty set to store the result community after the community expansion procedure. Line two calculates the degree of each node in the graph, which is the measurement of node similarity. Line seven calculates the neighboring nodes  $N(v_{temp})$ . Line 10 calculates the similarity between the seed and each node in the neighboring nodes of the seed. Lines 8–17 search the seed based on the two conditions among  $N(v_{temp})$ . The seed will be replaced iteratively by the node searched by the program above until no node meets the conditions (Lines 5–18).

**Community initialization.** The community initialization procedure generates an initial community based on the seed. The initial community consists of the seed and the potential community of the seed that has the highest similarity to the seed. In Algorithm 3, Line three calculates the potential community sets  $P(v_{seed})$  of the seed based on Definition 5. Line five calculates the similarity between the seed and each potential community in  $P(v_{seed})$ . The seed and the potential community with the highest similarity to the seed form the initial community (Lines 4–10).

**Community expansion.** The community expansion procedure adds eligible suspicious nodes to the initial community to form the resulting community. The eligible suspicious nodes should satisfy the condition that the fittest community of the node is the detected

community expanded from the initial community. In **Algorithm 4**, the initial community generated by the community initialization procedure is assigned to the temporary community (Line 1). Line two initializes a list of suspicious nodes to the empty set. Line five obtains the community neighbors  $N(C_{temp})$  of community  $C_{temp}$  based on **Definition.3**. Line one assigns  $N(C_{temp})$  to the list of suspicious nodes. For each node in the list of suspicious nodes, *LCDPC* gets the potential community set (Line 9) and calculates the fittest community (Line 10). If the fittest community is the detected community, the node will be added to the detected community (Lines 11–12), and Line 13 adds the neighboring nodes that are outside the detected community to the list of suspicious nodes. The program stops when there is no suspicious node meeting the condition, which means that the community does not change anymore (Line 16).

**Input:** Graph  $G = \{V, E\}$ , link set  $E$ , node set  $V$ , node  $v$ .

**Output:** Potential communities  $Ps$ .

**Process:**

- 1: Calculate neighboring nodes  $N(v)$  of  $v$  based on **Definition.1**;
- 2: **for all**  $v_i \in N(v)$  **do**
- 3: Initialize  $P = \emptyset$
- 4: Initialize  $list = v_i$
- 5: **do**
- 6: Initialize  $v_{temp} =$  pull out the first node from list
- 7: Calculate neighboring nodes  $N(v_{temp})$  of  $v_{temp}$  based on **Definition.1**;
- 8:  $P = P \cup (N(v) \cap N(v_{temp}))$
- 9:  $list = list \cup (N(v) \cap N(v_{temp}))$
- 10: **while**  $list \neq \emptyset$
- 11: Add  $P$  to  $Ps$
- 12: **end for**
- 13: return  $Ps$

#### Algorithm 1. Potential communities exploration.

**Input:** Graph  $G = \{V, E\}$ , link set  $E$ , node set  $V$ , seed node  $v_{seed}$ .

**Output:** candidate seed  $v_{seed}$ .

**Process:**

- 1: Initialize a community  $C$ ,  $C = \emptyset$ ;
- 2: Calculate the degree  $d_{v_i}$  of each node  $v_i \in V$ ;
- 3: Set  $v_{temp} = v_{seed}$ ;
- 4: Set  $max\_similarity = 0$ ;
- 5: **do**
- 6:  $v_{seed} = v_{temp}$ ;
- 7: Calculate neighboring nodes  $N(v_{temp})$  of  $v_{temp}$  based on **Definition.1**;
- 8: **for all**  $v_i \in N(v_{temp})$  **do**
- 9: **if**  $d_{v_i} > d_{v_{temp}}$  **then**
- 10: Calculate the node similarity  $NS(v_i, v_{temp})$  between  $v_i$  and
- 11:  $v_{temp}$  based on **Definition.6**;
- 12: **if**  $NS(v_i, v_{temp}) > max\_similarity$  **then**
- 13:  $v_{temp} = v_i$ ;
- 14:  $max\_similarity = NS(v_i, v_{temp})$ ;

- 15: **end if**
- 16: **end if**
- 17: **end for**
- 18: **while**  $v_{temp} \neq v_{seed}$
- 19: return  $v_{seed}$

#### Algorithm 2. Seed selection.

**Input:** Graph  $G = \{V, E\}$ , link set  $E$ , node set  $V$ , seed node  $v_{seed}$ .

**Output:** Initial Community  $C_{initial}$ .

**Process:**

- 1: Set  $similarity\_max = 0$ ;
- 2: Calculate the neighboring community  $N(v_{seed})$  of  $v_{seed}$  based on **Definition.2**;
- 3: Calculate the potential community set  $P(v_{seed})$  based on **Definition.5**;
- 4: **for all**  $P(v_{seed})_i \in P(v_{seed})$  **do**
- 5: Calculate the node community similarity  $NCS(v_{seed}, P(v_{seed})_i)$  between  $P(v_{seed})_i$  and  $v_{seed}$  based on **Definition.6**;
- 6: **if**  $NCS(v_{seed}, P(v_{seed})_i) > similarity\_max$  **then**
- 7:  $C_{initial} = P(v_{seed})_i \cup v_{seed}$ ;
- 8:  $similarity\_max = NCS(v_{seed}, P(v_{seed})_i)$ ;
- 9: **end if**
- 10: **end for**
- 11: return  $C_{initial}$

#### Algorithm 3. Community initialization.

**Input:** Graph  $G = \{V, E\}$ , link set  $E$ , node set  $V$ , Initial Community  $C_{initial}$ .

**Output:** Community  $C$ .

**Process:**

- 1: Set  $C_{temp} = C_{initial}$ ;
- 2: Set  $suspicious\_list = \emptyset$ ;
- 3: **do**
- 4:  $C = C_{temp}$ ;
- 5: Get the community neighbors  $N(C_{temp})$  of community  $C_{temp}$  based on **Definition.3**
- 6:  $suspicious\_list = N(C_{temp})$ ;
- 7: **while**  $suspicious\_list \neq \emptyset$  **do**
- 8: Pull  $v_i$  from  $suspicious\_list$ ;
- 9: Get the potential community set  $P(v_i)$  based on **Definition.5**
- 10: Calculate the fittest community  $FC(v_i)$  of  $v_i$  based on **Definition.8**
- 11: **if**  $FC(v_i) = C_{temp}$  **then**
- 12: Add node  $v_i$  to  $C_{temp}$ ;
- 13: update  $suspicious\_list = N(v_i) - C_{temp}$ ;
- 14: **end if**
- 15: **end while**
- 16: **while**  $C \neq C_{temp}$
- 17: return  $C$

#### Algorithm 4. Community expansion.



### 3.4.1 Time complexity analysis

We analyze the time complexity of *LCDPC* on a network  $G$  with an average degree of  $\bar{d}$  in the following. The analysis is performed in three steps of our algorithm.

The first step is to execute the [Algorithm 2](#) to find candidate core members of the seed node; this step requires  $O(\bar{d}^4)$  [18]. In the second step, we initialize the detected community based on the output of [Algorithm 2](#), as shown in the first phase of [Algorithm 3](#). Detecting all suspicious nodes requires  $O(\bar{s})$  where  $\bar{s}$  is the mean size of the potential community. Then [Algorithm 4](#) is executed to identify each suspicious node; this algorithm takes  $O(\bar{d}^3)$ . Accordingly, the time complexity of step 2 is  $O(\bar{s}\bar{d}^3)$ . The last step is to expand the community from the initial community. This requires  $O(r\bar{s}\bar{d}^3)$  where  $r$  is the maximum length of the path from the seed node to the community fringe. Finally, the overall time complexity of our method is  $O(\bar{d}^4 + r\bar{s}\bar{d}^3)$ .

## 4 Experiments and analyses

All the proposed algorithms and the comparison algorithms in this paper are written in JAVA and run on a computer with Intel (R) Core (TM) i5-4590 CPU, 3.3 GHz, and 16 GB RAM.

### 4.1 Evaluation criteria

To verify the performance of the proposed algorithms, we use the following two common evaluation criteria of community detection: *NMI* [44] (the normalized mutual information) and the *F-measure* [45].

#### 4.1.1 Normalized mutual information

*Danon et al.* proposed normal mutual information (*NMI*) measure [44] based on information entropy to measure the similarity between real-world communities and detected communities. This measure defines a *confusion matrix*  $\mathbf{N}$  with the rows denoting real-world communities and the columns denoting detected communities. Element  $N_{ij}$  in matrix  $\mathbf{N}$  represent the numbers of nodes that exist in both community  $i$  and community  $j$  [44]. The formula of *NMI* is:

$$NMI(A, B) = \frac{-2 \sum_{i=1}^{c_A} \sum_{j=1}^{c_B} N_{ij} \log(N_{ij}N / N_i N_j)}{\sum_{i=1}^{c_A} N_i \log(N_i / N) + \sum_{j=1}^{c_B} N_j \log(N_j / N)} \quad (7)$$

where  $c_A$  denotes the number of real-world communities and  $c_B$  denotes the number of detected communities.  $N_i$  and  $N_j$  denote the sum of elements in row  $i$  and column  $j$  respectively [44].

*NMI* is used to assess the performance of algorithms in dividing communities. A good partition means a great *NMI* value. The extreme case is *NMI* value is one when the partition is correct absolutely.

#### 4.1.2 F-measure

*F-Measure* is a widely used evaluation criterion to assess the performance of community detection algorithms. *F-Measure* is defined as:

$$F = 2 \times \frac{\text{Precision} \times \text{Recall}}{\text{Precision} + \text{Recall}} \quad (8)$$

The recall and precision are as follows:

$$\text{Recall} = \frac{|C_G \cap C_D|}{|C_G|} \quad (9)$$

$$\text{Precision} = \frac{|C_G \cap C_D|}{|C_D|} \quad (10)$$

where  $C_G$  denotes the real-world community and  $C_D$  denotes the detected community,

*F - Measure* is the weighted harmonic average of *Recall* and *Precision*.

## 4.2 Datasets

### 4.2.1 Artificial networks

We used a set of artificial networks generated by *LFR* [46] (Lancichinetti Fortunato Radicchi) benchmark networks to test the performance of the proposed algorithms. *LFR* is a common method to generate artificial networks that have the properties of real-world networks. The topology of the generated artificial networks is controlled by the following parameters:  $\mu$  is a mixing parameter that is used to control the difficulty of revealing the community structure;  $|C|_{\min}$  is the minimum size of the community, and  $|C|_{\max}$  is the maximum size of the community;  $\bar{d}$  is the mean node degree and  $d_{\max}$  is the maximum node degree  $O_n$  is the number of overlapping nodes and  $O_m$  is the average number of node overlaps. The parameter settings of *LFR* benchmark networks are listed in [Table 2](#), where the expression  $[a: b: c]$  means the value of the parameter ranges from  $a$  to  $c$  with a span of  $b$ . As displayed in [Table 2](#), we generate three groups of benchmark networks: *LFR- $\mu$* , *LFR- $\alpha_{size}$* , *LFR- $\alpha_{degree}$* . *LFR- $\mu$*  aims to examine the performance of the algorithm with the change in community identifiability. *LFR- $\alpha_{size}$*  aims to examine the performance of the algorithm with the change of community size. *LFR- $\alpha_{degree}$*  aims to examine the performance of the algorithm with the change in node degree. To ensure the accuracy of the experiments, we generated 10 artificial networks for each group of parameters and took the average value as the result.

The symbols mentioned in this section and their descriptions are displayed in [Table 3](#).

### 4.2.2 Real-world networks

[Table 4](#) displays the characteristics of six widely used real-world networks involved in this paper. *Karate* is a network of a karate club [42]. The nodes of the network represent members of the club and the links between two nodes denote a friendship between the two members. *Dolphins* is a network of bottlenose dolphins living in New Zealand [47]. Each node represents a bottlenose dolphin and if two dolphins are in frequent contact, there will be a link between the nodes representing them. *Books* is a network of political books [48]. The nodes represent political books on the Amazon website, and there is a link between the nodes representing two books if they are often bought together. *Football* is a network of college football teams in America [6]. Each node of the network represents a college and the links indicate that two football teams which have played against each other. *Amazon* is a network of products on the Amazon website [33]. *LastFM* is a social network of music website users, where the nodes of the network denote users of LastFM and links denote mutual follower relationships between them [49].

TABLE 2 The parameter settings of LFR benchmark networks.

Networks	$n$	$\bar{d}$	$d_{\max}$	$ C _{\min}$	$ C _{\max}$	$\mu$	$O_n$	$O_m$
LFR- $\mu$	1,000	5	25	10	100	[0.1:0.1:0.8]	20	2
LFR- $\alpha_{\text{size}}$	1,000	5	25	$5 \times [1 : 1 : 8]$	$50 \times [1 : 1 : 8]$	0.1	20	2
LFR- $\alpha_{\text{degree}}$	1,000	[4:1:11]	$5 \times [4 : 1 : 11]$	10	100	0.1	20	2

TABLE 3 Symbols and descriptions.

Symbols	Descriptions (for network $G$ )
$n$	The number of nodes
$m$	The number of links
$\bar{d}$	The mean degree
$d_{\max}$	The maximum degree of node
$ C _{\min}$	The minimum size of the community
$ C _{\max}$	The maximum size of the community
$\overline{ C }$	The average size of the community
$\mu$	The mixing parameter
$O_n$	The number of overlapping nodes
$O_m$	The average number of node overlaps
$n_c$	The number of communities

### 4.3 Experimental settings

We name the proposed algorithm that performs the process of potential community exploration to be *LCDPC1* and the one that does not perform the process as *LCDPC2*. To verify the performance of the proposed algorithm, we compared it to eight state-of-the-art local community detection algorithms: *RTLCD* (a Robust Two-stage Local Community Detection algorithm) [18], *Clauset* [50], *LWP* (Luo, Wang, and Promislow) [51], *Chen* [52], *LS* (Link Similarity) [53], *VI* (Vertex Influence) [54], *LCD* (Local Community Detection based on Maximum Cliques) [55] and *LCDDCE* (Local Community Detection method on line graph through Degree Centrality and Expansion) [56].

*Ding et al.* [18] proposed a robust community detection algorithm *RTLCD* that consists of two stages: seed selection stage and community expansion stage. *RTLCD* searches the core member of the community

where the given node is located in the seed selection stage, which solves the seed-dependent problem. *RTLCD* expands from the core member to the local community based on the relationship strength between nodes and communities in the community expansion stage, which solves the seed-invalid problem.

Based on the concept of *Newman's* [57] modularity, *Clauset et al.* [50] introduced a local community quality function  $\Delta R$ , which can be expressed as the ratio of the links within the community to the links with one or more endpoints outside the community. The *Clauset* algorithm adds nodes that cause the maximum increment of the quality function  $\Delta R$  to the community.

Based on the *Clauset* algorithm, *LWP et al.* [51] proposed an improved quality function  $M$ , which can be expressed as the edges within the community divided by the number of edges between communities. The *LWP* algorithm expands the community by adding nodes that cause the maximum increment of  $M$  to the community and deletes nodes that cause the maximum increment of  $M$  but are separated from the community. Different from *Clauset*, *LWP* has definite termination criteria.

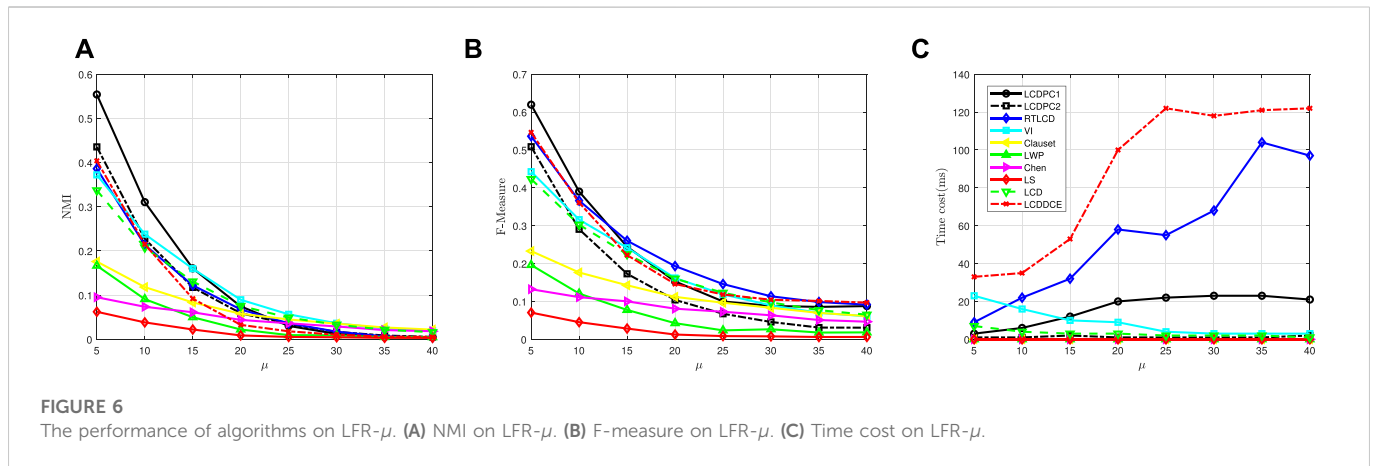
*Chen et al.* [52] introduced a novel quality function  $L$ , which takes the connection among nodes in the community and the connection between communities into consideration. To solve the outliers problem, the *Chen* algorithm checks for the changes in the quality function of the community after removing nodes.

*Wu et al.* [53] proposed a link similarity algorithm (*LS*) that can be defined as the intersection of a node's neighbors and the nodes and neighborhoods of the community. First, greedy optimization of link similarity is performed, which adds nodes with the largest similarity to the community. Second, whether the nodes on the boundary should continue to remain in the community is checked. Third, nodes that have more neighbors in the boundary than in the community are removed.

*Fanrong et al.* [55] took advantage of the maximum clique expansion and proposed the *LCD* algorithm, which takes the maximum clique as the seeds. *LCD* detects all the maximum cliques in the network as seeds and expands the community from these seeds according to greedy optimization until all the maximum cliques are assigned to communities.

TABLE 4 The characteristics of real-world networks.

Network	$n$	$m$	$\bar{d}$	$n_c$	$\overline{ C }$	$\mu$	$O_n$	$O_m$	References
<i>Karate</i>	34	156	4.58	2	17.00	0.128	0	--	[42]
<i>Dolphins</i>	62	318	5.12	2	31.00	0.038	0	--	[47]
<i>Football</i>	115	1,226	10.66	12	9.58	0.357	0	--	[6]
<i>Books</i>	105	440	8.38	3	35.0	0.159	0	--	[48]
<i>LastFM</i>	7,624	27806	7.29	18	423.56	0.126	0	--	[49]
<i>Amazon</i>	16716	97478	5.83	1,163	15.16	0.005	867	2.06	[33]



**FIGURE 6**  
The performance of algorithms on LFR- $\mu$ . (A) NMI on LFR- $\mu$ . (B) F-measure on LFR- $\mu$ . (C) Time cost on LFR- $\mu$ .

Yao et al. [54] proposed a variable influence local community detection algorithm (VI) based on a mechanism of influence attenuation, which is capable of detecting communities with variable scale according to the demands.

Wang et al. [56] introduced a line graph model to local community detection, and proposed an algorithm based on node centrality and PageRank. First, edges are transferred into nodes based on the line graph model. Second, nodes are ranked by a novel node similarity and PageRank and seeds are determined by this ranking. Third, the community is expanded by a fitness function.

In our experiments, all the algorithms were run in six real-world networks and three groups of LFR artificial networks, and the average of the experimental results was recorded.  $\alpha$  in LCDDDCE is set to 1.5. Note that all the algorithms that ran for more than 24 h were terminated. We take each node in the network as a given node and execute the algorithm with this given node. Finally, the results of all nodes are averaged as the performance of the algorithm on this network.

## 4.4 Experimental results on artificial networks

### 4.4.1 Experimental results on LFR- $\mu$

The purpose of LFR- $\mu$  is to examine the performance of algorithms with the change in community identifiability. Figures 6A, B display the NMI and F-Measure metrics for the proposed algorithms and the comparison algorithms on LFR- $\mu$ . From the pictures, we can observe that the trend of all results is downward with the increase of mix parameter  $\mu$ . The reason for this phenomenon is as follows. The definition of parameter  $\mu$  is the sharing ratio between the node and nodes in other communities; in other words, the parameter  $\mu$  represents the ratio between links outside the community and all links of the node. Therefore, as the parameter  $\mu$  increases, the community structure becomes more difficult to detect.

As shown in Figures 6A, B, when  $\mu \leq 0.5$ , except for Chen and LS, the results show a significant downward trend, and the performance of LCDPC has always been at a high level. Although the NMI and F-Measure metrics of Chen and LS are stable, they are always at a low level. When  $\mu > 0.5$ , the results of all algorithms are stable at a low level, because the mixed parameter  $\mu$  is so great that the community structure is very complex, and all algorithms cannot effectively detect the community structure.

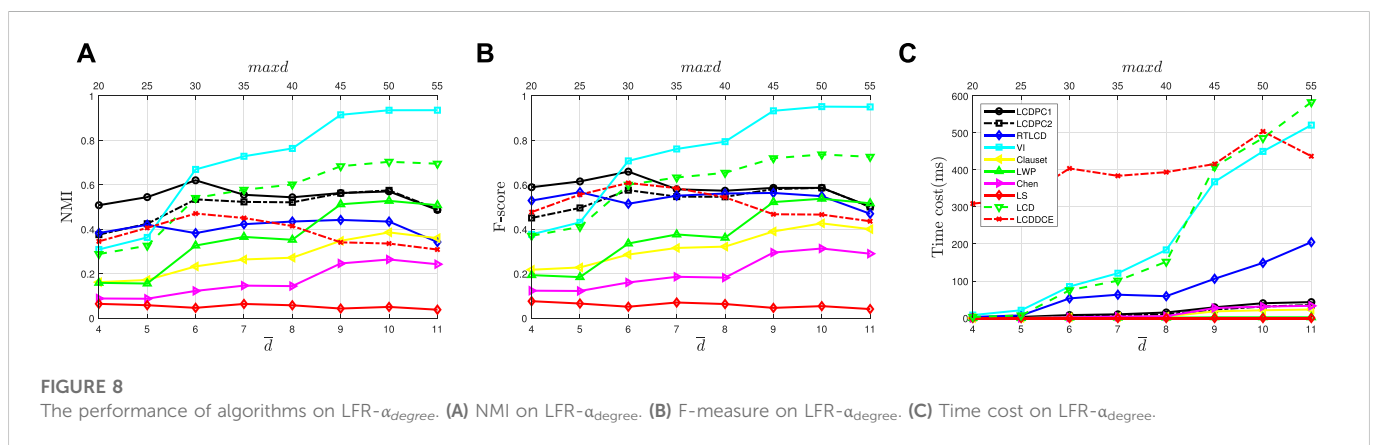
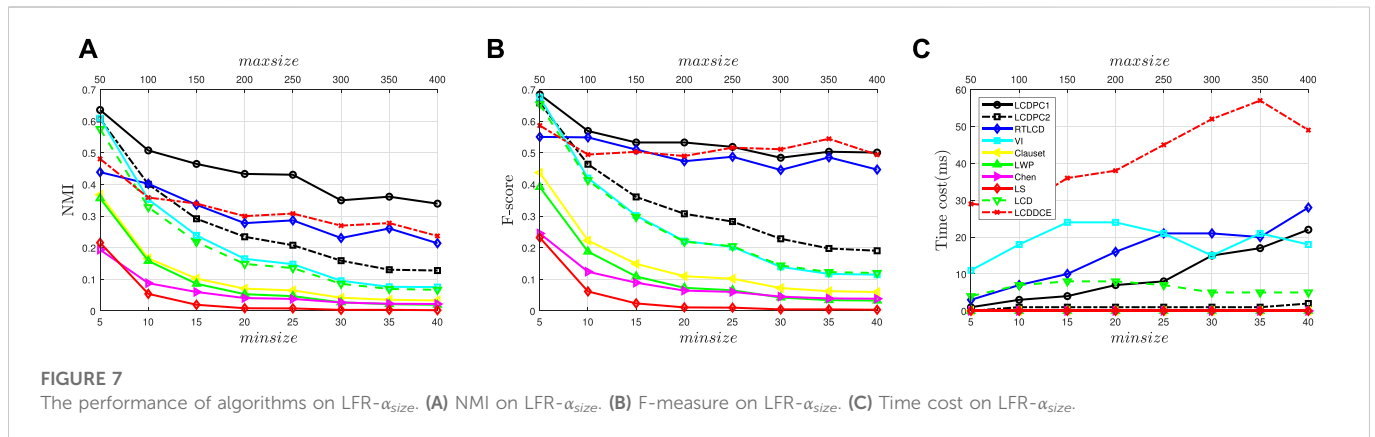
In addition, the performance of LCDPC1 is better than that of LCDPC2 on each value of parameter  $\mu$ . As the parameter  $\mu$  increases, the difference between LCDPC1 and LCDPC2 in performance decreases. The reason for this outcome is as follows. With the increase in the parameter  $\mu$ , the links between the node and the other communities increase. In this condition, it is easy for external adjacent nodes of the node to form the potential community. The scale of the external community formed by LCDPC1 and LCDPC2 is almost the same, which leads to a decrease in the gain effect of the application of potential community exploration. Therefore, the exploration of potential communities plays a better role in community detection in simple networks than in complex networks.

Figure 6C shows the time cost index of all the algorithms. We can observe that the majority of results remain stable at a low. RTLCD shows a significant upward trend with the growth of  $\mu$ . VI reaches a peak at  $\mu = 0.1$  and decreases gradually. LCDPC1 increases slightly and remains stable at  $\mu = 0.4$ .

### 4.4.2 Experimental results on LFR- $\alpha_{size}$

The purpose of LFR- $\alpha_{size}$  is to examine the performance of algorithms with the change of community size. Figures 7A, B display the NMI and F-Measure metrics for the proposed algorithms and the comparison algorithms on LFR- $\alpha_{size}$ . The top and bottom  $x$ -label in the graph represent the maximum and minimum community sizes, respectively. As seen from the figures above, the results of algorithms on LFR- $\alpha_{size}$  decrease with the increase in  $\alpha_{size}$ . This is because communities in the network become more diverse with the increase of parameter  $\alpha_{size}$ , which makes the boundaries of the community more difficult to identify.

From Figures 6A, B, we can observe that LCDPC1 outperforms the other algorithms on LFR- $\alpha_{size}$  and shows a smooth declining trend. When  $minc = 5$ , LCDPC2, VI and LCD perform as well as LCDPC1. However, when  $minc > 5$ , the performance of LCDPC2, VI and LCD decrease sharply. The reason for this result is as follows. As mentioned above, the community of the network becomes more diverse with the increase in the parameter  $\alpha_{size}$ . Therefore, the similarity between a neighboring community to a node becomes diverse. The application of potential community exploration helps LCDPC1 subdivide the similarity from a neighboring community to a node. The detailed similarity division makes LCDPC1 have a better effect than other algorithms. This explanation is confirmed in Figures 7A, B,



where the results gap between *LCDPC1* and *LCDPC2* becomes wider with the increase of the parameter  $\alpha_{size}$ .

### 4.4.3 Experimental results on LFR- $\alpha_{degree}$

The purpose of *LFR- $\alpha_{degree}$*  is to demonstrate the performance of algorithms with the change of node degrees. Figures 8A, B display the *NMI* and *F-Measure* metrics for the proposed algorithms and the comparison algorithms on *LFR- $\alpha_{degree}$* . The top and bottom  $x$ -label in the graph represent the maximum and mean values of node degree, respectively. From the figures above, we find that the efficiency in community detection of algorithms on *LFR- $\alpha_{size}$*  improves with the increase of parameter  $\alpha_{size}$ . The reason for this outcome is as follows. In the *LFR* network, the parameter  $\alpha_{degree}$  represents the node diversity. As the parameter  $\alpha_{degree}$  increases, the richer node diversity can bring more node information, which makes it easier for the algorithm to identify nodes.

From Figures 8A, B, we find that the performance of *VI*, *LCD* improves significantly with increasing  $\alpha_{degree}$ . However, as shown in Figure 8C, these algorithms consume much more time on node information processing with the increase of  $\alpha_{degree}$ . Furthermore, the difference between *LCDPC1* and *LCDPC2* in terms of the gain effect of potential community application, decreases with the increase in the parameter  $\alpha_{degree}$ . As mentioned above, with the increase of parameter  $\alpha_{degree}$ , nodes are easier to identify, which weakens the gain effect of potential communities application. Therefore, the application

of potential communities is effective in identifying nodes with poor degrees.

### 4.5 Experimental results on real-world networks

Table 5 displays *NMI*, *Recall*, *Precision*, *F-Measure* and Time metrics of the proposed algorithms with other comparison algorithms on five real-world networks. Table 6 lists the performance differences between *LCDPC1* and *LCDPC2* in terms of *NMI*, *Recall*, *Precision* and *F-Measure* metrics. From Table 5, we find that *LCDPC1* shows better average performance in detecting communities of real networks than the other comparison algorithms. This result shows that the proposed algorithm outperforms state-of-the-art local community detection algorithms in community identification.

From Table 6, we can observe that *LCDPC1* has improved in terms of *NMI*, *Recall*, *Precision* and *F-Measure* metrics compared to *LCDPC2*. This outcome verifies the effectiveness of the potential community application in community identification. Especially for the *Karate* and *Dolphins* networks, *LCDPC1* has significant improvement compared to *LCDPC2* in community identification. The common characteristic of the *Karate* and *Dolphins* networks is that they all have low mean degree values. Furthermore, we find that

**TABLE 5** The algorithm results on real-world networks. The maximum value in a criterion is marked in bold.

Network	Criteria	LCDPC1	LCDPC2	RTLCD	Clauaset	LWP	Chen	LS	VI	LCD	LCDDCE
Karate	NMI	0.9186	0.6815	<b>1.0000</b>	0.2992	0.5160	0.1552	0.1688	0.4041	0.4093	0.3635
	Recall	0.9722	0.8252	<b>1.0000</b>	0.5527	0.6912	0.2071	0.2339	0.6556	0.6182	0.8758
	Precision	0.9446	0.9338	<b>1.0000</b>	0.9088	0.8019	0.6345	0.5588	0.9014	0.8449	0.5958
	F-Measure	0.9580	0.8717	<b>1.0000</b>	0.6474	0.7179	0.2949	0.3171	0.7317	0.6918	0.7089
	Time(ms)	2	0	2	1	2	3	0	4	2	0
Dolphin	NMI	<b>0.4553</b>	0.2895	0.4526	0.1857	0.2809	0.0959	0.0709	0.2326	0.2616	0.4175
	Recall	0.6352	0.4831	0.6399	0.3013	0.3696	0.1517	0.0980	0.3717	0.3853	<b>0.6497</b>
	Precision	<b>0.9846</b>	0.9785	0.9647	0.9694	0.5271	0.7043	0.4032	0.9667	0.9546	0.8537
	F-Measure	<b>0.7365</b>	0.6274	0.7376	0.4287	0.4173	0.2364	0.1458	0.4985	0.5086	0.7266
	Time(ms)	2	0	9	1	0	2	0	4	1	0
Football	NMI	0.6269	0.6226	0.5146	0.5712	0.6023	0.5863	0.5714	<b>0.8389</b>	0.5638	0.0261
	Recall	0.8058	0.7921	<b>0.9209</b>	0.7133	0.6409	0.6665	0.5956	0.8907	0.7280	0.9506
	Precision	0.6896	0.6957	0.5568	0.6466	0.6257	0.6456	0.6461	<b>0.8963</b>	0.6354	0.0922
	F-Measure	0.7404	0.7379	0.6639	0.6689	0.6301	0.6479	0.6180	<b>0.8916</b>	0.6708	0.1673
	Time(ms)	4	3	15	3	0	5	0	4	3	0
Books	NMI	<b>0.4924</b>	0.5029	0.4881	0.2687	0.2925	0.0905	0.0106	0.3862	0.3594	0.4770
	Recall	0.8368	0.8044	0.8681	0.4387	0.4710	0.1532	0.0195	0.6256	0.6032	<b>0.8835</b>
	Precision	0.7579	0.7774	0.7049	0.7656	0.4643	0.5720	0.1705	<b>0.7778</b>	0.7554	0.7074
	F-Measure	<b>0.7851</b>	0.7822	0.7640	0.4982	0.4619	0.2219	0.0307	0.6398	0.6210	0.7766
	Time(ms)	4	3	33	8	3	6	0	58	25	
LastFM	NMI	<b>0.3716</b>	0.3301	0.3480	0.0189	0.0167	0.0079	0.0034	---	---	0.2176
	Recall	<b>0.5621</b>	0.4438	0.6471	0.0164	0.0168	0.0062	0.0026	---	---	0.4178
	Precision	<b>0.5967</b>	0.6366	0.5605	0.8183	0.1949	0.4015	0.2694	---	---	0.4469
	F-Score	<b>0.5466</b>	0.4978	0.5528	0.0279	0.0218	0.0109	0.0049	---	---	0.3968
	Time(ms)	883	563	8,717	19	8	53	0	---	---	5
Amazon	NMI	<b>0.7547</b>	0.7470	0.7254	0.5668	0.6261	0.4235	0.3918	0.6837	0.6888	0.7543
	Recall	0.7266	0.7183	0.6966	0.5192	0.5977	0.3772	0.3641	0.6499	0.6551	<b>0.7332</b>
	Precision	0.9929	0.9930	0.9914	0.9964	0.8783	0.8307	0.6776	0.9938	<b>0.9958</b>	0.9450
	F-Measure	<b>0.7861</b>	0.7784	0.7570	0.6138	0.6531	0.4638	0.4122	0.7157	0.7213	0.7852
	Time(ms)	2	2	0	1	0	1	0	10	4	16

**TABLE 6** The difference between LCDPC1 and LCDPC2.

Criteria	Karate	Dolphin	Football	Books	LastFM	Amazon
NMI	+0.2371	+0.1658	+0.0043	-0.0105	+0.0415	+0.0077
Recall	+0.1470	+0.1521	+0.0137	+0.0324	+0.1183	+0.0083
Precision	+0.0108	+0.0061	-0.0061	-0.0195	-0.0399	-0.0001
F-Measure	+0.0863	+0.1091	+0.0025	+0.0029	+0.0488	+0.0077

*LCDPC1* has limited improvement compared to *LCDPC2* in identifying communities on the *Football* and *Books* networks. Both the *Football* and *Books* networks have high mean degree values. These findings indicate that the potential community application has a more significant improvement in the efficiency of community identification on the network with a high mean node degree than that with a low mean node degree. In addition, *LCDPC1* has good improvement on the *LastFM* network and has limited improvement on the *Amazon* network compared to *LCDPC2*. The *LastFM* and *Amazon* networks have the same low mean node degree, but the difference is that the *LastFM* has a great average community size and the *Amazon* network is poor. This outcome verifies that the potential community application has a more significant improvement in identifying communities on the network with a greater average community size than those with a lesser average community size.

## 5 Conclusion

Local community detection is efficient in detecting local community structures based on seeds. However, the existing algorithms ignore the effect of community structure in undetected networks.

This paper introduces the abstract concept of the potential community into the local community detection algorithm to help determine the membership of suspicious nodes. We propose a three-stage algorithm based on potential community exploration, which performs seed selection, community initialization, and community expansion in order. First, the seed selection searches for a suitable seed to replace the given node. Second, the community initialization process forms the initial community by combining the seed and its suitable potential community. Finally, the community expansion process adds eligible nodes to the initial community for community expansion.

The proposed algorithm is compared to eight state-of-the-art local community detection algorithms on artificial networks and real-world networks. The experimental results show the following conclusions. First, the potential community application can improve the efficiency of community identification. Second, the improvement in potential community application on community identification is related to the

parameters  $\mu$ ,  $\alpha_{size}$ , and  $\alpha_{degree}$ . The improvement of potential community application increases with increasing  $\alpha_{size}$  and decreases with increasing  $\mu$  and  $\alpha_{degree}$ .

However, the experimental results show that the potential community application is not obviously better for some networks, especially networks with a low mean node degree. Therefore, in future work, we will improve the algorithm to expand the role of potential community applications.

## Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material further inquiries can be directed to the corresponding authors.

## Author contributions

SW write and revise the manuscript. JY, XD, JZ, and MZ contributed to paper ideas and data curation, analysis. All authors approved the submitted version.

## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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