

Finding community structure

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1 Introduction to `igraph`'s community detection algorithms

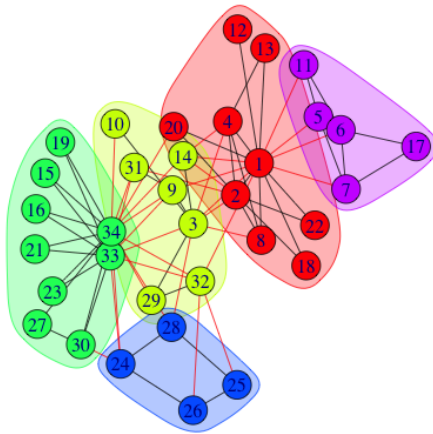
In this session you will run and compare different community finding algorithms. In the `igraph` package there are a few already implemented, including some we have seen in theory class:

- `edge.betweenness.community` [Newman and Girvan, 2004]
- `fastgreedy.community` [Clauset et al., 2004]: modularity greedy optimization method.
- `label.propagation.community` [Raghavan et al., 2007]: This is a fast, nearly linear time algorithm for detecting community structure in networks. It works by labeling the vertices with unique labels and then updating the labels by majority voting in the neighborhood of the vertex.
- `leading.eigenvector.community` [Newman, 2006]
- `multilevel.community` [Blondel et al., 2008]: This is a multi-level modularity optimization algorithm (the Louvain method).
- `optimal.community` [Brandes et al., 2008]: Works by maximizing the modularity measure over all possible partitions.
- `spinglass.community` [Reichardt and Bornholdt, 2006]: tries to find communities in graphs via a spin-glass model and simulated annealing.
- `walktrap.community` [Pons and Latapy, 2005]: tries to find densely connected subgraphs (communities) in a graph via random walks. The idea is that short random walks tend to stay in the same community.

- `infomap.community` [Rosvall and Bergstrom, 2008]: Find community structure that minimizes the expected description length of a random walker trajectory.

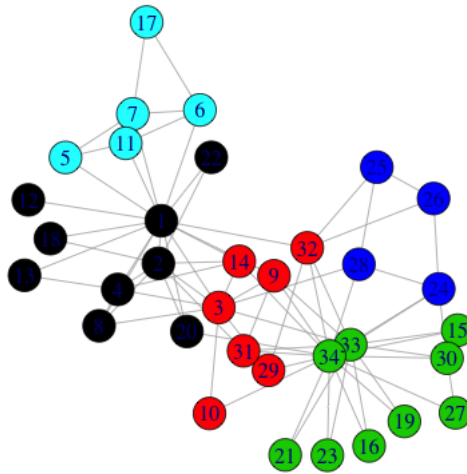
All of these methods return a `communities` object, which you can then use to explore, plot, and compute metrics on. As an example, consider the following snippet of code:

```
> karate <- graph.famous("Zachary")
> wc <- walktrap.community(karate)
> modularity(wc)
[1] 0.3532216
> membership(wc)
[1] 1 1 2 1 5 5 5 1 2 2 5 1 1 2 3 3 5 1 3 1 3 1 3 4 4 4 3 4 2 3 2 2 3 3
> plot(wc, karate)
```



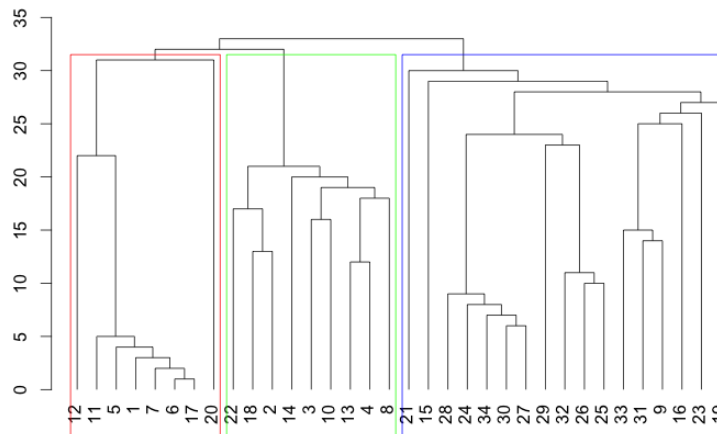
An alternative way of plotting communities without the shaded regions is:

```
> plot(karate, vertex.color=membership(wc))
```



For those algorithms that output communities with hierarchical structure, this information can be visualized using the `dendPlot` function, which displays the corresponding dendrogram:

```
> karate <- graph.famous("Zachary")
> fc <- fastgreedy.community(karate)
> dendPlot(fc)
```



Next, to do an agglomerative hierarchical clustering of 34 stocks belonging to the main Spanish market index IBEX, see the R script in R Example 3.5 at http://computationalfinance.lsi.upc.edu/?page_id=74 Use the dataset `Ibex0809` provided. It contains the 34 stocks' series of daily returns (as columns), observed from 1/12/2008 to 1/2/2009.

Be aware the underlying network of dissimilarity is a complete graph, hence modularity optimization methods most likely give not so good community partitions, e.g. all singleton clusters or a single community (Why? Can you give an explanation and propose a fix?).

2 Tasks

2.1 Task 1

Write an R function that, given an undirected graph, it outputs for each available community finding algorithm the value achieved by the output partition for each of the following criteria: 'Triangle Partition Ratio' (TPT), 'expansion', 'conductance' and 'modularity' (see theory lecture notes for their definitions). Notice also that these definitions are for a single community, in order to compute the metrics for the whole network, you can use a weighted average of each of the communities, where the weight is given by the "volume" (i.e. fraction of number of nodes) in each community.

Use your code to find graphs for which different community detection methods beat on different criteria. In particular, you should build matrices like the following for a minimum three different networks of your choice.

	<i>TPT</i>	<i>expansion</i>	<i>conductance</i>	<i>modularity</i>
<code>edge.betweenness</code>				
<code>fastgreedy</code>				
<code>:</code>				
<code>infomap</code>				

Beware that community detection algorithms can be very time consuming, so stay away from large networks. Unless you have a *very* powerful computer.

Regarding what networks to consider, you can be as creative as you want. Simple suggestions could be to use the famous *Zachary* karate network, or generate your own using `igraph`'s generation tools. A way to generate a small network with very clear community structure is, for example:

```
g <- graph.full(10) + graph.full(10)
g <- g + edges(sample(V(g), 10, replace=TRUE))
```

However, in such simple networks the outputs of the community detection al-

gorithms are going to be the same so it is not going to be very interesting. However, you can build up on this idea to generate more complex networks.

Alternatively, you can use networks from network repositories available in the web.

2.2 Task 2

Load the network `wikipedia.gml` provided¹. It is in gml format, which can be imported into `igraph` using the following command

```
read.graph("wikipedia.gml", format="gml")
```

The vertices of this network are wikipedia pages. The label of each vertex is the title of the wikipedia page.

Now use any community detection algorithm. Do you think the communities found make sense? You can use the vertex labels to check this.

3 Deliverables

You have to prepare a report describing your findings and results while solving this lab, especially emphasizing any difficulties you encountered and the solution you found to overcome them.

To deliver: You must deliver the report explained above. The formats accepted for the report are, in principle, pdf, Word, OpenOffice, and Postscript. You also have to hand in the source code in R (or any other language) that you have used, including some minimal comments that can help the reader. This work can be done in pairs; in that case it is enough that one of you submits the work as long as both names are clearly visible in the report.

Procedure: Submit your work through the `raco` platform as a single compressed file.

Deadline: Work must be delivered within 2 weeks from the lab session you attend. Late deliveries risk being penalized or not accepted at all. If you anticipate problems with the deadline, please tell us as soon as possible.

References

[Blondel et al., 2008] Blondel, V. D., Guillaume, J.-l., Lambiotte, R., and Lefebvre, E. (2008). Fast unfolding of community hierarchies in large networks. *Networks*, pages 1–6.

¹Thanks to Lada Adamic for providing this in her course *Social Network Analysis*.

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- [Clauset et al., 2004] Clauset, A., Newman, M. E. J., and Moore, C. (2004). Finding community structure in very large networks. *Physical Review E*.
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