1 Community Structure

Many real networks have a natural community structure, where disjoint subgroups of nodes exchange more connections within their subgroup than between subgroups. Formally, we want to compute the optimal division of the network that minimizes the number of links between subgroups (also called communities). The raw number of links across boundaries of communities does not give a good partition of the network. For example, the community structure can be a consequence of random variations in the density of links. A more reliable approach uses the configuration model [5] as a null model to assess the quality of a given network partition. Newman and Girvan [4] define the modularity as follows:

$$Q = \frac{1}{2m} \sum_{ij} \left(A_{ij} - \frac{k_i k_j}{2m} \right) \delta(g_i, g_j) \tag{1}$$

where $m = \frac{1}{2} \sum_{ij} A_{ij}$ is the number of links and g_i indicates the label of the community the node i belongs to. Notice that maximizing the above function yields a partition that minimizes the expected number of links falling between different communities, i.e., when $\delta(g_i, g_j) = 0$. Modularity Q takes values between -1 and 1: low modularity indicates the number of links between distinct communities is not significantly different from the random distribution and high modularity indicates there is a strong community structure.

2 Community detection in a single layer

We analyzed the real Twitter data using two the binary and the weighted version of the adjacency matrix using a Matlab version of the Louvain algorithm [1]. The results are (confirming Elisa's values):

	Binary	Weighted
c	27	37
Q	0.2610	0.4082

As we can see, the modularity value increase considerably from binary to weighted representation. However, because the representation of the network is different (binary vs weighted), the modularity value is not enough to say that the community structure is different. The reason of the difference in values may be only because we chose different representations. However, we will show in the following subsection that the community partition is also different.

2.1 Distinguishing binary vs weighted community partition

By just looking at the number of modules in each representation, we can be tempted to say that the community structure is different. However, as the next table shows, the percentage of total nodes inside the biggest five nodes is very high. In other words, the first five biggest modules dominates the community structure.

	1	2	3	4	5	Total %
Binary	1092	941	410	292	12	97.76
Weighted	1098	938	220	293	140	95.69
Shared	876	777	66	244	0	69.86

First, notice that the total size (last column) of the five biggest modules very high (> 95 %) as already described. Second, notice that for the binary row we sorted the modules in decreasing order. However, for the weighted row this is not the case (see module 3 and 4). We relabel the weighted partition in order to maximize similarity structure with the binary partition.

2.1.1 Jacquard's index

The reader may already infer that the value at the right-bottom is an estimation (based in the fact that > 95% of nodes are inside the five biggest modules) of the Jacquard index, which in this case can be translated as:

$$J = \frac{\text{Number of nodes inside the same modules}}{\text{Total number of nodes}}$$
 (2)

3 Community detection across time snapshots

In addition to study the community structure a static single network, we studied how the community structure changes across time snapshots. The main reason of this analysis is that, in real Twitter data, the mutual information between users may vary across time, and therefore the community structure of the mutual information network may change across time. As a specific example, this type of analysis could potentially be a method to detect the birth of trending topics in the Twitter social network.

3.1 Algorithms

In order to perform the analysis we used two modularity algorithms specifically designed either for time snapshots or multiplex networks (a time varying network can be described as a multiplex network in which each time snapshot is a multiplex layer). These two algorithms are described on the articles published by Kawadia and Sreenivasan [2], and Mucha et Al [3]. Both algorithms are available online as open source in Python and Matlab, respectively.

3.1.1 Kawadia and Screenivasan algorithm

The idea of this algorithm is to find a new community partition P_t for a graph snapshot G_t , such that the estrangement (a distance metric that the authors defined) between the new partition and the previous one is smaller than δ . In other words, the algorithm solves the following constrained optimization problem:

$$\begin{array}{ll}
\text{maximize}_P & Q(P) \\
\text{subject to } E(P) \le \delta,
\end{array} \tag{3}$$

where E(P) represents the estrangement between the current and the last community partition. The authors make use a Lagrange multiplier to solve this problem (for more details see [2]). Unfortunately, even that the algorithm worked for the synthetic data, it did not for the real Twitter data. The reason is that the Python library that they use to solve the constraint satisfaction problem becomes really slow for networks of big size as is the case of the real Twitter data that we present here.

3.1.2 Mucha et al. algorithm

This algorithm is based in optimizing a modularity definition that includes extra terms for the existence of different network layers, which in this case consist of different time snapshots. The new modularity definition that the authors propose to optimize in order to find the change of modularity across layers is:

$$Q_{multilayer} = \frac{1}{2\mu} \sum_{ijsr} \left\{ \underbrace{\left(A_{ijs} - \gamma_s \frac{k_{is} k_{js}}{2m_s}\right) \delta_{sr}}_{\text{within layer contribution}} + \underbrace{\delta_{ij} C_{jsr}}_{\text{between layer contr.}} \right\} \delta(g_{is}, g_{jr})$$
(4)

where A_{ijs} is the element i, j of the adjacency matrix of layer s, k_{is} the degree of node i inside layer s, m_s the total weight of layer s, g_{is} the module index of node i inside layer

s. Finally, the user parameter γ_s balances the within layer contribution and C_{jsr} balances the between layer contribution. C_{jsr} can be seen as a virtual link between node j's of two different layers s and r.

In order to optimize the modularity equation, the authors use a generalization of the Louvain algorithm [1], which is one of the most used when dealing with large scale networks. The released Matlab library of this algorithm outperformed the previous one in terms of performance and speed. Therefore, this algorithm is the one that we used to find how modularity changes across time.

3.2 Data

We create seven different time snapshots of the Twitter data, each corresponding to a different week.

3.3 Results

Figure 1 represent the modularity across the seven different weeks for a single combination of parameters. This Figure shows, that in essence, the structure does not vary across time. This fact tell us that for this data it may be better to just evaluate the modularity of a single snapshot across the entire seven weeks.

References

- [1] Vincent D Blondel, Jean-Loup Guillaume, Renaud Lambiotte, and Etienne Lefebvre. Fast unfolding of communities in large networks. *Journal of Statistical Mechanics: Theory and Experiment*, 2008(10):P10008, 2008.
- [2] Vikas Kawadia and Sameet Sreenivasan. Sequential detection of temporal communities by estrangement confinement. *Scientific reports*, 2, 2012.
- [3] Peter J. Mucha, Thomas Richardson, Kevin Macon, Mason A. Porter, and Jukka-Pekka Onnela. Community structure in time-dependent, multiscale, and multiplex networks. *Science*, 328(5980):876–878, 2010.
- [4] Mark EJ Newman and Michelle Girvan. Finding and evaluating community structure in networks. *Physical review E*, 69(2):026113, 2004.
- [5] Mark EJ Newman, Steven H Strogatz, and Duncan J Watts. Random graphs with arbitrary degree distributions and their applications. *Physical Review E*, 64(2):026118, 2001.

Community Structure across time

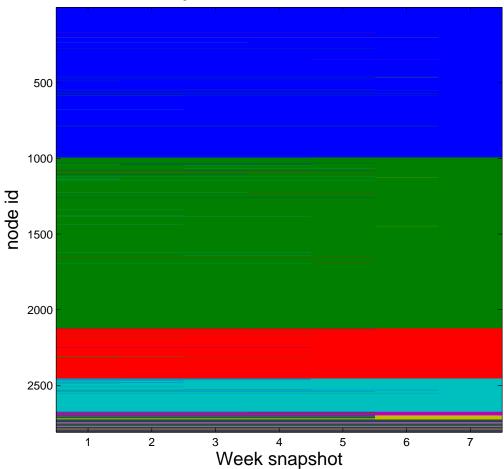


Figure 1: Modularity across time with parameters $\gamma_s = 1$ for all s and $C_{jrs} = \omega = 0.25$ for all j, s, r. Different colors represent the module id to which each node (rows in this figure) belong to. We can observe, that the structure is dominated by the existence of four big modules plus a lot of small size modules (bottom of the Figure). Further, the modularity structure remains almost constant across time in the sense that only a very small number of nodes (represented as colored lines inside the big modules) switch to other modules as time develops.