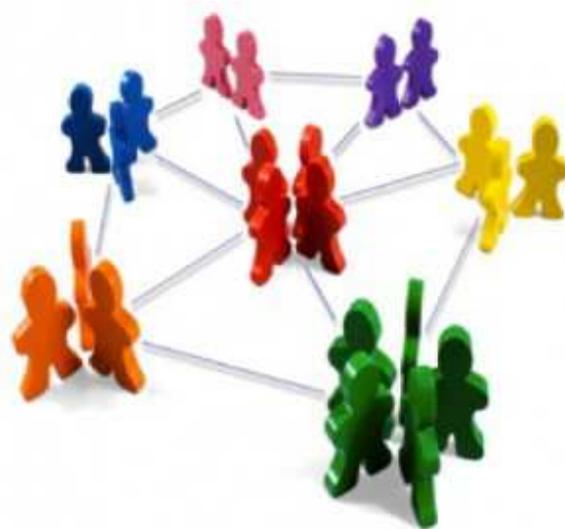


# **Quick Start Instructions for the Community Detection Toolbox v. 0.9**

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## 1 Installation

Installation of the CDTB is extremely simple. You can download the file CDTB.zip from <http://mathworks.com>. When the download is complete, unzip the file in a folder of your choice. From now on we will assume that it is the folder `c:\CDTB`. When unzipped, the folder will contain several subfolders, e.g., `c:\CDTB\Algorithms`, `c:\CDTB\Auxiliary` etc. Now you are ready to go.

## 2 A Simple Demo

Start Matlab, go to `c:\CDTB` and in the command line type

```
>> PathAdd
```

and hit [Enter]. This will add to the *Matlab Path* the subdirectories which contain the Matlab \*.m files which do the actual community detection work<sup>1</sup>. In the command line type

```
>> CDTBDemo01
```

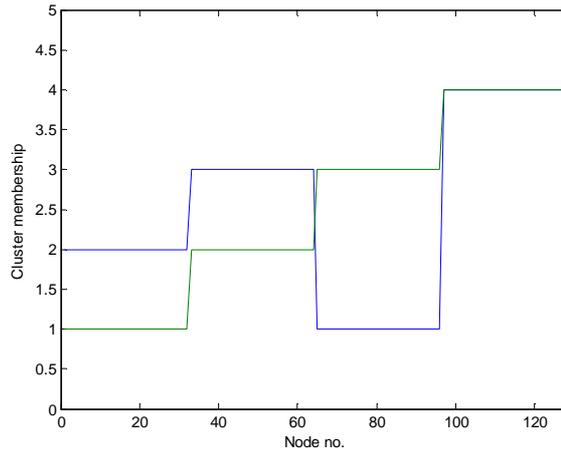
and hit [Enter]. The *Matlab script* CDTBDemo01.m will run and in the command window you will get a message which says

```
The NMI metric between V0 and Vest is 1
```

NMI is the *normalized mutual information* index of partition (i.e., clustering) similarity. It takes values in the interval [0,1]. The maximum value 1 indicates maximum similarity, i.e., identity. The partitions compared are  $V_0$ , the true partition, and  $V_{est}$ , the partition estimated by a *modularity maximization algorithm*. Hence in this experiment  $V_{est}$  is identical to  $V_0$ . This can also be seen in the plot which Matlab presented. It looks like this

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<sup>1</sup> Note that the new path is not saved, so you must retype PathAdd every time you start a new Matlab session. Or, you can go to the Matlab menu File/Set Path and click the button Save, which will make Matlab memorize the path for all future sessions.



**Figure 2.1**

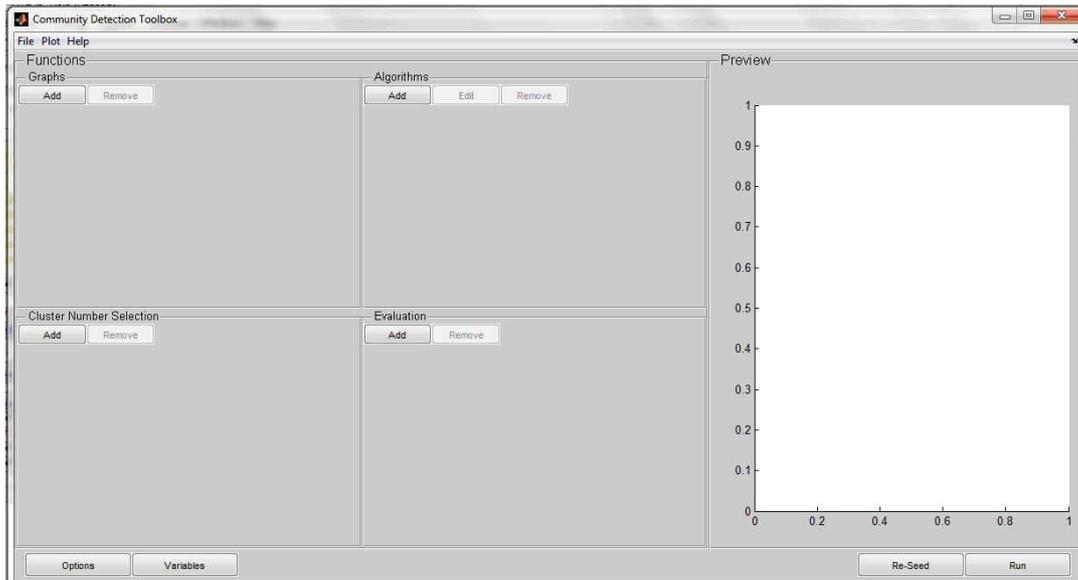
The horizontal axis gives the node id. numbers. The demo utilized a *Girvan-Newman* graph [NewmanGirvan2004] with 128 nodes. These nodes originally were partitioned (by their edge patterns) into four *communities* (i.e., clusters) as follows: nodes 1, 2, ..., 32 go into community no.1, nodes 33, 34, ..., 64 go into community no.2 and so on; this partition is  $V_0$  and is plotted by the green line in Fig.1.  $V_{est}$ , on the other hand, assigns nodes 1, 2, ..., 32 to community no.2, nodes 33, 34, ..., 64 to community no.3 and so on. While the community *labels* are different between  $V_0$  and  $V_{est}$ , the actual division into clusters is the same. In short, the modularity maximization algorithm obtained the correct solution, modulo a relabeling of clusters.

### **3 The GUI**

Let us also run a simple demo of the Graphical User Interface (GUI). In the Matlab command line type

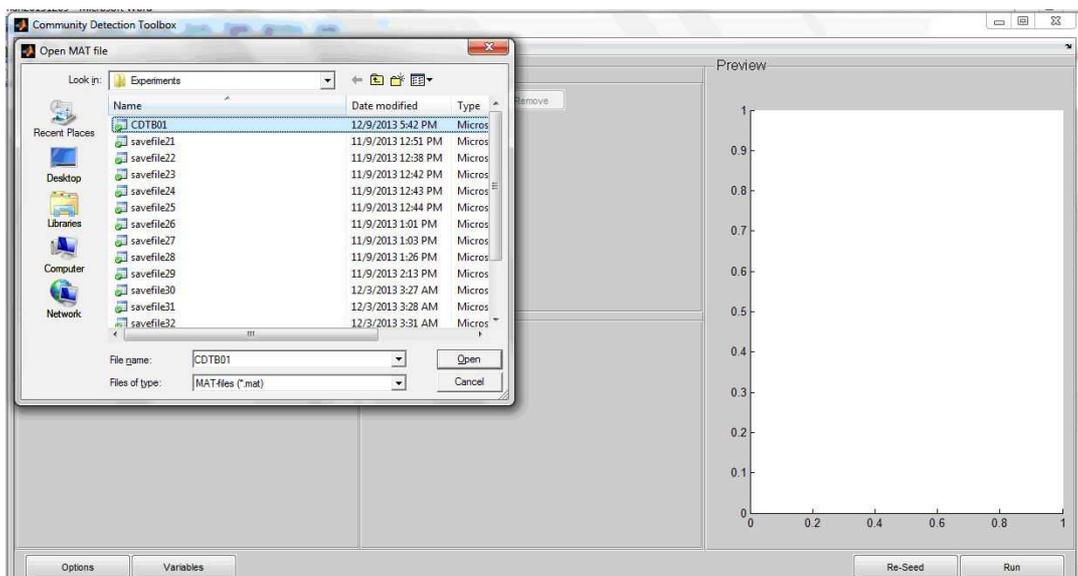
```
>> gui
```

and hit [Enter]. You will get the following standard Matlab GUI.



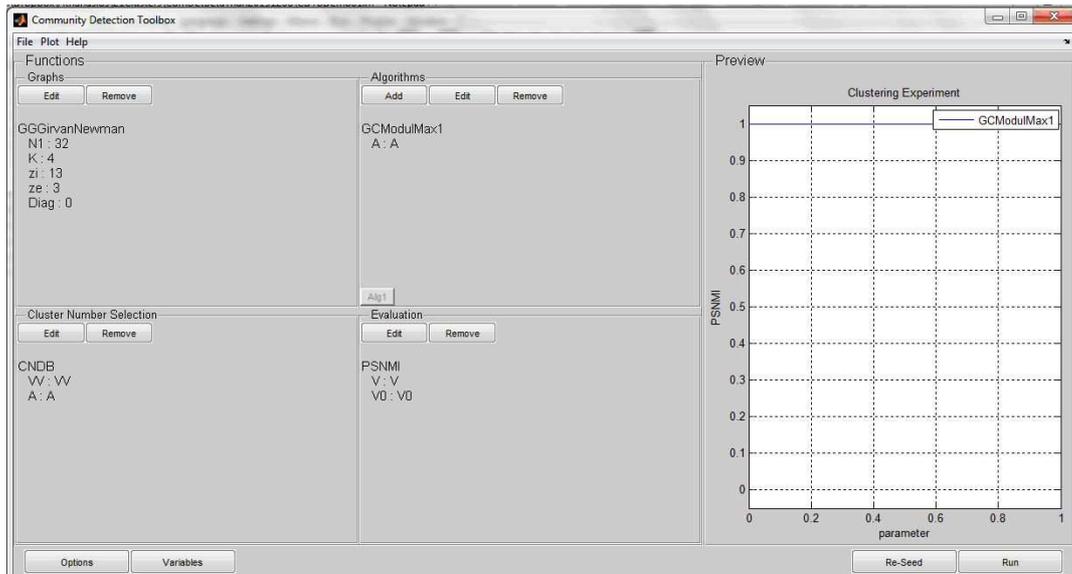
**Figure 2.2**

The GUI, makes it easy to design and perform community detection experiments. To perform exactly the same experiment that CDTBDemo01 performed, do the following. Go to the GUI menu File/Import Data. A standard *File Dialog* will open, like this



**Figure 2.3**

Select the file CDTB01 and click on the Open button. The GUI will change and now look like this:



**Figure 2.4**

Now click on the Run button. A *progress bar* will flash for a while and then you will get a plot (actually a straight line). The GUI has run the experiment specified by the choices appearing in Fig. 2.4. You can access these results by selecting the menu option `Plot/Results to CLI`: if you switch to the Matlab command line (it is always available, e.g. by `[Alt]-[Tab]`-ing) you will see that a new variable has been loaded in your *Matlab workspace*, called `Results`. We will later discuss which results are contained in `Results`. We can now take a break.