

A simple and fast exact clustering algorithm defined for complex networks and based on the properties of primes

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Abstract

In this paper a new clustering method based on primes is proposed. This method define a nodes cluster of any complex network, considering the nodes with same input/output number and same number of paths with equal length, so all the network nodes with analogous functions will be possible to identify. The clustering algorithm proposed, results very efficient because it is defined on simple computations with primes. For example, with our algorithm the analysis of a network with 500 nodes and 124750 connections is performed in 80 seconds on Pentium 4 with CPU 2Ghz and 1Gb ram.

Keywords: Complex network, clustering method, graph theory, bidirectional network, complete path.

1. Introduction

In the last years the networks applications are frequently been used to study the complex systems. There are many types of complex network: Information Networks, Technological Networks, Biological Networks, Social Network etc, (M. E. J. Newman 2003, S. Broh e et al. 2008) but essentially they are all characterized by nodes and connections among the nodes. The nodes or vertexes are objects, with one or more input and one or more output, instead the connections define the information flow among the nodes, which can be bidirectional or unidirectional. The graph theory (F. Harary 1995, Bornholdt & Schuster 2002, Bondi J.A. & Murty U.S.R 2008) is the appropriate tool in the study and representation of complex networks. A bidirectional (unidirectional) network is represented with an *undirected (directed) graph* $G = (V, E)$ that consists of two sets V and E , such that $V \neq \emptyset$ and E is a set of unordered (ordered) pairs of elements of V . The elements of $V \equiv \{N_1, N_2, \dots, N_h\}$ are the *nodes* (or *vertices*, or *points*) of the graph G , instead the elements of $E \equiv \{e_1, e_2, \dots, e_k\}$ are its *connections* (or *links*, or *lines*, or *edges*). According with graph theory we consider a *path* from node i to node j is an alternating sequence of adjacent nodes and edges that begins with i and ends with j , in which each node is considered only once. Since the network nodes may represent proteins, cells, computers, web pages, individuals or animals etc, the individuation of nodes with analogous functions is important, so a simple methodology to define a exact clustering method for any bidirectional network, is proposed. This paper is organized as follows: in Section 2 we introduce the definitions of complete path and complete paths set, that are fundamentals to define the clustering algorithm. In Section 3, the new clustering method is described. In Section 4 there are the conclusions for this paper.

2. Complete Paths

We consider in Figure 1, the bidirectional complex network represented by the graph $G = (V, E)$, with $V \equiv \{A, B, C, D\}$ and unordered set $E \equiv \{AB, AD, BC, BD, CD\}$. We give the following definitions:

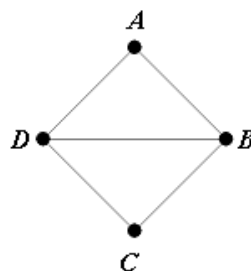


Figure 1. *Bidirectional Complex Network*

Definition 2.1. A *complete path (CP)* for a network node is a possible path which starts from the node and

include all the possible nodes network only once in a direction.

Definition 2.2. Two complete paths are equal if they will have same length or number of nodes, same nodes and same consecutive nodes without considering the verse.

Definition 2.3. Two complete paths are similar if they will have same length or nodes number, same nodes but they haven't same consecutive nodes, without considering the verse.

Example 2.1. The complete path ABCD (Figure 1) is equal to: BCDA, CDAB, DABC, and DCBA, CBAD, BADC, ADCB, instead it is similar to BACD.

Definition 2.4. The complete paths set (CPS) of a node V is the set $\wp_V = \{p_1, p_2, \dots\}$ of all complete paths p_i of a network, which start from V.

Remark 2.1. We want underline that a maximal path is complete path too, but no vice versa is true. For example we consider the path that start from B and ended to C (Figure 1). The maximal path is BADC instead the complete path are: BADC, BDC.

Example 2.2. In Figure 2 we show the complete paths set $\wp_B = \{BADC, BCDA, BDC, BDA\}$ of the network of Figure 1,

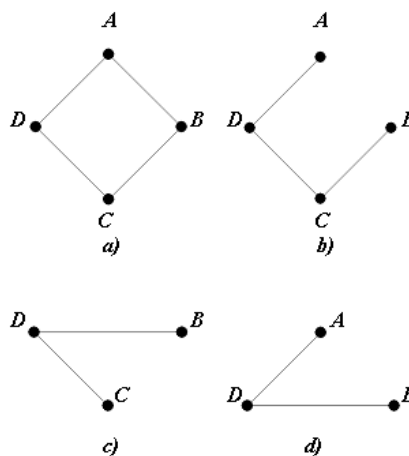


Figure 2: Complete paths set of the node B of the bidirectional complex network in Figure 1.

Considered \wp_H and \wp_K two CPS for H and K, if we indicate with $|\wp_H|$ and $|\wp_K|$ the number of complete paths of \wp_H and \wp_K respectively, we have the following definition:

Definition 2.5. \wp_H and \wp_K will be said equal, if $|\wp_H| = |\wp_K|$ and $\forall p_i$ (CP), if $p_i \in \wp_H$, than there is a $p_j \in \wp_K$ such a that p_j and p_i are equal and vice versa.

Example 2.3. We consider the CPS \wp_B and \wp_D of the nodes B and D (Figure 1),

$$\wp_B = \{BCDA, BACD, BDA, BDC\}$$

$$\wp_D = \{DABC, DCBA, DBC, DCA\}$$

for definition 2.5, they are equal.

Definition 2.6. Considered \wp_H and \wp_K the CPS of H and K, they will be said similar, if $|\wp_H| = |\wp_K|$ and $\forall p_i$ (CP), if $p_i \in \wp_H$, than there is $p_j \in \wp_K$ such a that p_j and p_i are similar and vice versa.

Remark 2.2. Two equal CPS are similar too, but two similar CPS are not equal.

Definition 2.7. Considered \wp_H and \wp_K the CPS of H and K, they will be said almost-similar, if they will have in common only some CP equal/similar.

Example 2.4. We consider the network in Figure 1 and the CPS \wp_A and \wp_B of the nodes A and B respectively:

$$\wp_A = \{ABCD, ADCB, ABDC, ADBC\}$$

$$\wp_B = \{BCDA, BACD, BDA, BDC\}$$

in this case, we observe that the first and second \wp_A -CP with the first \wp_B -CP are equals and the second \wp_B -CP is similar with the first and second \wp_A -CP, besides the set $\{ABDC, ADBC\} \not\subset \wp_B$ and the set $\{BDC, BDA\} \not\subset \wp_A$, therefore \wp_A and \wp_B are almost-similar.

3. Clustering Method

The clustering is a method that defines a number k of subsets separated of a initial set, according with an

opportune measure of likelihood or similarity. Such operation can be used for classifying the kind, to reassume results or still to analyze images, etc. Therefore the clustering represents in many cases an essential operation of statistic analysis of the experimental data. Further advances in this field are given by Anderberg, M. R. 1973, Hartigan, J. 1975, Dubes, R. C. et al. 1988, White D.R. 2001, Brades U. et al. 2003, Shamir R et al. 2004, Schaeffer S.E. 2007, Xing B. et al. 2007, Xu X. et al. 2007, Kim C. et al. 2008, Tan, L. et al. 2009, Gorke R. et al. 2010, Ahmed A. et al. 2012. The mathematical language that we have introduced, allows us to define a clustering algorithm. This algorithm divide in classes the nodes of a bidirectional complex network, according their *CPS*. To proceed to the clustering of the network nodes, for each *CPS* of a node will be encoded with an univocal integer. We will call this integer, *CPS-code*. Therefore it is possible to define the clusters for the network nodes, with equal *CPS-code* (or *CPS-code* of the same order of greatness), codifying the length of every *CP* of a node with a prime number (*CP-code*), according to the associations in Table 1.

Table 1. *CP-code for bidirectional complex network.*

| | |
|--|---------------------------------|
| $N_1N_2 = 2;$ | <i>CP-code</i> with two nodes |
| $N_1N_2N_3 = 3;$ | <i>CP-code</i> with three nodes |
| $N_1N_2N_3N_4 = 5$ | <i>CP-code</i> with four nodes |
| $N_1N_2N_3N_4N_5 = 7$ | |
| $N_1N_2N_3N_4N_5N_6 = 11;$ | |
| $N_1N_2N_3N_4N_5N_6N_7 = 13;$ | |
| $N_1N_2N_3N_4N_5N_6N_7N_8 = 17;$ | |
| $N_1N_2N_3N_4N_5N_6N_7N_8N_9 = 19;$ | |
| $N_1N_2N_3N_4N_5N_6N_7N_8N_9N_{10} = 23$ | <i>CP-code</i> with ten nodes |
| | |

with N_1, N_2, \dots the nodes of the generic *CP*. For example codifying all the *CP* of the node *A* (Figure 1), we will have:

$$\wp_A = \{ABCD, ADCB, ABDC, ADBC\}$$

CP-code(*ABCD*) = 5, *CP-code*(*ADCB*) = 5, *CP-code*(*ABDC*) = 5, *CP-code*(*ADBC*) = 5, i.e. we have four common *CP* of same length. so the \wp_A -code will be defined by:

$$\wp_A\text{-code} := \prod_{i=1}^{n=4} \text{CP-code } p_i \Rightarrow 5 \times 5 \times 5 \times 5 = 5^4 = 5 \tag{3.1}$$

where n is the number of the *CP* associated to the node *A*. The integer \wp_A -code will individualize a nodes cluster. Analogous for the nodes *B*, *C* and *D*, we have: $\wp_B = \{BCDA, BACD, BDA, BDC\}$, $\wp_C = \{CBAD, CBDA, CDAB, CDBA\}$, $\wp_D = \{DABC, DCBA, DBC, DCA\}$ with \wp_B -code = $3^2 \times 5^2$, \wp_C -code = 5^4 , \wp_D -code = $3^2 \times 5^2$.

Remark 3.1. We have codified the *CP* with a prime number, because this defines an univocal *CPS-code* for a generic network node.

Definition 3.1. We will define a *node cluster* ζ_A the nodes set of a network, with same \wp_A -code, so $\zeta_A = \{A, C\}$.

Remark 3.2. We observe that, if two nodes belong to the same complex network (Figure 1), it doesn't implicate that they belong to the same class, because each node class is defined by the *CPS-code*. Particularly two similar *CPS* define two different clusters in a unidirectional networks and same clusters in a bidirectional network.

If we consider two *CPS*, \wp_H and \wp_K with $\wp_K \subset \wp_H$ i.e. if \wp_K -code is a integer divisor of \wp_H -code, then ζ_K is a subcluster of ζ_H , therefore it is possible to individualize all *subclusters* of the node cluster *H*, considering all the nodes with *CPS-code* integer divisor of ζ_H -code. We observe that, if the Greatest Common Divisor (*GCD*) between two *CPS-codes* associated to ζ_H and ζ_K , is equal to *CPS-code* of ζ_K , then ζ_K is a *subcluster* of ζ_H .

Remark 3.3. If we consider *GCD* between \wp_A -code and \wp_B -code (Figure 1): $\text{GCD}(5^4, 3^2 \times 5^2) = 5^2$, then the *GCD factors*, define the common number of *complete paths* between \wp_A and \wp_B with *same length*. Now, if we divide \wp_A -code with $\text{GCD}(\wp_A, \wp_B)$, we have a integer rest. If we factorize this rest, the primes define the length of the *complete paths* of \wp_A not belonging to \wp_B .

We can verified that the nodes network *A* and *B* are *almost-similar* (Figure 1, *Example 2.4*), using *CPS-code*. In fact \wp_A -code = 5^4 , \wp_B -code = $3^2 \times 5^2$ and the $\text{GCD}(\wp_A\text{-code}, \wp_B\text{-code}) = 5^2$, i.e. \wp_A and \wp_B have two common *CP* of length 5, therefore \wp_A and \wp_B are *almost-similar* and the nodes *A* and *B* define two different node clusters, ζ_A and ζ_B for the bidirectional network in Figure 1.

Finally in Table 2, we have considered 10 bidirectional networks with different number of nodes and of connections and evaluated the networks analysis times, with our clustering algorithm, using a Pentium 4 with CPU 2Ghz and 1Gb ram.

Table 2. *Algorithm performance on different bidirectional networks.*

| Nr. Nodes | Nr. Connections | Time (sec) |
|-----------|-----------------|------------|
| 10 | 45 | ≈ 0.05 |
| 20 | 190 | ≈ 0.05 |
| 40 | 780 | ≈ 0.05 |
| 80 | 3160 | ≈ 0.22 |
| 100 | 4950 | ≈ 0.35 |
| 150 | 11175 | ≈ 0.95 |
| 200 | 19900 | ≈ 3.00 |
| 300 | 44850 | ≈ 13.00 |
| 400 | 79800 | ≈ 36.00 |
| 500 | 124750 | ≈ 80.00 |

5. Conclusion

The clustering method proposed for the analysis of the bidirectional complex networks, is a fast method that could be applied in different scientific sectors. The analysis is founded on idea to gather in classes the network nodes or more networks with same number of connections and paths with equal length. Each class of nodes is characterized by a code obtained codifying each complete path (CP) with a prime number. The simplicity of the clustering algorithm introduced, gives a strong computational implementation and therefore of great applicability to every network type. Finally, we want underline that by generic *CPS-code*, it is possible to individualize the nodes with more connections, i.e. the nodes with more input/output and the root nodes with paths more long. To start these information, this clustering method could be used as routing algorithm in the telecommunication networks too. Finally the object of future paper will be develop the present research to describe the clustering method based on primes for unidirectional complex networks.

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