University of Szeged

Embedded network structures of transaction graphs

Master Thesis

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# Abstract:

A prototype console application has been implemented to detect social networks and clustering transactional type of the graphs. Several algorithms have been implemented to run against the given databases, such as Girvan-Newman, Markov chain, etc. The application is able to map a graph (also known as network) with any given delimiter, in any text format furthermore there is an option to connect to database if necessary. The results of the algorithms are plotted and displayed after the run. The coloring logic has been implemented in a fairly naive and greedy colormap. The application is scalable and modularized.

# Introduction

World Wide Web, blogging platforms, instant messaging and Facebook can be characterized by the interplay between rich information content, the millions of individuals and organizations who create and use it, and the technology that supports it. This thesis will cover the processing of recent research on the structure and analysis of large social and transaction networks and on models and algorithms that abstract their basic properties. Unusual ways have been explored how to practically analyze large scale network data and how to reason about it through models for network structure. Topics include methods for network community detection and their connection with transactional graphs. [1]

Community detection and analysis is an important methodology for understanding the organization of various real-world networks and has applications in problems as diverse as consensus formation in social communities. Currently used algorithms that identify the community structures in large-scale real-world networks require a priori information such as the number and sizes of communities or are computationally expensive. I intend to rely more on algorithms, which use the network structure as their guide instead of this priori information. Finding community structures in networks is another step towards understanding the complex systems they represent [2]. Social networks are represented by people as nodes and their relationships by edges therefore they contain more triangles than a random graph with similar edge density or degree properties. In contrast technological or transaction graphs contain fewer triangles and often display tree-like structures [15].

I gained knowledge about some of the prerequisites of social graph studies, for example algorithms of community detection have been used, along with the extracted data from an earlier implemented software, the Sixtep program, as well.

Based on the research written in article [15] and some earlier studies I intend to give an explanation about the implementation of algorithms for generalized coloring of transactional graphs. As the study goes, I follow the heuristics and possible solutions for the clustering problem.

*There was an attempt to find the embedded pattern in the provided transaction-like graphs.*

# Used Technologies:

* Python, PyEnv, PyCharm
* NetworkX
* Sixtep software
* Git
* Graph Databases

## Python, PyEnv, PyCharm

Python is a programming language that makes possible to work more quickly and integrate the designed system more effectively. Perfect choice to develop proof of concepts, make prototypes. It can be easy to pick up whether the developer makes a first time project or has experience with other languages. The well documented packages help the programmer along the learning way and the gains in productivity can bee seen almost immediately. It also reduces maintenance costs of the targeted application. The packages used in this thesis are developed under open source license, making it freely usable and distributable not only educational purposes but even for commercial use. [16]

PyEnv makes it possible to switch between multiple versions of Python. It is unobstructive and follows the UNIX tradition of single-purpose tools. PyEnv-virtualenv is a PyEnv plugin that provides features to manage virtualenvs for Python on UNIX-like systems.

PyCharm is a Python IDE with intelligent code completion, on-the-fly error checking, quick fixes, built in support for Numpy, Matplotlib and other scientific libraries while offering graphs, array viewers and much more.

These tools have been chosen for thesis purposes not only because they are open source, but the documentations and tutorials are in good quality. The tools have wide range of support layer and make the development ready to start. Perfect choice when the scope is widely changes during the development and each features have to be prototyped at first.

## Git

Git is a free and open source distributed version control system designed to handle everything from small to very large project with speed and efficiency. It has wide range of supported features, like branching and merging strategy, distribution, data assurance, staging area, however during the app development for the thesis, it has only been used for version control.

## NetworkX

NetworkX is a Python package for the creation, manipulation, and study of the structure, dynamics, and functions of complex networks. It supports a variety of features, provides classes for graph objects, generators to create standard graphs, IO routines for reading in existing datasets, algorithms to analyze the resulting networks and some basic drawing tools.

Most of the NetworkX API is provided by functions which take a graph object as an argument. Methods of the graph object are limited to basic manipulation and reporting. This provides modularity of code and documentation. It also makes it easier to learn about the package in stages.

In NetworkX, a graph is represented by a collection of edges that are pairs of nodes. Attributes are often associated with nodes and/or edges. NetworkX graph objects come in different flavors depending on the properties of the network, like is the graph directed, or are multiple edges allowed. Based on that, the basic graph classes are Graph, Directed Graph, Multi Graph, Multi-Directed Graph. Graph objects can be created by importing data from pre-existing (usually file) sources. In this thesis, graph algorithms provided by NetworkX include shortest path and breadth first search and clustering algorithm. While NetworkX in not designed as a network drawing tool, it provides a simple interface to drawing packages and some simple layout algorithms. In the application drawing has been done using this package and the Matplotlib Python package. The basic drawing functions essentially place the nodes on a scatterplot using the positions which have been provided via a dictionary or the positions are computed with a layout function. The edges are represented as lines between the dots. NetworkX uses dictionaries as the basic network data structure. This allows fast lookup with reasonable storage for large sparse networks. [4]

* Data structures for graphs, digraphs, and multigraphs
* Many standard graph algorithms
* Network structure and analysis measures
* Generators for classic graphs, random graphs, and synthetic networks
* Nodes can be "anything" (e.g., text, images, XML records)
* Edges can hold arbitrary data (e.g., weights, time-series)
* Open source 3-clause BSD license
* Well tested with over 90% code coverage
* Additional benefits from Python include fast prototyping, easy to teach, and multi-platform

Algorithms:

Bipartite module provides functions and operations for Graph and DiGraph classes. The algorithms are not imported into the NetworkX namespace at the top level, so it has to be added manually. The convention is to use a node attribute with value 0 or 1 to identify the sets each node belongs to. The functions in this package do not check that the node set is actually correct nor that the input graph is actually bipartite.

*For bipartition computing, the Kernighan-Lin bipartition algorithm is a good choice. This algorithm partitions a network into two sets by iteratively swapping pairs of nodes to reduce the edge cut between the two sets.*

Finding the largest clique in a graph is NP- complete problem, so the find\_cliques() function probably has an exponential running time, however it returns all maximal cliques in an undirected graph.

*K-Clique. A k-clique community is the union of all cliques of size k that can be reached through adjacent k-cliques.*

Modularity-based communities: greed\_modularity\_communities()

Greedy modularity maximization begins with each node in its own community and repeatedly joins the pair of communities that lead to the largest modularity until no further increase in modularity is possible.

*Helper Functions.*

*Edges\_equal()*

*Write to file*

## Sixtep software:

The software has been released in 2007 by network theory researchers and CRM advisors. It is able to load a graph and visualize it. After adding a graph, an attribute file can be attached to it, provide information about the nodes with it. For example, if the nodes are representing people, such information like name, age, city can be added.

Several algorithms are implemented to clusterize or detect communities such as the Newman-Girvan algorithm or the Markov chain model. The UI representation of the graph is user friendly; the location of the nodes can be easily modified by clicking one by one or select a targeted area. Several built-in functions help to make the graph more interpretable. The user can select unique modules like clusters or communities and display only the selected ones.

3 algorithms have been used in this thesis to classify the nodes:

* Markov chain clustering
* Maximized modularity
* Community detection

The source code of the software cannot be accessed; therefore, the software is not expandable, cannot be further developed. However, it is possible to get the calculated cluster and community data with the export function. The exported data needed further processing, since the data structure was not in the right format to manufacture the NetworkX Graph object. An import function processes the data from the export then a factory function provides the Graph object that is needed. After that, a colormap is assigned to the nodes by the exported labels. The result can be compared with the built in NetworkX functions via modularity and/or the number of the used colors.

Sadly, there is no information provided in the software about the edges between the clusters, however valuable data can be found while exploring that area. See more at graph condensation.



## Graph Databases

A graph database is defined as a specialized, single-purpose platform for creating and manipulating graphs. Graphs contain nodes, edges and properties, all of which are used to represent and store data in a way that relational databases are not equipped to do. Graph analytics is another commonly used term and it refers specially to the process of analyzing data in a graph format using data points as nodes and relationships as edges. Graph analytics requires a database that can support graph formats. This could be a dedicated database, or a converged database that supports multiple data models, including graphs.

There are two popular models of graph databases: property graphs and RDF graphs. The property graph focuses on analytics and querying, while the RDF graph emphasizes data integration. Both types of graphs consist of a collection of points (nodes) and the connection between those points (edges). But there are differences as well.

Property Graphs.

These types of graphs are used to model relationships among data and they enable query and data analytics based on these relationships. A property graph has nodes that can contain detailed information about a subject and edges that denote the relationship between nodes. The nodes and edges can have attributes, called properties, with which they are associated. Because they are so versatile, property graphs are used in a broad range of industries and sectors, such as finance, manufacturing, public safety, retail and many others.

RDF graphs.

RDF stand for Resource Description Framework. These graphs conform to a set of W3C standard designed to represent statements and are best for representing complex metadata and master data. They are often used for linked data, data integration and knowledge graphs. They can represent complex concepts in a domain, or provide rich semantics and inferencing on data. In the RDF model a statement is represented by three elements: two nodes connected by an edge reflecting the subject, predicate and object of a sentence – this is known as an RDF triple. Every node and edge is identified by a unique URI, or Unique resource Identifier. The RDF model provides a way to publish data in a standard format with well-defined semantics, enabling information exchange. Government statistics agencies, pharmaceutical companies and healthcare organizations have adopted RDF graphs widely.

How graph databases work.

Graphs and graph databases provide graph models to represent relationships in data. They allow users to perform “traversal queries” based on connections and apply graph algorithms to find patterns, paths, communities, influencers, single points of failure, and other relationships, which enable more efficient analysis at scale against massive amounts of data. The power of graphs is in analytics, the insights they provide, and their ability to link disparate data sources.

When it comes to analyzing graphs, algorithms explore the paths and distance between the vertices, the importance of the vertices, and clustering of the vertices. For example, to determine importance algorithms will often look at incoming edges, importance of neighboring vertices, and other indicators.

Graph algorithms—operations specifically designed to analyze relationships and behaviors among data in graphs—make it possible to understand things that are difficult to see with other methods. When it comes to analyzing graphs, algorithms explore the paths and distance between the vertices, the importance of the vertices, and clustering of the vertices. The algorithms will often look at incoming edges, importance of neighboring vertices, and other indicators to help determine importance. For example, graph algorithms can identify what individual or item is most connected to others in social networks or business processes. The algorithms can identify communities, anomalies, common patterns, and paths that connect individuals or related transactions.

Advantages of graph databases.

The graph format provides a more flexible platform for finding distant connections or analyzing data based on things like strength or quality of relationship. Graphs let you explore and discover connections and patterns in social networks, IoT, big data, data warehouses, and also complex transaction data for multiple business use cases including fraud detection in banking, discovering connections in social networks, and customer 360. Today, graph databases are increasingly being used as a part of data science as a way to make connections in relationships clearer.

Because graph databases explicitly store the relationships, queries and algorithms utilizing the connectivity between vertices can be run in subseconds rather than hours or days. Users don’t need to execute countless joins and the data can more easily be used for analysis and machine learning to discover more about the world around us.

Graph databases are an extremely flexible, extremely powerful tool. Because of the graph format, complex relationships can be determined for deeper insights with much less effort. Graph databases generally run queries in languages such as Property Graph Query Language (PGQL). The example below shows the same query in PGQL and SQL.

Because graphs emphasize relationships between data, they are ideal for several different types of analyses. In particular, graph databases excel at:

* Finding the shortest path between two nodes
* Determining the nodes that create the most activity/influence
* Analyzing connectivity to identify the weakest points of a network
* Analyzing the state of the network or community based on connection distance/density in a group

# Graphs, Data sources

## Social:

### IWIW

Number of nodes: 190

Number of edges: 1216



The graph above represents the connections of a person on the IWIW social platform. This is the only data which has not been anonymized. The nodes represent people and the edges the friendships. An attribute file is available for this graph, holding information about names, locations and other valuable information. With this data, a possibility was given to check the efficiency of the implemented community detector algorithms. The results showed that the communities appeared in certain period of time.

### Facebook

This dataset is collected from the website of Stanford University. Despite the data is anonymized, this source opened a possibility to handle large social type of graphs and examine the structure of them. Based on what is known about the IWIW graph, the same patterns can be deduced in other anonymized graphs like the ones collected from Facebook or Twitter.

The datasets from Stanford are using a wide range of delimiters to separate the nodes of the edges, so most of them are not eligible to open with the Sixtep software that has been provided to use during the research. The application had to be implemented that way to be able to use any delimiter given by the user. This parameter comes from outside, and not depends on the implementation.

Number of nodes: 128

Number of edges: 100



Figure 1 Newman-Girvan run on the slice of a FB graph

### Karate club graph

A social network of a karate club was studied by Wayne W. Zachary. The network became a popular example of community structure in networks after its use by Michelle Girvan and Mark Newman. It captures 34 members of a karate club, documenting links between pairs of members who interacted outside the club. During the study a conflict arose between the administrator and instructor, which led to the split of the club into two. Half of the members formed a new club around the instructor; members from the other part found a new instructor or gave up karate. Based on collected data Zachary correctly assigned all but one member of the club to the groups they actually joined after the split. The coloring of the graph represents the two new community. [5]

Number of nodes: 34

Number of edges: 78



Figure : The well-known karate club community is divided into 2 main part due to a conflict of interest.

## Transaction:

### OTP graph

This graph has been provided by the *Department of Computal Optimization* from the University of Szeged and despite the data is covered with a total anonymization, it is strictly confidential. The graph originally has been cut into 3 pieces and even smaller slices were made from the main partitions to make the detection easier even to the naked eye. It is not possible to decide obviously whether the graph is social or transactional type, there are areas with each kind. See layers after condensation.

Main partitions:

Relevant first quarter year:

* Nodes: 34992
* Edges: 72440

Relevant second quarter year:

* Nodes: 37008
* Edges: 79098

Relevant third quarter year:

* Nodes: 39155
* Edges: 86110



Figure 3 Largest connected components clustered by MCL in a slice of OTP first quarter graph

### Word-Graph

# Algorithms, experiences

## Community detection algorithms

### Newman-Girvan



The Girvan-Newman algorithm for the detection and analysis of community structure relies on the iterative elimination of edges that have the highest number of shortest paths between nodes passing through them. By removing edges from the graph one-by-one, the network breaks down into smaller pieces, so-called communities. The algorithm was introduced by Michelle Girvan and Mark Newman. The idea was to find which edges in a network occur most frequently between other pairs of nodes by finding edges betweenness centrality. The edges joining communities are then expected to have a high edge betweenness. The underlying community structure of the network will be much more fine-grained once the edges with the highest betweenness are eliminated which means that communities will be much easier to spot.

The Girvan-Newman algorithm can be divided into four main steps:

1. For every edge in a graph, calculate the edge betweenness centrality.
2. Remove the edge with the highest betweenness centrality.
3. Calculate the betweenness centrality for every remaining edge.
4. Repeat steps 2-4 until there are no more edges left.

### Maximized modularity

A képen égbolt látható

Automatikusan generált leírás

This algorithm finds communities in graph using Clauset-Newman-Moore greedy modularity maximization. This method currently does not consider edge weights. Greedy modularity maximization begins with each node in its own community and joins the pair of communities that most increases modularity until no such pair exists.

Source code:

<https://networkx.org/documentation/stable/_modules/networkx/algorithms/community/modularity_max.html>

Modularity

One of the most sensitive detection methods is optimization of the quality function known as modularity over the possible divisions of a network, but direct application of this method using, for instance, simulated annealing is computationally costly. A community structure in a network corresponds to a statistically surprising arrangement of edges, can be quantified using the measure known as modularity [9]. The modularity is, up to a multiplicative constant, the number of edges falling within groups minus the expected number in an equivalent network with edges placed at random. The modularity can be either positive or negative, with positive values indicating the possible presence of community structure [10].

Modularity is defined in [11] as

Where m is the number of edges, A is the adjacency matrix of G, is the degree of i, γ is the resolution parameter, and is 1 if i and j are in the same community, else 0.

According to [12] (and verified some algebra) this can be reduced to

Where the sum iterates over all communities c, m is the number of edges, is the number of intra-community links for community c, is the sum of degrees of the nodes in community c, and γ is the resolution parameter.

The resolution parameter sets an arbitrary tradeoff between intra-group edges and inter-group edges. More complex grouping patterns can be discovered by analyzing the same network with multiple values of gamma and then combining the results [13]. That said, it is very common to simply use gamma=1. More on the choice of gamma is in [14].

This NetworkX version has been used in the community detector repository. The parameters are the following: G represent the NetworkX graph. Communities are a list or iterable of set of nodes. These node sets must represent a partition of G’s nodes. Weight is an edge attribute that holds the numerical value used as a weight. It is an optional parameter, if the value is None or an edge does not have that attribute, then that edge has weight 1. Resolution is an optional parameter as well. If resolution is less than 1, modularity favors larger communities. Greater than 1 favors smaller communities. The function returns Q, the modularity of the partition. In case of the communities are not a partition of G, the function raises NotAPartition exception.

The second formula is the one actually used in calculation of the modularity. For directed graphs the second formula replaces with .

### Cliques

Clique problem.

In computer science, the clique problem is the computational problem of finding cliques (subsets of vertices, all adjacent to each other, also called complete subgraphs) in a graph. It has several different formulations depending on which cliques, and what information about the cliques should be found. Common formulations of the clique problem include finding a maximum clique (a clique with the largest possible number of vertices), finding a maximum weight clique in a weighted graph, listing all maximal cliques (cliques that cannot be enlarged) and solving the decision problem of testing whether a graph contains a clique larger than a given size.

The clique problem arises in the following real-world setting. Consider a social network, where the graph's vertices represent people, and the graph's edges represent mutual acquaintance. Then a clique represents a subset of people who all know each other, and algorithms for finding cliques can be used to discover these groups of mutual friends. Along with its applications in social networks, the clique problem also has many applications in bioinformatics, and computational chemistry.

Most versions of the clique problem are hard. The clique decision problem is NP-complete (one of Karp's 21 NP-complete problems). The problem of finding the maximum clique is both fixed-parameter intractable and hard to approximate. Listing all maximal cliques may require exponential time as there exist graphs with exponentially many maximal cliques. Therefore, much of the theory about the clique problem is devoted to identifying special types of graphs that admit more efficient algorithms, or to establishing the computational difficulty of the general problem in various models of computation.

To find a maximum clique, one can systematically inspect all subsets, but this sort of brute-force search is too time-consuming to be practical for networks comprising more than a few dozen vertices. Although no polynomial time algorithm is known for this problem, more efficient algorithms than the brute-force search are known. For instance, the Bron–Kerbosch algorithm can be used to list all maximal cliques in worst-case optimal time, and it is also possible to list them in polynomial time per clique. [19]

Clique maximization:

For each node n, a maximal clique for n is a largest complete subgraph containing n. The largest maximal clique is sometimes called the maximum clique.

This function returns an iterator over cliques, each of which is a list of nodes. It is an iterative implementation, so should not suffer from recursion depth issues.

This function accepts a list of nodes and only the maximal cliques containing all of these nodes are returned. It can considerably speed up the running time if some specific cliques are desired.

A list output of the function find\_cliques(G) has been used to obtain all maximal cliques. However, in the worst-case scenario, the length of this list can be exponential in the number of nodes in the graph. This function avoids storing all cliques in memory by only keeping current candidate node lists in memory during its search.

This implementation is based on the algorithm published by Bron and Kerbosch (1973) [6], as adapted by Tomita, Tanaka and Takahashi (2006) [7] and discussed in Cazals and Karande (2008) [8].

This algorithm ignores self-loops and parallel edges, since cliques are not conventionally defined with such edges.



Figure It is important to determine the size of the cliques should be detected. For example, a clique with size 3 has lower importance than 5 or above.

## Clustering algorithms on transaction networks

### Markov-chain



The MCL algorithm is short for the Markov Cluster Algorithm, a fast and scalable unsupervised cluster algorithm for graphs (also known as networks) based on simulation of (stochastic) flow in graphs. The algorithm was invented/discovered by Stijn van Dongen at the Centre for Mathematics and Computer Science (also known as CWI) in the Netherlands. [4]

### H-avoiding coloring

Embeddedness of bipartite subgraphs in transaction networks

Certain bipartite graphs, for example pollinator networks or trade networks suggest the presence of different structures, like the notion of embeddedness. The vertices of each color class can be ordered and the smaller ranked vertex neighbourhood contains the neighbourhood of any higher ranked one. A binary matrix *A* is fully nested if its rows and columns can be reordered such that the ones are in echelon form. Let be the bipartite graph whose adjacency matrix is A. Then A being fully nested is equivalent to satisfying embeddedness.

Let X (the columns) and Y (the rows) be the bipartition of . The matrix A and the graph are each said to be k-nested with respect to X if X can be partitioned as such that all subgraphs spanned by are fully nested for . The quantity of interest for any is smallest k for which is k-nested.

Given a “forbidden graph” H and an integer k, an H-avoiding k-coloring of a graph G is a k-coloring of the vertices of G such that no maximal H-free subgraph of G is monochromatic.

A monochromatic graph is a colored graph (either vertex-colored or edge-colored, depending on the context) in which each of the vertices or edges is assigned the same color.



Figure . Proper 2K2 avoiding coloring of a bipartite graph

*Colormap: {'1': 0.82, '2': 0.44999999999999996, '4': 0.72, '6': 0.94, '8': 0.15000000000000002, '10': 0.15000000000000002, '3': 0.82, '5': 0.82}*

A new kind of clustering of general (that is, not necessarily bipartite) transaction graphs has been presented via a certain class of proper colorings. The clusters are the color classes, since no edges are desired inside a cluster. The structure of the edges is restricted between the pairs of classes. The above examples suggest that in some cases there should be a fully nested or, equivalently, embeddedness relation among any two-color classes. This notion is generalized to an arbitrary host graph G and a forbidden bipartite subgraph H as follows.

Definition 1. Fix a bipartite graph H. A proper coloring of a graph G is an H-avoiding coloring if the union of any two-color classes spans an induced H-free graph. Let be the minimum number of colors in an H-avoiding coloring of G.

Observation 2. For any graphs H and G, . If G is H-free, then

The computation of is NP-hard for some graphs and polynomially computable for others. The most interesting case is, when , gives back embeddedness as described above. For these generalized chromatic numbers some theoretical extremal results have been derived as well as results on complexity.

## Other algorithms

### Condensation

Condensed graphs



Each community/cluster has been contracted to single node.

If there is a path between the modules, the algorithm puts an edge between the nodes.

The algorithm works well, if the source graph is strongly connected.



### Small-worldness

A small-world network is a type of mathematical graph in which most nodes are not neighbors of one another, but the neighbors of any given node are likely to be neighbors of each other and most nodes can be reached from every other node by a small number of hops or steps. Specifically, a small-world network is defined to be a network where the typical distance L between two randomly chosen nodes (the number of steps required) grows proportionally to the logarithm of the number of nodes N in the network, that is:

while the clustering coefficient is not small. In the context of a social network, this results in the small world phenomenon of strangers being linked by a short chain of acquaintances. Many empirical graphs show the small-world effect, including social networks, wikis such as Wikipedia, gene networks, and even the underlying architecture of the Internet. [19]

Small-world functions are used for estimating the small-worldness of graphs. A small world network is characterized by a small average shortest path length and a large clustering coefficient. Small-worldness is commonly measured with a coefficient, which compares the average clustering coefficient and shortest path length of a given graph against the same quantities for an equivalent random or lattice graph.

The coefficient is defined as:

Where C and L are respectively the average clustering coefficient and average shortest path length of the graph. [3]

The graph is commonly classified as small-world if .

### Colormap

# Measuring, results

Experiences:

The NetworkX implementations of the algorithms have been used for the python script. The code of the algorithms can be found in the coloring.py file. The colormap and the coloring logic is self-implemented and it can be found in the Utils.py file.

The Newman-Girvan algorithm is fairly slow on medium sized graphs, but the result is more accurate and all the nodes are classified.

Measuring results:

## Modularities

### Network: Facebook, 0.edges dataset

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Application | Community detector app | | | |
| Algo | NG | NG-comm\* | Markov | Maximized modularity |
| Nr. of colors | 2 | 30 | 24 | 8 |
| Modularity | 0.3615 | (6) 0.2523 | -0.8080 | **0.4429** |
| (20) 0.3729 |
| (30) 0.4136 |
| (35) 0.4129 |
| (40) 0.4111 |
| (50) 0.4043 |

|  |  |  |  |
| --- | --- | --- | --- |
| Application | Sixtep | | |
| Algo | communities | Markov | Maximized modularity |
| Nr. of colors | 208 | 7 | 8 |
| Modularity | Cannot be calculated\*\* | 0.3306 | **0.3891** |



Figure 6: Facebook communities detected by greedy modularity algorithm

Number of nodes: 324

Number of edges: 2514

The value of the number of edges is different from the value given by the Sixtep software. The reason behind this is that the edges are represented in both direction in the text file. In conclusion, the value is double.

\* The numbers before the modularity values represent the number of communities the algorithm took.

\*\* The communities that has been provided by the Sixtep software are not a valid partition of the graph, therefore the NetworkX modularity function cannot be run against the data.

### Network: transaction graphs of the OTP database

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Application | Community detector app | | | | | |
|  | Newman-Girvan | | Markov | | Greedy modularity | |
|  | # colors | Modularity | # colors | Modularity | # colors | Modularity |
| OTP 100 | 10 | 0.6981 | 14 | 0.5413 | 9 | 0.7042 |
| First 600 edges of first quarter year largest cc | 19 | **0.7570** | 80 | 0.5725 | 15 | 0.7632 |
| First quarter year 1k | 19 | 0.7977 | 113 | 0.6182 | 20 | 0.7981 |
| Middle slice | - | - | 3810 | 0.5509 | 92 | 0.8227 |
| total | - | - | 0.3900 | | - | |
| Second quarter year | - | - | 0.3794 | | - | |
| Third quarter year | - | - | 0.3690 | | - | |



Figure First 600 edges of first quarter year largest cc, 19 with 0.7570 mod



|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Application | Sixtep | | | | |
|  | communities | Markov | | Greedy modularity | |
|  | # colors | # colors | Modularity | # colors | Modularity |
| OTP 100 | 2 | 9 | 0.6128 | 12 | 0.7580 |
| First 600 edges of first quarter year largest cc | 39 | 25 | 0.7346 | 19 | 0.7697 |
| First quarter year 1k | 56 | 39 | 0.7751 | 25 | 0.7965 |
| Middle slice | 862 | 1587 | 0.7632 | 879 | 0.8349 |
| First quarter year | 3583 | 1688 | 0.5792 | - | |
| Second quarter year | 3814 | 1587 | 0.5764 | - | |
| Third quarter year | 4303 | 1738 | 0.5707 | - | |

First quarter year:

* Number of nodes: 34992
* Number of edges: 72440, 76668 in Sixtep

Second quarter year:

* Number of nodes: 37008
* Number of edges: 79098, 83926 in Sixtep

Third quarter year:

* Number of nodes: 39155
* Number of edges: 86110, 91862 in Sixtep

OTP midslice of a first quarter year:

* Number of nodes:18032
* Number of edges: 25686

First 1K edges of first quarter year largest cc:

* Number of nodes:675
* Number of edges: 1000

First 600 edges of first quarter year largest cc:

* Number of nodes: 398
* Number of edges: 600 in Sixtep

First 100 edges of first quarter year largest cc:

* Number of nodes: 96
* Number of edges: 100

Is the OTP graph directed?

Usually, the difference between the value of the edges represents the direction of the graph. X -> Y and Y -> Y show that the nodes are mutually connected to one another. Difference between edges caused by graph vs. largest cc.The smaller OTP graph has been represented by 96 nodes, the first quarter year by 675 nodes and the middle slice is about 18032 nodes. The smaller one is a subset of the first quarter year. The total amount of nodes in this dataset is 34992. The graph of the second quarter year contains 37008 and the third one has 39155. At this size there is not much sense to visualize the graphs in 2D. No information can be gained with naked eyes or manual clustering.



Figure 8. First quarter year of the OTP transactions clustered by Markov chain algorithm. 675 nodes have been colorized.

### Wordgraph

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Application | Community detector app | | | | | |
|  | Newman-Girvan | | Markov | | Greedy modularity | |
|  | # colors | Modularity | # colors | Modularity | # colors | Modularity |
| wordgraph | - | - | 3786 | 0.2241 | 106 | 0.5349 |
| 1200 | 17 | **0.8663** | 52 | 0.7264 | 19 | 0.8661 |

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Application | Sixtep | | | | |
|  | communities | Markov | | Greedy modularity | |
|  | # colors | # colors | Modularity | # colors | Modularity |
| wordgraph | null | 236 | 0.4303 | 114 | 0.5303 |

* Number of nodes: 11381
* Number of edges: 31784



Figure slice of wordgraph, colored by NG

OTP transaction leveling by condensation

This graph is a slice of the OTP transactions from the first quarter year.



And the condensated one…



This is interesting because technical graphs usually represent tree-like structure (like said before), but this seems more like a social graph. Some of the layers in the otp graph look like communities. Originally the suppliers supposed to be in a competitive relation by restrict the edges between them.



But based on this, some cooperation can be supposed.

The modularity of the original graph is 0.7632.

Patterns!

Efficiency, solutions, etc.

# Further studies:

Make a ui to represent small graphs, but the software should provide opportunity to manually color graph nodes, modify the location of the nodes.

This solution is scalable and modularized which makes further implementation more easier. The code and be found in the github repository liked below:

<https://github.com/daniellanikov/Community-detector>

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* Cliques
* Small-world graphs