Structural Analysis of a Core Periphery Network: Co-Authorship on Dengue

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Abstract. We find in the literature empirical results and models to explain the growth of Small World and Scale Free networks but very little on Core Periphery networks. The present work focuses on a co-authorship network based on scientific publications for Dengue Fever, a so-called "Neglected Disease". We found that for this network the Core Periphery structure presented resembles a crossover between Small World and Scale Free networks, with a heterogeneous behavior of its authors. As a further result of this research we demonstrate, based on the network cores, how the authors divide themselves into groups and their influence over the network.

Resumo. A literatura apresenta uma série de pesquisas e modelos que explicam o crescimento de redes "Small World" e "Scale Free". No entanto, pouco é discutido sobre redes "Core Periphery". O presente trabalho utiliza uma rede de coautoria baseada em publicações científicas da Dengue, conhecida como uma das doenças "Negligenciadas". Para esta rede foi analisado o quanto que a estrutura "Core Periphery" se assemelha a estrutura de redes "Small World" e "Scale Free", demonstrando o comportamento heterogêneo dos seus autores. Como resultado complementar da pesquisa é apresentada, com base nas comunidades centrais, a forma como os autores se dividem em grupos e a sua influencia na rede.

1. Introduction

The use of articles and research from journals and scientific databases has, for a long time, been a good source of information for analysis on complex networks, providing an overview on the patterns of collaboration within the academic community and its authors. "Co-authorship of a paper can be seen as a documented form of collaboration between two or more authors. These collaborations form a co-authorship network, in which the network nodes represent authors, and the lines between the nodes the collaboration of authors that have coauthored one or more papers together. The structure

of such networks turns out to reveal many interesting features of academic communities" (M. E. J. Newman 2004).

In the past few years, co-authorship analysis has been applied in scientific research and publications quite often and in many different fields. Some of the first researchers on the area of complex network, such as Albert Barabási and Mark J. Newman have used co-authorship networks on their studies (Barabasi et al. 2002; M. E. J. Newman 2004; M. E. J. Newman 2001), which provided a good basis for the work been pursued today. In regards to the medical fields and specially on specific diseases we have also encountered research applied with this type methodology and corpus (Morel et al. 2009; Ramos, Gonzalez-Alcaide, e Bolanos-Pizarro 2013).

The use of complex network analysis methodologies have been often used to find communities within research fields. The use of dynamic or temporal analysis is also another area within co-authorship research where the analysis focuses on the network structural change. In our study we have been looking on formation and maintenance of relations within and between communities, taken into consideration the dynamics of the network, in order to understand how positional control might come into place and why.

For our research the data defined for the analysis is based on research publications for Dengue Fever (also know as Dengue), a disease contained in the so-called neglected tropical diseases (NTD). The term NTD, which has been used since the mid 1990s, has become a "brand-name" referring to a group of diseases that are especially endemic in low-income populations living in tropical and subtropical countries (Ramos, Gonzalez-Alcaide, e Bolanos-Pizarro 2013). However, up to this date, there are no clear or agreed definitions for what constitute a Neglected Disease.

Our intentions at this point with this research is to understand the structure of such networks and how this structure and the influence of key actors might affect the understanding of the network, with the possibility to aid or provide valuable information for government agencies or public policies makers on regards to NTD research.

1.1. Network Structures

One of the issues been faced in statistical physics was to define new concepts and measures to try to infer the structural properties of large empirical networks. "The main outcome has been the identification of a series of unifying principles and statistical properties shared by most of the real-world networks examined" (Gay 2012).

Real-world networks have been found to differ from the classical random graph theory of Erdös and Rényi (1959), which carried statistical regularities not been anticipated. For those networks, the small-world property (relatively short paths between any two nodes and a large clustering coefficient) and scale-free degree profile (power-law scaling for the probability distribution of the number of links at a node) were a common structure. New models were developed to reproduce those structural properties observed in real topologies. Networks with high clustering coefficient and small average path length can be generated with an evolution by the small-world model of Watts and Strogatz (1998), while networks with power-law degree distribution can be generated with an evolution by the scale-free model of Barabási and Albert (1999).

Another common but informal notion in social network analysis and other fields is the concept of a core/periphery structure. The intuitive conception entails a dense, cohesive core and a sparse, unconnected periphery. According to Borgatti (2000) "the notion of a Core Periphery network can be seen as a center with many cores and a peripheral community that are connected to those different cores."

1.2. Network Concepts

"A network can be seen in its simplest form as a collection of points, called nodes or vertices, joined together in pairs by lines, called edges" (M. Newman 2010). Networks can be used to model systems where entities could be seen as the nodes and their relations the edges on the network. In order to understand and analyze the network a series of measures have been developed and some of the most common and widely used is regarded as node centrality. Some of those measures are degree, betweenness and eigenvector, which have been used on our study. The degree of a node is the number of lines that are incident with it; betweenness is a measure of how nodes might influence the interaction of other nodes on the network; "eigenvector centrality gives each vertex a score proportional to the sum of the scores of its neighbors, in a sense that as more influential nodes a node has as its neighbors higher its own score" (Wasserman e Faust 1994; M. Newman 2010).

In regards to structure the concepts of cluster coefficient also know as transitivity and average path length or geodesic distance play an important roll. There are many types of relations on a network where the simplest is "connected by an edge." If the "connected by an edge" relation were transitive it would mean that if vertex u is connected to vertex v, and v is connected to w, then u is also connected to w. As for geodesics it's definition implies the shortest path between two nodes and the average of all the paths on a given network (Wasserman e Faust 1994; M. Newman 2010).

For last this study have made use of the concepts of k-core and community detection as a form to understand the structure and its subdivisions. A k-core is a subgraph in which each node is adjacent to at least a minimum number, k, of the other nodes in the subgraph (M. Newman 2010). For community detection there is a concept called modularity (M. Newman 2006) that measures how well a network is subdivided.

1.3. Methodology

The advance of technology has permitted a wider analysis of some networks especially on large data sets. Analysis of data sets with tens or hundreds of thousands of vertices and edges were not common and restricted to small groups of researchers. For the data treatment we used the software Tetralogie and R programing language and for network manipulation and visualization we used the package Igraph on R and Gephi.

Tetralogie allows users to conduct strategic analysis from heterogeneous textual data through the use of conventional and innovative methods. Another feature of the software is that it allows graphical visualization for understanding human activities and their interactions as well as their evolution in a decision-making perspective (Gay e Dousset 2006).

The R software is a free descriptive programming language and environment for statistical computing and graphics. It allows the treatment and analysis of large data sets including network analysis by the use of packages such as igraph and sna. Gephi is an interactive visualization and exploration platform for all kinds of networks and complex systems, dynamic and hierarchical graphs.

For the data collection we have used PubMed¹ database, with a search term/descriptor "dengue" for papers from 1981 to 2012 included. PubMed is a free database accessing references and abstracts on life sciences and biomedical topics.

2. Data on Scientific publications

The research has been made taking into consideration authors of articles, countries where those authors were from and journals used for the scientific publications. Data files were retrieved from PubMed with the term/descriptor "dengue" for title and abstract. The database was than converted into a co-authorship network taken into consideration the authors and their co-appearances on the respective articles.

2.1. Statistical Data

The numbers concerning the research publications on Dengue has grown with the past years as concerns with the NTDs have become more common. Countries that had many cases of the diseases became key research players on this growth. Over the 30 years span for this research the number of publications have jumped from 67 on 1981 to 1,053 publications on 2012. For the total number of publications we have retrieved 9,496 articles containing a total of 21,083 authors from 102 different countries. The number of journals added up to 1,257, however only 309 of those had 5 or more articles in the area and 175 journals had 10 or more publications. The most common journals retrieved from the database and their respective number of publications on Dengue for the period were: American Journal of Tropical Medicine and Hygiene with 506 publications; Southeast Asian Journal of Tropical Medicine and Public Health (304); Journal of Virology (282); PLOS Neglected Tropical Disease (192); Virology (180).



FIGURE 1 - WORLD MAP FOR PUBLICATIONS ON DENGUE(1981-2013)

As for the countries of authors on the research we have noticed a great change on participation over the years. Some countries have been responsible for the research for much of the period studied but others became key players as the time progressed. Countries such as USA (3,386) and France (918) falls on the first category and

¹ <u>http://www.ncbi.nlm.nih.gov/pubmed</u> accessed on october 2012

emerging countries happen to be Brazil (1,911), India (889) and Thailand(836). Those were the top 5 countries in terms of number of publications and the respective numbers in parenthesis. Figure 1 shows a world map for publications. The liter the color for the country the greater the number of publications assigned to that country.

2.2. Network Data

Network information is important in the sense that it gives insights and helps clarify the structure of the network, the main goal of our research. Because of the size of the network, visualization at this point is meaningless once there is no clear understanding of its positioning. One point we can make at this stage is that, using a centrality-based distribution such as Frunchterman Reingold, there is a very dense center on the network and a periphery of less connected vertices. Table 1 gives some of the information regarding the network with a comparison over a random network based on Erdös and Rény theory of random graphs.

TABLE 1. COMMUNITIES AND NODE DISTRIBUTION

	Nodes on Main	Edges on Main	Mean Path	Mean Path Length	Cluster	Cluster Coefficient
	Component	Component	Length	Random Network	Coefficient	Random Network
Values	18,023	119,148	4.85	4.06	0.38	0.00068

The first column on Table 1 shows the number of nodes on the main component (biggest connected component) as 18,023 and the number of edges as 119,148. Those numbers have been used to create a random network. The mean path length of the network is similar to the random network denoting a possibility for small-world structure. As for the Cluster Coefficient the network studied shows a much greater score. This denotes how well connected a scientific collaboration network might be in comparison to a random network.

The number of edges and its cluster coefficient implies a very dense network with large number of connections amongst the different authors. The average degree found is 13, which means that on average each author collaborates with 13 other authors. The maximum degree for the network was 386.

3. Network Structure

The network structure defines how the interactions amongst the different actors might occur and several conclusions can be drawn for the understanding of this structure in regards to scientific research collaboration. In a heterogeneous behavior, some vertices on the network will act in a very different way from the rest of the network, not in a sense of an outlier but in log scale distribution. Networks on this form are called scalefree once this is an attribute of a true log log distribution curve. Few vertices have a very large number of connections and the majority of vertices have a very small number of connections.

Many networks studied over the past years have been called scale-free but some of those have not presented true evidences of its claims. According to a study on powerlaw distribution (Clauset, Shalizi, e Newman 2009) half of the networks analyzed by them failed to behave as such. On our research we have applied the algorithm of Clauset (2009) in order to validate our assumptions. Both Publications and Degree distribution had a true log log behavior as stated by the algorithm with a high p-value, although the tail cut-off or x-min for the distributions happened to be quite high.

Because of the somehow similar behavior of both curves we can assume that the number of publications is a direct influence of the degree of its authors, although there are some cases that it does not hold true. In one particular case a single article had 94 authors, which made the degree of the authors on that article very high in comparison to other authors with similar number of publications. The average number of authors per paper has increased over the years from 2.5 authors per paper in 1981 to 4.3 in 2012.

Figure 2 shows the two distributions in terms of number of publications and number of degree on x axis and the number of authors on a cumulative distribution on y axis. As we can see, there are few authors with very high number of publications (above 100) and degree (above 200) and more than 50% of authors with less than 5 publications or 10 degrees.



FIGURE 2 - AUTHORS PUBLICATIONS AND DEGREE DISTRIBUTION (1981-2013)

At this point we can assume that the network follow a power-law distribution with a heterogeneous behavior of its authors. Some of the researchers on the filed have a very high concentration and influence over the network based on the number of publications they had up to this date and the number of relations as a consequence.

As for the Small World characteristics we have seen a low average path length and a high cluster coefficient. If we use the betweenness centrality to analyze some of the authors we can understand how those authors might connect the different pockets or groups.

The question now arises as how this possible Core Periphery structure interacts with the Scale Free and Small World structures and characteristics. Can we say that the Core Periphery is a mix of the two others? The idea with a Core-periphery structure is that there are a core structure with highly cohesive nodes and a periphery structure with low cohesion among the nodes but with a considerable number of connections to the core of the network. Some nodes on the core of the network connect themselves to other core networks but do also connect themselves to the peripheral nodes, which makes the structure as it is.

The first step we did to define the core of the network was to use a k-core algorithm. K-core helps identify people that have a high number of connections within a group that also have a high number of connections among themselves. This definition could be used to understand part of our core structure in the network.

4. Network Core and Communities

One important concept over the network analysis is the idea of a cohesive or core group. An easy form to find this group is to make use of a k-core algorithm. In this way one can understand how groups of people connect themselves with others that have similar degrees. This definition is very adequate to the understanding of a core periphery network, although it is not enough by itself.

For this research we have applied a filtering to find the k-core nodes that would help explain a core periphery structure and control. At first we filtered all the nodes that have only one publication to avoid the problem of getting authors that only published once but with many co-authors in the same article. Second we used the software to find the k-core levels close to 10% and 1% of the nodes on the network but in the last case with the concern that it would not disrupt or disconnect the clusters entirely. The k-cores and number of nodes were k-core 5 with 1,820 authors, approximately 10% of the network, and k-core 20 with 279 authors representing 1.55% of the network.

Another important concept for our analysis is the idea of network clusterization or modularization. How should we separate groups within a network based on their connections with other nodes? Many studies have been made in regards to the division of networks by algorithms and we have decided to use the latest Newman's leading eigenvector algorithm (M. E. J. Newman 2006) implemented on R by Igraph package. The decision was made based on the possibility of application on large size networks, not possible for other algorithms with the same level of modularity. The idea of the algorithm is to break the network into groups based on their eigenvector values where a node would become the leading eigenvector for a particular group and other nodes on the network with similar eigenvector values would fall on the same group.

After applying the algorithm to the network we were able to get 22 communities with its sizes shown on Table 2. Within each community we analyzed how the nodes on k-core 5 and nodes on k-core 20 were distributed within the communities.

Node Distribution within Communities												
Community #	Nodes	K-core 5	K-core 20	Community #	Nodes	K-core 5	K-core 20					
1	2488	371	95	12	381	40	8					
2	348	45	19	13	59	0	0					
3	3736	431	62	14	401	42	1					
4	350	53	7	15	727	55	21					
5	6127	355	0	16	132	11	5					
6	1207	188	47	17	83	12	0					
7	161	26	0	18	68	9	0					
8	278	34	1	19	13	1	0					
9	251	22	1	20	9	0	0					
10	482	37	3	21	85	19	0					
11	618	68	9	22	19	1	0					

TABLE 2. COMMUNITIES AND NODE DISTRIBUTION

As the first analysis of the communities we can see that the division is not regular in terms of size, with community 5 with 6,127 nodes and community 20 with 9 nodes for the two high and low extremes. It is important at this point to understand how the algorithm behaves in terms of eigenvector centrality. Because the highest

eigenvector values found are appointed to the primary groups, some authors on communities 1 through 4 might have a higher importance than those authors on community 5 as we will see later on. If we follow this concept one can see that on k-core 20 there are no authors on community 5, which implies that these community, although has a great number of nodes, those nodes do not carry a great significance in terms of degree.

The second step on the analysis is the distribution of nodes within the communities at the k-core level. What we aim to see is how the k-core behaves in terms of distribution within each community once there is a clear differentiation of sizes. Do some communities carry a greater number of important, or higher k-core, authors or not. We found that the distribution of k-core nodes in the communities follows the same percentage as the entire network, close to 10% for k-core 5 and close to 1.5% for k-core 20. This means that for each community, independent of its size, the number of authors belonging to k-core 5 is approximately 10% varying between 8% and 12% on average with higher percentages for communities 1, 4, 6 and 7 but not more than 16%. This implies that the algorithm, based on eigenvector centrality, do not discriminate the communities in terms of degrees. If we check the distribution with k-core 20 that has only 279 authors, the average of 1,55 % remains very close with communities 1, 2 and 6 with higher percentages and some communities without any author at this level, such as community 5, 7, 13 and 17 to 20.

The distribution of the k-core nodes happened to be a very interesting one. If we think in terms of research groups we could understand how they come to be formed. Some researchers, as head of research groups, hold a greater number of publications and degrees, but their group of people, divided by the algorithm, does not put them together with other authors just based on the degree.

5. Relations and Control

The last attempt of our study was to understand how those core groups, defined by the k-core algorithm, relate themselves to the rest of the network. Going back to the definition of a core periphery network, a node to be considered core must have a high number of connections with nodes at the same level, found by the k-core, and connections with the rest of the network as seen by the division of the groups, or the periphery on other groups.

Applying a one level node neighborhood we could find how many connections or degree each author has. We used the same concept for all the nodes in each k-core and subtracted the repeated node' neighbors to find the level of influence for each of the k-cores. For the k-core 5 we found that with only 10% of the nodes (1820) that level could influence or connect themselves to more than 82% of the network. For the k-core 20 the 279 authors, or 1.55% of the network, had a direct influence over almost 37% of the network. If we take into consideration that close to 50% of the network has fewer than 5 publications we could assume that a very few group of authors might have influenced the entire field over 3 decades.

What we have found up to now is that the network divides itself into groups that are not related to degree centrality but to eigenvector centrality. We also have found that a very few number of authors have influence over a great portion of the network. But how are those nodes, divided into groups, related to themselves? In order to answer these question we have draw a network of communities and their relations. The idea was to understand the influence of the communities, their exchange or relations and the structures in regards to the core periphery characteristics.

Figure 3 shows the network of community relations. The size of each node is a proportional relation to the number of nodes on each community, normalized in order to be able to see all the communities. The size of the links is a direct proportion of the number of relations between each community. The number on each node is the respective community number that could be compared to Table 2. For the distribution it was applied the Fruchterman Reingold algorithm. The communities with less importance or size were mostly drawn on its periphery while the most important communities stayed at the center of the graph.

An important analysis of Figure 3 is the number of connections coming from community 1 and to which communities it most connects itself. If we relate this information with the number of k-cores 5 and 20 nodes we could understand the reason for this great exchange.



FIGURE 3 - NETWORK COMMUNITY RELATIONS

With this analysis we could conclude our assumption on a core periphery network. Within each community, there is a regular distribution of high degree nodes that holds the community together. Those high degree nodes connect themselves to lower degree nodes within each community. Between the communities there are a high number of relations, as seen on Figure 3, which enforces the idea of a small world attribute with few high betweenness score authors. There are also smaller communities that do not carry many high degree authors. Nevertheless, those smaller communities still well connected over the network defining its periphery.

6. Conclusions

For this research we have tried to understand the structural behavior of a scientific collaboration network based on a specific field, Dengue. We have found a clear core periphery structure with many central communities on the network that relate themselves to the periphery but also to other central communities and authors. Those communities were well distributed and the authors within those communities played an important roll in maintaining the network together with its distributed relations.

We have also found that this network carry characteristics of Scale Free networks once we have applied techniques to confirm its distribution and behavior. Over the scientific field there are few very well connected researchers that somehow controls the relations with the rest of the network. With the assumption of Small World attributes such as small short path lengths, high clustering coefficients and communities connecting themselves with important nodes relations we confirm all three network structures implying a possibility of interlayer characteristics among those structures.

As for the field we could conclude that it is very cohesive, as a scientific field should be, but is also well controlled by few authors on the network. Further studies on the structure could help understand how knowledge is distributed or dispersed in the network and how innovation, if any, happens on the network and where, on the core or on the periphery.

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