

Algorithm Design And Analysis  
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# Hierarchical Agglomerative Clustering to Extract Communities from Social Network Data

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**Abstract.** Relationships between individuals can not be seen from raw social network data. Can a Java implementation of hierarchical agglomerative clustering be used to extract information from a social network data set to provide knowledge of community relationships? This report describes a Java implementation of an hierarchical agglomerative clustering algorithm. The aim of this project is to carry out an independent study of this algorithmic problem in relation to social network data. The performance of the algorithm is discussed and the project results are presented.

## 1 Introduction

In today's environment of Social Networking websites, such as Facebook, large amounts of statistical data is collected. However, this data is often ambiguous, irregular and difficult to interpret. Using appropriate methods, hidden structures can be identified and information extracted and analysed from this complex networked data (Yuruk, Mete, Xu, & Schweiger, 2009). This project uses one such approach to analyse data from social network distributions by implementing an hierarchical agglomerative community detection algorithm, highlighting the importance of graph theory in data analysis.

## 2 Hierarchical Agglomerative Clustering Algorithm to Compute Social Network Communities

Hierarchical Agglomerative Clustering or HAC takes a bottom-up approach, initially treating each node in the network data set as a singleton cluster (or community) and then successively merging (or agglomerating) pairs of clusters until all clusters have been merged into a single cluster that contains all nodes (Manning, Raghavan, & Schütze, 2008).

Merges are determined by choosing the locally optimal choice at each stage with the hope of finding a global optimum. Therefore, it is a greedy algorithm. The results of hierarchical clustering are often presented in a dendrogram. This report, however, presents clustering with coloured edges between nodes, where the same colour represents belonging to a cluster. In this way, the relationship between nodes can easily be visualized in the graph.

A threshold value is provided to specify at above what value nodes will be merged into a cluster.

### 2.1 Mathematical Analysis

To perform cluster analysis an indicator of "closeness" is needed to decide whether it is possible to merge a specific group of nodes into one cluster. We denote:

$$C = \frac{e}{\frac{n(n-1)}{2}}$$

where  $e$  is the actual number of edges,  $n$  is the actual number of nodes and  $\frac{n(n-1)}{2}$  is the maximum possible edges in one community. We use the ratio between  $e$  and  $\frac{n(n-1)}{2}$  to calculate the closeness between each sub-community, where a larger  $C$  indicates a closer relationship. The largest ratio between actual edges and maximum possible edges is 1 as:

$$0 \leq C \leq 1$$

To analyse each cluster, we compare the result of  $C$  for all communities, and select the largest  $C$  to merge into the cluster and continue further cluster operations. Therefore, the time complexity is  $O(n^3)$ .

## 2.2 Pseudocode Hierarchical Agglomerative Clustering Algorithm

Hierarchical Cluster:

```
cluster(thresh)
    initialize all communities
    run community one
    run community two
    calculate cluster coefficient of community one and two
    merge the largest two communities
    remove the original communitie(s)
    return the new merged community
```

Calculate density:

```
Identity = e/(n*(n-1))/2
For all nodes in communities
    if node1 is connected to node2
        add existing edges
```

```
Density = existing edges / total
```

## 2.3 Design and Implementation

The HAC algorithm was implemented as a Java software application using the Netbeans Integrated Development Environment. The project contains the following eight classes:

Community.java

- Create initial communities, calculate the density and merge communities by selecting the largest density result in each round of comparisons.

DensityQualityFunction.java

- Calculate the coefficients of communities.

Digraph.java

- Implement Graphic User Interface

HClustering.java

- Cluster communities based on different value of identity (from the formula)

LoadGML.java

- Convert gml dataset into an arrayList data structure in java

MainActivity.java

- Implement a command-line User Interface

Node.java

- Create nodes

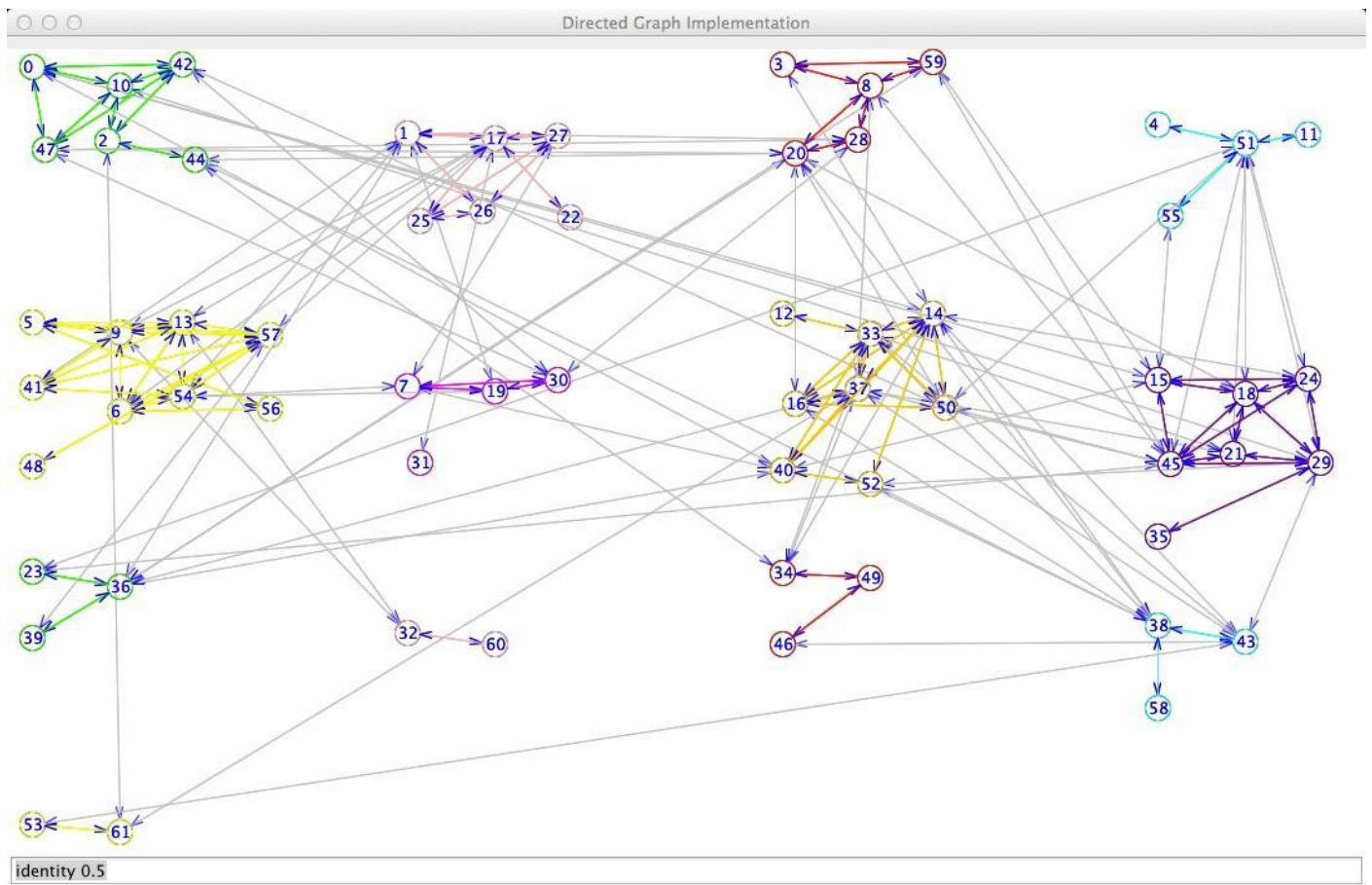
QualityFunction.java

- An Interface for calculating the cluster's coefficients

The data set primarily used for this project was a Graph Modeling Language (.gml) file of a [dolphin social network](#): (Lusseau et al., n.d.) This data set was chosen for the availability of previous clustering results for comparison and for it's New Zealand context.

A second .gml data set, [Zachary's Karate club](#) (Zachary, 1977), was also used to further test the algorithm's accuracy against previous analysis.

## 2.4 Results and Analysis



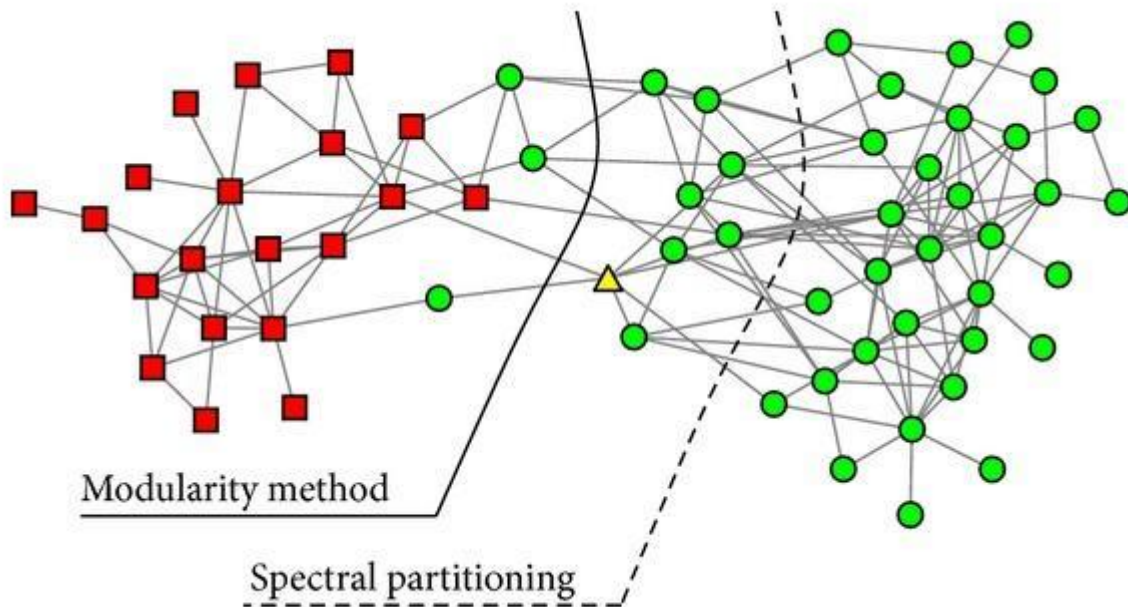
**Figure 1** Clustering of Dolphin set using HAC algorithm with threshold of 0.5

The advantage of this HAC algorithm is that community groups do not need to be defined and these communities can be automatically detected. A disadvantage of this algorithm is that it has a large time complexity of  $O(n^3)$ . The processing time increases polynomially as the number of input nodes increases. This means the algorithm is too slow for large data sets. ("Hierarchical clustering," 2014)

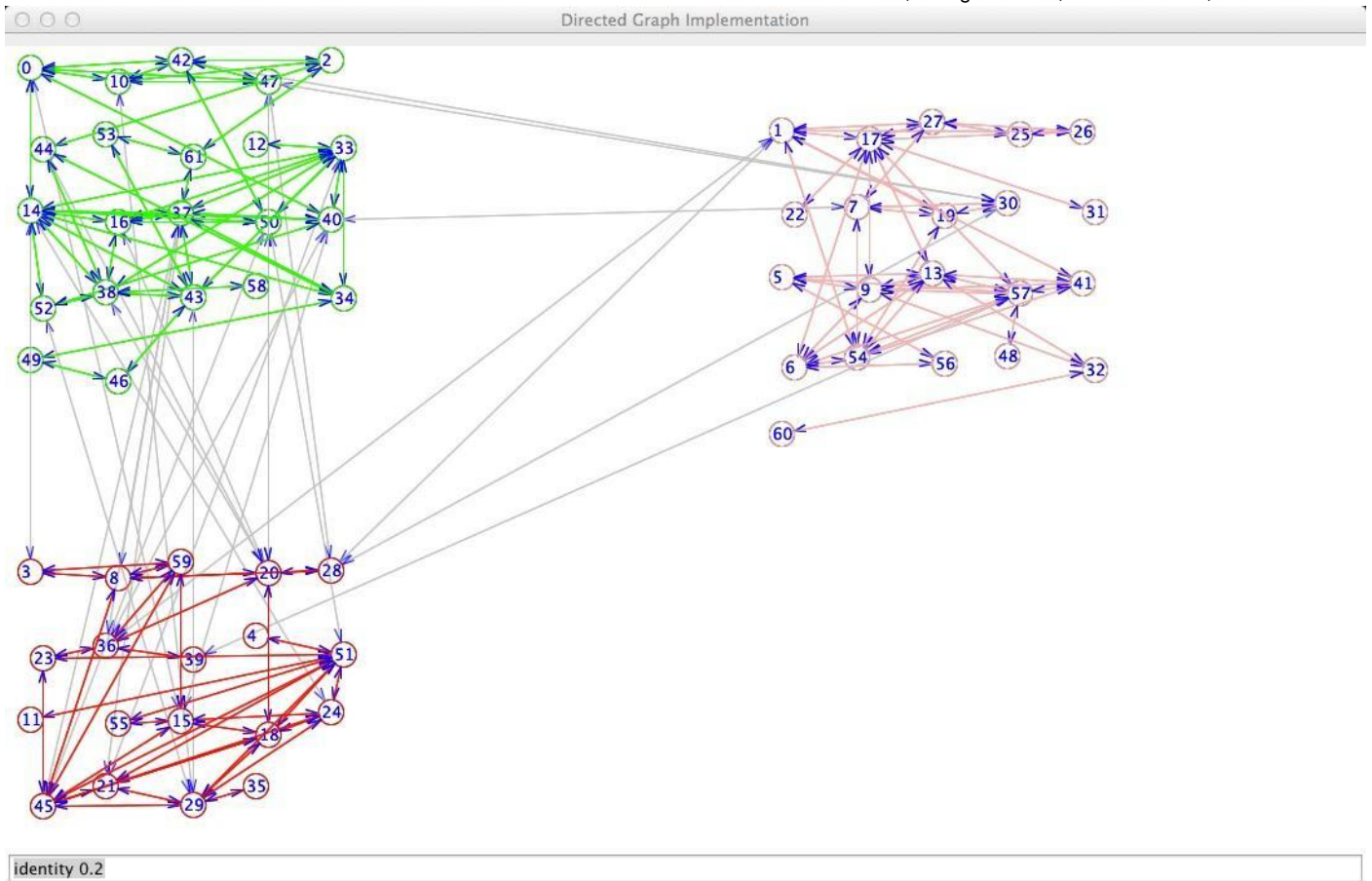
The dolphin set is a small set of 62 nodes, small enough to gain insight into the data set by hand. This is too small to see the full advantage of using an automated algorithm. The advantages of the automated algorithm would better be seen on large data sets, where hand processing would be practically impossible.

### 3 Performance Analysis of Hierarchical Agglomerative Clustering

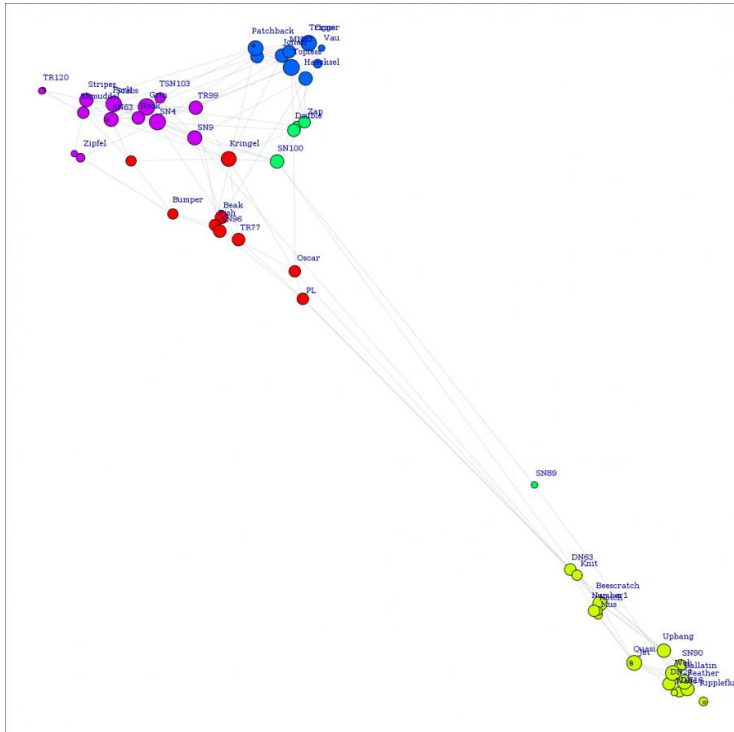
The following figures provide examples of previous analysis network data sets. These are presented as a comparison of how the implemented HAC algorithm compares with previous analyses.



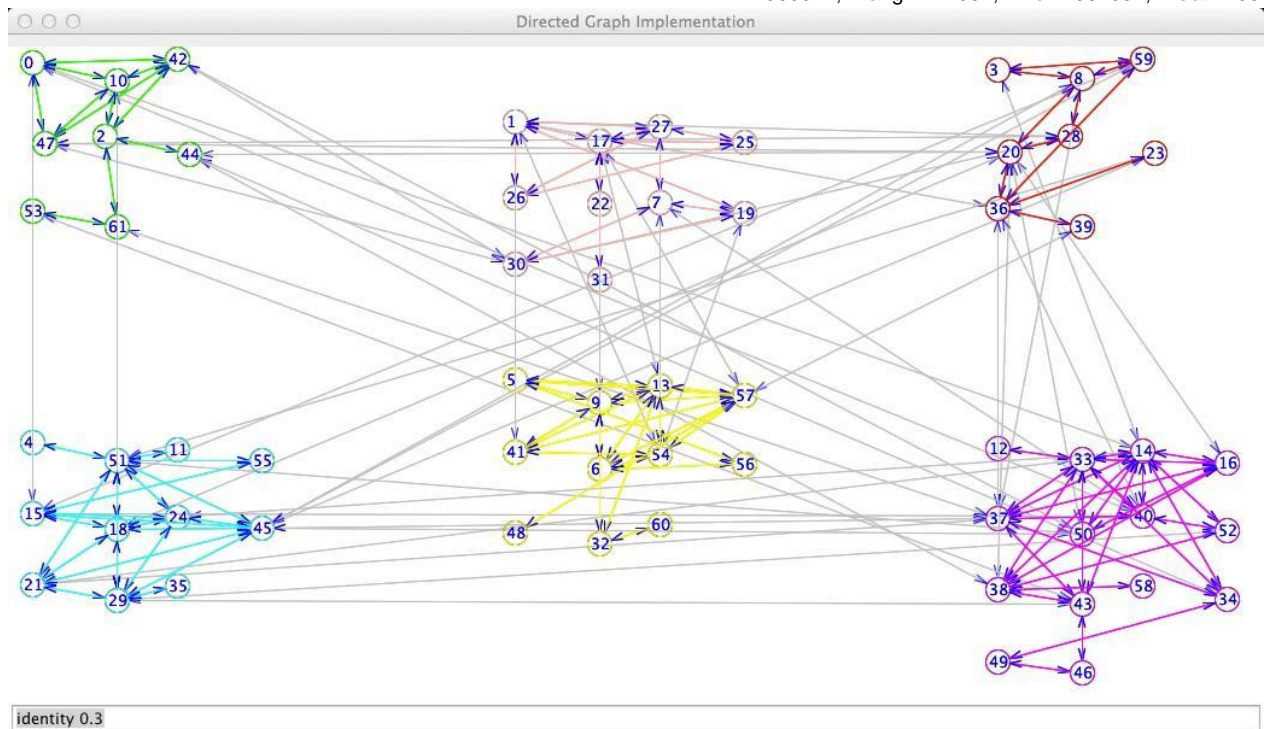
**Figure 2** Clustering of Dolphin set using centrality divided clustering (Wu, Qi, Fuller, & Zhang, 2013)



**Figure 3** Clustering of Dolphin set using HAC algorithm with threshold of 0.2

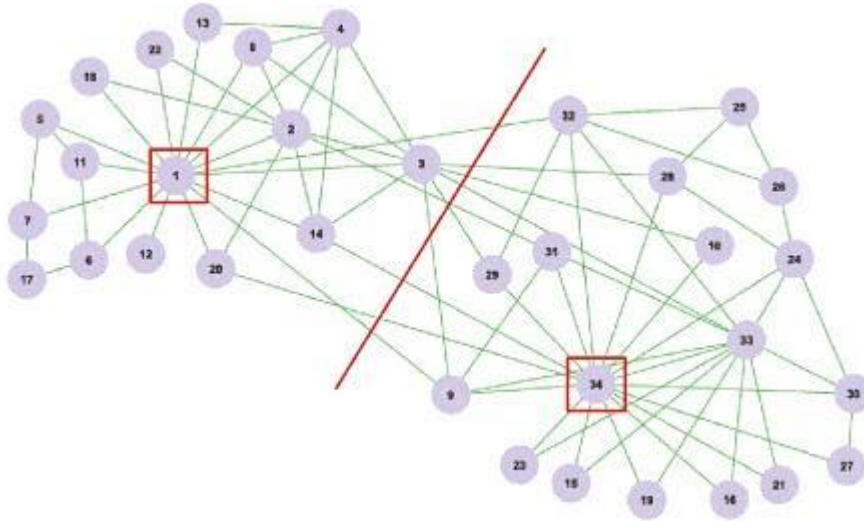


**Figure 4** Clustering of Dolphin data set using multi-level graph clustering (Rotta, 2008)

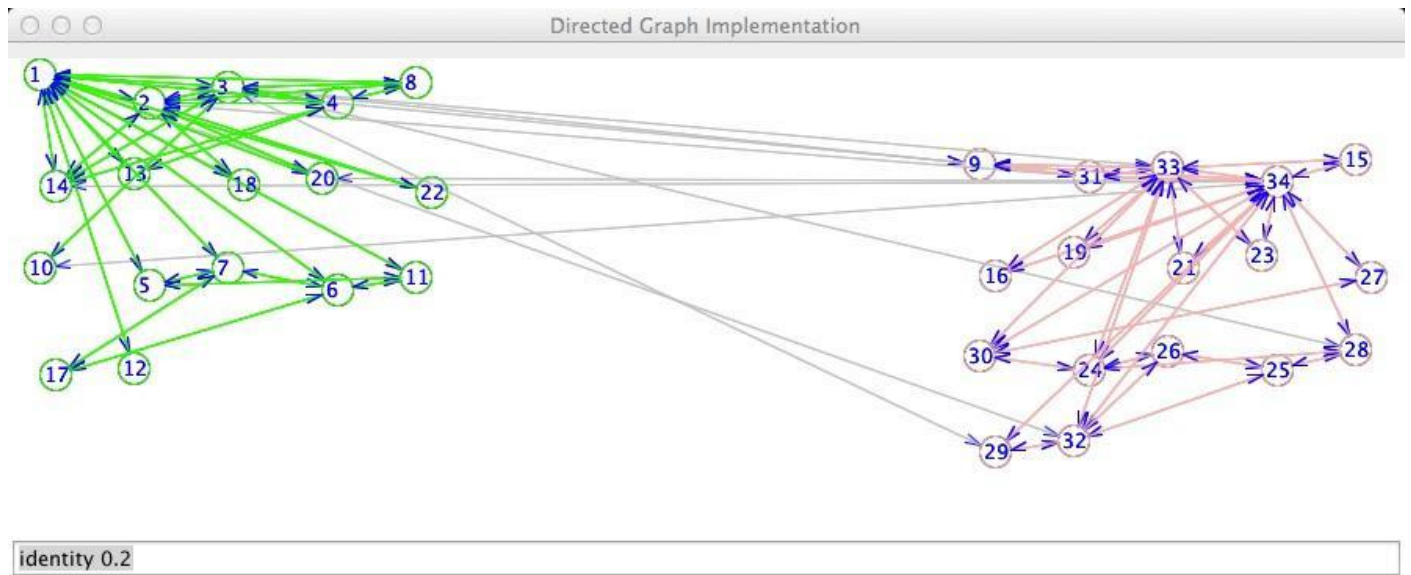


**Figure 5** Clustering of Dolphin set using HAC algorithm with threshold of 0.3





**Figure 6** Clustering of Zachary's Karate club using centrality method (Combinatorial Optimization and Applications - 4th International Conference, COCOA 2010, Kailua-Kona, n.d.)



**Figure 7** Clustering of Zachary's Karate club using HAC algorithm with threshold of 0.2

In this comparison of Zachary's Karate data set, two communities and two influential nodes, 1 and 34 can be seen in both the example graph (**Figure 6**) and the HAC algorithm graph (**Figure 7**)

## 4 Conclusions

To summarise, an hierarchical agglomerative clustering algorithm was successfully implemented as a Java application. Using this application, communities within a complex network dataset can automatically be detected and visualised. Future implementations could provide an easier interface to select different data sets. A more flexible .gml parser could also be implemented. Further directions for investigation could include measuring the processing time for (large data set) input values to confirm the time complexity experimentally.

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