# Matlab Tools for Network Analysis (2006-2011)

This toolbox was first written in 2006. The last version, posted here, is from November 2011. These routines are useful for someone who wants to start hands-on work with networks fairly quickly, explore simple graph statistics, distributions, simple visualization and compute common network theory metrics. The code is not object-oriented, and should be easy to use, read and improve upon. There is some overlap with the <a href="mailto:grTheory toolbox">grTheory toolbox</a> and the <a href="mailto:MatlabBGL">MatlabBGL</a> library published on <a href="mailto:MatlabBGL">MatlabBCL</a> contact gergana at alum dot mit.edu if you notice errors or have pointers to better routines.

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## **Current version**

The new and current version of this code is available in Octave (GNU Octave Version 3.4.0 and Gnuplot 4.2.5) <a href="https://examples.com/here.com

## **Archived Version (frozen Nov. 2011)**

The code is classified by functionality as follows (121 routines total, last version Nov 6, 2011). All functions can be downloaded here.

## Basic network routines (17)

- getNodes.m return the list of nodes for varying graph representations;
- getEdges.m return the list of edges for varying graph representations;
- numnodes.m number of vertices/nodes in the network;
- <u>numedges.m</u> number of edges/links in the network;
- link\_density.m the density of links of the graph;
- selfloops.m number of selfloops, i.e. nodes connected to themselves;
- <u>multiedges.m</u> number of arcs (i,j) with multiple edges across them;
- average degree.m the average degree (# links) across all nodes;
- num conn comp.m number of connected components (using algebraic connectivity);
- find conn comp.m the number of connected components in an undirected graph;
- giant component.m extract the giant component only (undirected graph);
- tarjan.m find the strongly connected components in a directed graph;
- graph\_complement.m the complement graph;
- graph dual.m the graph dual (or line graph, adjoint graph);
- subgraph.m return the subgraph adjacency given the graph and the subgraph nodes;
- leaf nodes.m nodes connected to only one other node;
- leaf edges.m edges with only one adjacent edge;

## Diagnostics (11)

- issimple.m check whether the graph has selfloops and multiple edges;
- isdirected.m directed or undirected graph (right now uses issymmetric.m);
- issymmetric.m check whether a matrix is symmetric;
- <u>isconnected.m</u> check whether a graph is connected;
- isweighted.m determine whether the graph has weighted links;
- isregular.m check whether it's a regular graph;
- iscomplete.m check whether the graph is complete;
- iseulerian.m find out whether it's an eulerian graph;
- istree.m check whether the graph is a tree;
- <u>isgraphic.m</u> check whether a sequence of numbers is graphic;
- isbipartite.m check whether a graph is bipartite;

## **Conversion routines (14)**

- adj2adjL.m, (adjL2adj.m) convert an adjacency matrix to an adjacency list;
- adj2edgeL.m, (edgeL2adj.m) convert an adjacency matrix to an edge list;
- adj2inc.m, (inc2adj.m) convert adjacency matrix to incidence matrix and vice-versa;
- adj2str.m, (str2adj.m) convert adjacency matrix to a string graph representation;
- adjL2edgeL.m, (edgeL2adjL.m) convert adjacency list to edge list;
- inc2edgeL.m convert incidence matrix to an edge list;
- adj2simple.m Remove selfloops and multiedges from an adjacency matrix;
- edgeL2simple.m remove selfloops and multiedges from an edge list;
- add\_edge\_weights.m adding edges that occur multiple times in an edgelist;

#### **Centrality measures & Distributions (15)**

- degrees.m total degree, in- and out-degree sequence of an arbitrary graph;
- rewire.m degree-preserving rewiring;
- rewire assort.m degree-preserving rewiring with increasing assortativity;
- <u>rewire\_disassort.m</u> degree-preserving rewiring with decreasing assortativity;

(for degree preserving rewiring, see also <u>code by Maslov</u>);

- ave\_neighbor\_deg.m the average degree of neighboring nodes for every vertex;
- closeness.m computes the closeness centrality for all vertices;
- node betweenness.m node betweenness, (number of shortest paths definition);
- <u>node\_betweenness.m</u> a faster node betweenness algorithm;
- edge betweenness.m edge betweenness, (number of shortest paths definition);
- eigencentrality.m eigenvector corresponding to the largest eigenvalue;

- clust coeff.m two clustering coefficients: based on loops and local clustering;
- weighted\_clust\_coeff.m weighted clustering coefficient;
- <u>pearson.m</u> pearson degree correlation;
- rich club metric.m
- s metric.m the sum of products of nodal degrees across all edges;

### Spectral properties (6)

- <u>laplacian matrix.m</u> the Laplacian of the graph: degree matrix minus the adjacency;
- graph\_spectrum.m the sorted eigenvalues of the Laplacian;
- <u>algebraic\_connectivity.m</u> the second smallest eigenvalue of the Laplacian;
- fiedler vector.m the eigenvector corresponding to the algebraic connectivity;
- <u>eigencentrality.m</u> see Centrality measures & Distributions
- graph\_energy.m

#### Distances (12)

- simple\_dijkstra.m simple Dijkstra, does not remember the path;
- dijkstra.m Dijkstra which also returns the shortest paths;
- <u>shortest\_pathDP.m</u> shortest path algorithm using dynamic programming;
- kneighbors.m returns the indices of all neighbors k links away (for a given node);
- kmin neighbors.m indices of all neighbors minimum k links away (for a given node);
- diameter.m the longest shortest path in the graph;
- ave\_path\_length.m the average shortest path in the graph;
- smooth\_diameter.m a smoothed definition of diameter (effective diameter);
- <u>closeness.m</u> see Centrality measures & Distributions;
- vertex eccentricity.m the maximum distance to any other vertex;
- graph\_radius.m the minimum vertex eccentricity;
- distance distribution.m fraction of nodes at a given distance;

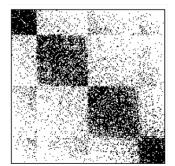
## Comparing graphs (1)

graph similarity.m - hubs and authorities based graph similarity matrix;

#### Modularity (6)

- <u>spectral\_partitioning.m</u> simple spectral partitioning;
- Newman-Girvan algorithm for community finding;
- <u>Newman eigenvector</u> modularity routine;
- <u>newman\_comm\_fast.m</u> faster community finding (Newman, 2003/4);

- modularity\_metric.m as defined in Newman/Girvan above;
- <u>louvain\_community\_finding.m</u>



**Example of Spectral Partitioning** 

#### Motifs (4)

- loops3.m count all loops of size 3 in the graph;
- <u>loops4.m</u> returns all loops of size 4;
- <u>num\_loops.m</u> number of independent loops;
- num\_star\_motifs.m number of star motifs of given size;

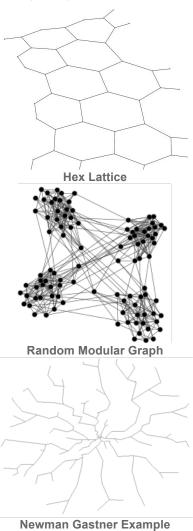
## **Building graphs (16)**

• random graph.m - generate a random graph adjacency matrix using various models;

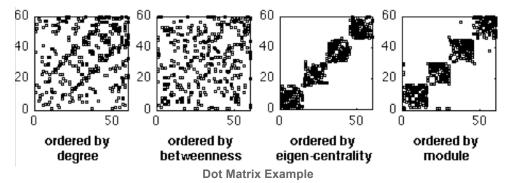
random\_graph(n) - Erdos-Renyi graph with **n** nodes and probability of attachment 0.5; random\_graph(n,p) - Erdos-Renyi graph with **n** nodes and probability of attachment **p**; random\_graph(n,[],E) - random graph with E number of edges; random\_graph(n,[],[],[],distribution) - nodal degrees are drawn from a particular distribution (uniform, normal, binomial, exponential); random graph(n,[],[],sequence',deg seq) - the degrees match a given sequence;

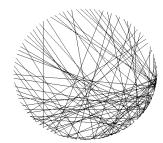
- <u>random\_directed\_graph.m</u> simple random directed graph routine;
- <u>random\_modular\_graph.m</u> random graph with modules;
- graph from degree sequence.m build deterministic graph given the nodal degrees;
- canonical nets.m simple graphs such as trees, lattices and hierarchical trees;
- kregular.m simple routine for building k-regular graphs;
- PriceModel.m Price's citations growth model;
- preferential attachment.m preferential attachment graph;
- exponential\_growth\_model.m;
- <u>master\_equation.m</u> a generalization of the degree-based random growth models;
- newmangastner.m Newman-Gastner spatial distribution growth model;
- <u>fabrikant\_model.m</u> another spatial distribution growth model;
- DoddsWattsSabel.m building randomized hierarchies, the Dodds-Watts-Sabel model;

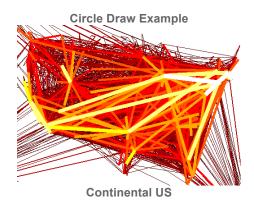
- <u>s-max graph</u> graph with the same degree sequence but maximum s-metric;
- forest fire model;
- <u>nested hierarchies model</u> by Sales-Pardo et al, 2007;



- pdf\_cdf\_rank.m pdf, cdf and rank distributions of a given sequence;
- adj2pajek.m, (pajek2adj.m) adjacency matrix to Pajek format: output .net file;
- edgeL2pajek.m, (pajek2edgeL.m) edge list to Pajek format: output .net file;
- <u>pajek2xyz.m</u> Extract node x,y,z coordinates from a Pajek .net file;
- el2geom.m plot an edge list geographically with color-coding of edge weights;
- <u>edgeL2cyto.m</u> convert edge list to Cytoscape text input;
- adj2dl.m adjacency to UCINET format; by D. Whitney;
- edgelist2dl.m edge list to UCINET format; by D. Whitney;
- draw\_circ\_graph.m draw a graph with all nodes in a circle, ordered by degree;
- <u>dot\_matrix\_plot.m</u> sparsity plots with nodes sorted by degree, betw-ness, modularity;
- <u>radial\_plot.m</u> best for trees and sparse graphs;









**Radial Plot Example** 

## Auxiliary (8)

- symmetrize.m symmetrize a matrix;
- symmetrize edgeL.m symmetrize an edge list (make it undirected);
- num\_conn\_triples.m the number of connected triples of nodes;
- <u>purge.m</u> remove elements from a list, while preserving the order;
- min\_span\_tree.m construct a minimum spanning tree given the adjacency matrix;
- BFS.m breadth-first tree;
- · sort nodes by the sum of neighbor degrees;
- sort nodes by the maximum neighbor degree;

## Links

- grTheory: Matlab Central
- MatlabBGL
- Bioinformatics toolbox: graph theory functions
- Matgraph by Ed Scheinerman
- Brain connectivity toolbox
- graphviz4matlab

## **Publications**

1. Bounova, G., de Weck, O.L. <u>"Overview of metrics and their correlation patterns for multiple-metric topology</u> analysis on heterogeneous graph ensembles", Phys. Rev. E 85, 016117 (2012).

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