Octave routines for network analysis

GB October 28, 2014



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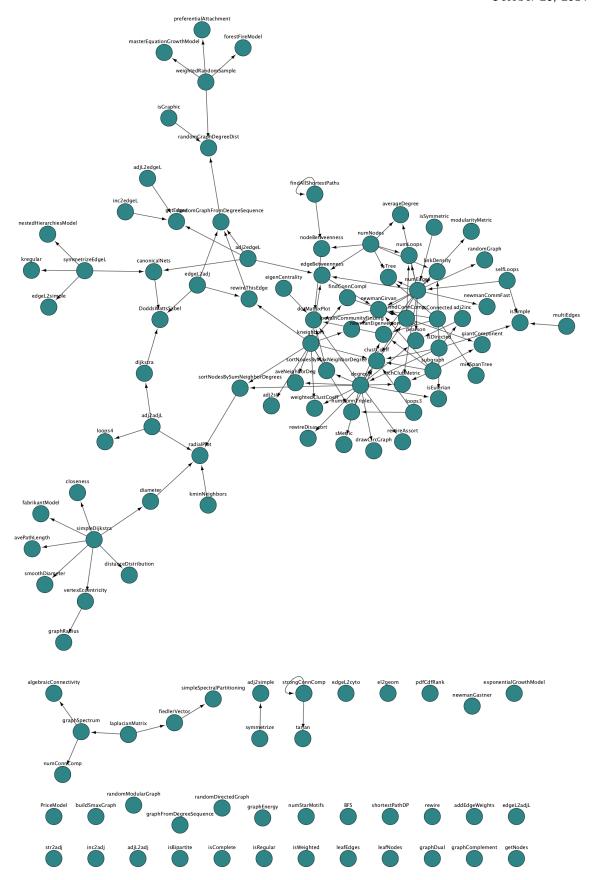


Figure 1: Graph of functional interdependencies in this toolbox. An edge points from routine A to routine B if routine A is called within routine B. For example, isConnected.m is called within minSpanTree.m

0 About this toolbox

(This is a copy of the README file.)

octave-networks-toolbox: A set of graph/networks analysis functions in Octave, 2012-2014

Quick description

This is a repository of functions relevant to network/graph analysis, organized by functionality. 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.

History

The original (2006-2011) version of these routines was written in Matlab, and is still hosted by strategic.mit.edu (http://strategic.mit.edu/downloads.php?page=matlab_networks). The octave-networks-toolbox inherits the original BSD open source license and copyright, provided at the end of this file. Many of the routines might still be compatible with Matlab. For Octave/Matlab differences, see http://en.wikibooks.org/wiki/MATLAB_Programming/Differences_between_Octave_and_MATLAB.

Installation

The code currently runs on GNU Octave Version 3.4.0 with Gnuplot 4.2.5. No specific library installation necessary. Interdependencies between functions are well-documented in the function headers. The routines can be called directly from the Octave prompt, either in the same directory or from anywhere if the toolbox folder is added to the path. For example:

```
% running numNodes.m
octave:1> numNodes([0 1 1; 1 0 1; 1 1 0])
ans = 3
```

Matlab compatibility

With newer versions of Matlab, the Octave branch may not always be Matlab-compatible, for example due to syntax changes. Consider exploring forks that focus on Matlab compatibility. There is currently no plan for the Octave original version to be synchronized with Matlab.

Authorship

This code was originally written and posted by Gergana Bounova. It is undergoing continuous expansion and development. Thank you for the many comments and bug reports so far! Contributions via email are usually given tribute to in the function header. Collaborators are very welcome. Contact via github/email for comments, questions, suggestions, corrections or simply fork.

Organization

The functions are organized in 11 categories: basic routines, diagnostic routines, conversion routines, centralities, distances, simple motif routines, linear algebra functions, modularity routines, graph construction models, visualization and auxiliary. These categories reflect roles/functionality and topics in the literature, but they are arbitrary, and mostly used for documentation purposes.

Documentation

Documentation is available in this Functions Manual. The manual contains general background information, function headers, code examples, and references. For some functions, additional background, definitions or derivations are included.

Citation

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1 Representing graphs in octave/matlab

1.1 Graph representations

Most succintly, a graph is a set of edges. For example, $\{(n_1, n_2), (n_2, n_3), (n_4, n_4)\}$ is a representation which stands for a 4-node graph with 3 edges, one of which is a self-loop. It is also easy to see that this graph is directed and disconnected, and it has a 3-node weakly connected component (see 2.1), namely $\{n_1, n_2, n_3\}$.

For larger graphs, text or visual representation does not suffice to answer even simple questions about the graph. Below are the definitions of some common graph representations, that could be used for computation. These should help with understanding and using the conversion routines in Section 1.2.

For the following discussion, assume that **n** is the number of nodes in a given graph, and **m** is the number of edges.

An **adjacency matrix** is a $n \times n$ matrix A, such that A(i,j) = 1 if i is connected to j and A(i,j) = 0, otherwise. The 1s in the matrix stand for the edges. If the graph is undirected, then the matrix is symmetric, because A(i,j) = A(j,i) for any i and any j. While usually this is a 0-1 matrix, sometimes edge weights can be indicated by using other numbers, so most generally the adjacency matrix has zeros and positive entries.

An **edge list** is a matrix representation of the set of edges. For the toy example $\{(n_1, n_2), (n_2, n_3), (n_4, n_4)\}$, the edge list representation would be $[n_1 \ n_2; n_2 \ n_3; n_4 \ n_4]$. Edge lists can have weights too, for example:

$$edge \ list = \left[\begin{array}{ccc} n_1 & n_2 & 0.5 \\ n_2 & n_3 & 1 \\ n_4 & n_4 & 2 \end{array} \right].$$

The **adjacency list** is the sparsest graph representation. For every node, only its list of neighbors is recorded. In Octave, one can use the cell structure to represent the adjacency list. In other languages this is known as a dictionary. The adjacency list representation of the 4-node example above is: $adjList\{n_1\} = [n_2]$, $adjList\{n_2\} = [n_3]$, $adjList\{n_4\} = [n_4]$.

The **incidence matrix** I is a table of nodes (n) versus edges (m). In other words, the rows are node indices and columns correspond to edges. So if edge e connects nodes i and j, then I(i,e) = 1 and I(j,e) = 1. For directed graphs I(i,e) = -1 and I(j,e) = 1, if i is the source node and j the target. For the above example:

$$I = egin{array}{c|cccc} & e_1 & e_2 & e_3 \ \hline n_1 & -1 & 0 & 0 \ n_2 & 1 & -1 & 0 \ n_3 & 0 & 1 & 0 \ n_4 & 0 & 0 & 1 \ \end{array}.$$

There can be other representations depending on purpose, understanding, or algorithm implementation. Suppose the graph information has to be stored as text. Here is an example **string representation** that could be easily read and stored in a text file. It is essentially the adjacency list, with some string nomenclature. Nodes are indexed from 1 to n, and every node has a list of neighbors (could be empty). Nodes and their lists are separated by commas (,), new neighbors by dots (.). Of course, this is arbitrary, but it is quite clear. The toy example representation is:

Four commas mean four nodes. Node 1 has one neighbor, namely node 2. Node 2 connects to node 3, node 3 has no neighbors (adjacent commas), and node 4 connects to itself. As an additional example, here is the representation of an undirected three-node cycle: "2.3,.1.3,.1.2,".

So there are many ways to write down and store a graph structure. Figure 2 shows one more example of all representations described above.

1.2 Routines

The functions in this section are conversion routines from one graph representation to another.

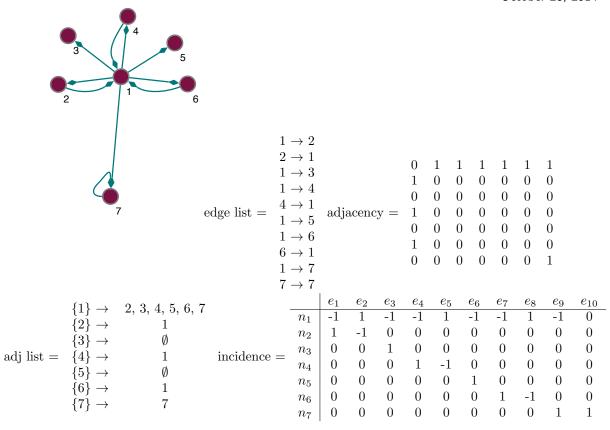


Figure 2: Most common graph representations: edge list, adjacency matrix, adjacency list and incidence matrix. Example of 7-node directed graph, with one self-loop. The string representation of this graph is ".2.3.4.5.6.7,.1,..1,..1,..7,".

1.2.1 adj2adjL.m

Convert an adjacency matrix to an adjacency list. This is the inverse function of adjL2adj.m (1.2.2).

```
% Convert an adjacency graph representation to an adjacency list. % Note 1: Valid for a general (directed, not simple) graph. % Note 2: Edge weights (if any) get lost in the conversion. % % INPUT: an adjacency matrix, nxn % OUTPUT: cell structure for adjacency list: x\{i_-1\}=[j_-1,j_-2,\ldots] % GB: last updated, September 24 2012
```

```
% undirected binary tree with 3 nodes
octave:1> adj2adjL([0 1 1; 1 0 0; 1 0 0])
ans =
{
   [1,1] =
        2      3
   [2,1] = 1
   [3,1] = 1
}
```

1.2.2 adjL2adj.m

Convert an adjacency list to an adjacency matrix. This is the inverse function of adj2adjL.m (1.2.1).

```
% Convert an adjacency list to an adjacency matrix.
% INPUTS: adjacency list: length n, where L\{i_1\}=[j_1,j_2,...]
% OUTPUTS: adjacency matrix nxn
% Note: Assume that if node i has no neighbours, then L\{i\}=[];
% GB: last updated, Sep 25 2012
octave:48> aL = { [2,3],[1,3],[1,2]};
octave:49> adjL2adj(aL)
ans =
   0
      1
           1
       0
           1
   1
      1
           0
```

1.2.3 adj2edgeL.m

Convert an adjacency matrix to an edge list. This is the inverse routine of edgeL2adj.m (1.2.4).

```
% Convert adjacency matrix (nxn) to edge list (mx3)
% INPUTS: adjacency matrix: nxn
% OUTPUTS: edge list: mx3
%
% GB: last updated, Sep 24, 2012
```

Example:

1.2.4 edgeL2adj.m

Convert edge list to adjacency matrix. This is the inverse routine of adj2edgeL.m (1.2.3).

```
% Convert edge list to adjacency matrix.
%
% INPUTS: edge list: mx3, m - number of edges
% OUTPUTS: adjacency matrix nxn, n - number of nodes
%
% Note: information about nodes is lost: indices only (i1,...in) remain
% GB: last updated, Sep 25, 2012
```

```
# a single directed edge
octave:1> edgeL2adj([1 2 1])
ans =

0    1
0    0
```

1.2.5 adj2inc.m

Convert an adjacency matrix to an incidence matrix. This is the inverse function of inc2adj.m 1.2.6.

```
% Convert adjacency matrix to an incidence matrix
% Note: Valid for directed/undirected, simple/not simple graphs
%
% INPUTs: adjacency matrix, nxn
% OUTPUTs: incidence matrix: n x m (number of edges)
%
% Other routines used: isDirected.m
% GB: last updated, Sep 25 2012
```

Example:

```
octave:4> adj2inc([0 1 1; 1 0 0; 1 0 0])
ans =
    1    1
    1    0
    0    1
```

1.2.6 inc2adj.m

Convert an incidence matrix to an adjacency matrix. This is the inverse function of adj2inc.m (1.2.5).

```
% Convert an incidence matrix representation to an
% adjacency matrix representation for an arbitrary graph.
%
% INPUTs: incidence matrix, nxm (num nodes x num edges)
% OUTPUTs: adjacency matrix, nxn
%
% GB: last updated, Sep 25, 2012
```

Example:

```
octave:10> inc = [1 0 1; 1 1 0; 0 1 1];
octave:11> inc2adj(inc)
ans =
0 1 1
1 0 1
1 1 0
```

1.2.7 adj2str.m

Convert an adjacency matrix to a string (text) graph representation. This is the inverse function of str2adj.m (1.2.8).

```
% Other routines used: kneighbors.m % GB: last updated, Sep 25 2012
```

```
% undirected binary tree

adj2str([0 1 1; 1 0 0; 1 0 0])

ans = .2.3, .1, .1,
```

$1.2.8 ext{ str2adj.m}$

This is the reverse routine of adj2str.m (1.2.7). Convert a string (text) graph representation to an adjacency matrix.

Example:

1.2.9 adjL2edgeL.m

Convert an adjacency list to an edge list. This is the inverse routine of edgeL2adjL.m (1.2.10).

```
% Convert adjacency list to an edge list.
%
% INPUTS: adjacency list
% OUTPUTS: edge list, mx3 (m - number of edges)
%
% GB: last updated, Sep 25 2012
```

1.2.10 edgeL2adjL.m

Convert an edge list to an adjacency list. This is the inverse routine of adjL2edgeL.m (1.2.9).

```
% Convert an edge list to an adjacency list.
%
% INPUTS: edge list, mx3, m - number of edges
% OUTPUTS: adjacency list
%
% Note: Information about edge weights (if any) is lost.
% GB: last updated, September 25, 2012
```

Example:

```
octave:1> edgeL2adjL([1 2 1; 1 3 1]) ans = {
    [1,1] = 
        2      3
        [2,1] = [](0x0)      [3,1] = [](0x0)
}
```

${\bf 1.2.11 \quad inc2edgeL.m}$

Convert an incidence matrix to an edge list.

```
% Convert an incidence matrix to an edge list.
%
% Inputs: inc - incidence matrix nxm (number of nodes x number of edges)
% Outputs: edge list - mx3, m x (node 1, node 2, edge weight)
%
% Example: [-1; 1] <=> [1,2,1], one directed (1->2) edge
% GB: last updated, Sep 25 2012
```

Example:

```
octave:2> inc = [1 0; 1 1; 0 1];
octave:3> inc2edgeL(inc)
ans =

1 2 1
2 3 1
2 1 1
3 2 1
```

${\bf 1.2.12}\quad adj2simple.m$

Remove self-loops and multi-edges from an adjacency matrix. Also symmetrizes the matrix and removes edge weights to produce the matrix of the corresponding simple graph.

```
% Convert an adjacency matrix of a general graph to the adjacency matrix of a simple graph (symmetric, no loops, no double edges, no weights)
% INPUTS: adjacency matrix, nxn
% OUTPUTs: adjacency matrix (nxn) of the corresponding simple graph
```

```
% Other routines used: symmetrize.m
% GB: last updated, Sep 6 2014
```

```
octave:1> adj2simple([1 2 1; 2 0 1; 1 1 0])
ans =

0 1 1
1 0 1
1 1 0
```

1.2.13 edgeL2simple.m

Remove self-loops and multi-edges from an edge list. Also symmetrizes the edge list and removes edge weights to produce the edge list of the corresponding simple graph.

```
% Convert an edge list of a general graph to the edge list of a
% simple graph (no loops, no double edges, no edge weights, symmetric)
%
% INPUTS: edge list (mx3), m - number of edges
% OUTPUTs: edge list of the corresponding simple graph
%
% Note: Assumes all node pairs [n1,n2,x] occur once; if else see addEdgeWeights.m
% Other routines used: symmetrizeEdgeL.m
% GB: last updated, Sep 25, 2012
```

Example:

```
octave:2> edgeL2simple([1 1 1; 1 2 1; 1 3 2])
ans =

1 2 1
1 3 1
2 1 1
3 1 1
```

1.2.14 symmetrize.m

Symmetrize a matrix. In this context, this means convert a directed graph representation to its equivalent undirected representation.

```
% Symmetrize a non-symmetric matrix,
% i.e. returns the undirected version of a directed graph.
% Note: Where mat(i,j)~=mat(j,i), the larger (nonzero) value is chosen
%
% INPUTS: a matrix - nxn
% OUTPUT: corresponding symmetric matrix - nxn
%
% GB: last updated: October 3, 2012
function adj_sym = symmetrize(adj)
adj_sym = max(adj,transpose(adj));
```

1.2.15 symmetrizeEdgeL.m

This function is similar to 1.2.14. For a general edge list, perhaps representing a directed graph, it checks whether the reverse edges of all edges are present. If not, they are added so the resulting graph is undirected.

```
% Making an edge list (representation of a graph) symmetric,
% i.e. if [n1,n2] is in the edge list, so is [n2,n1].
%
% INPUTs: edge list, mx3
% OUTPUTs: symmetrized edge list, mx3
%
% GB: last updated, October 3, 2012
```

Alternative to symmetrizeEdgeL.m using edgeL2adj.m, symmetrize.m and adj2edgeL.m.

```
def symmetrizeEdgeL(el):
    adj=edgeL2adj(el);
    adj=symmetrize(adj);
    el=adj2edgeL(adj);
    return el
```

Example:

1.2.16 addEdgeWeights.m

Adding edges that occur multiple times in an edge list; summing weights.

```
% Add multiple edges in an edge list
%
INPUTS: original (non-compact) edge list
% OUTPUTS: final compact edge list (no row repetitions)
%
% Example: [1 2 2; 2 2 1; 4 5 1] -> [1 2 3; 4 5 1]
% GB: last updated, Sep 25 2012
```

Example:

2 Basic network routines

2.1 Basic network theory

A graph is a set of nodes, and an associated set of links between them.

Networks are instantiations of graphs. They often represent real world systems that can be modeled as a set of connected entities.

Network theory is a modern branch of **graph theory**, concerned with statistics on practical instances of mathematical graphs. Graph theory and network theory references are abundant. Social science is probably the most

recent instigator of the trend to see the world as a network. In 1967, Milgram conducted his famous small world experiment [1], and found that Omahans are on average six steps away by acquaintance from Bostonians. Other prominent first sources are Price's work on the graph of scientific citations in 1965 [2] and in 1998, Watts and Strogatz's paper on dynamics of small-world networks [3].

Nowadays, there is no shortage of books and reviews on networks. Below is a non-exhaustive list of good reads [4] [5] [6] [7].

- S. Wasserman and K. Faust, Social network analysis, Cambridge University Press, 1994
- Duncan J. Watts, Six degrees: The science of a Connected Age, W. W. Norton, 2004
- M. E. J. Newman, The structure and function of complex networks, SIAM Review 45, 167-256 (2003)
- Alderson D., Catching the Network Science Bug: ..., Operations Research, Vol. 56, No. 5, Sep-Oct 2008, pp. 1047-1065

Here are some basic notions about graphs that are useful to understand the routines in Section 2.2.

Figure 1 illustrates a general **directed** graph. The nodes are functions from this toolbox. An edge points from function A to function B if function A is called within function B. For example, strongConnComp is used within tarjan. Notice, also that strongConnComp points to itself, i.e. strongConnComp contains a recursion. Stand-alone functions, that use no other function, are **single nodes** in the graph, such as leafNodes, getEdges and graphDual.

A directed graph is a graph in which the links have a direction. In the functions graph one function can call another, but the call is usually not reciprocated.

A **single node** is a node without any connections to other nodes. *graphDual* is an example of a single node in Figure 1.

A **self-loop** is an edge which starts and ends at the same node. $(strongConnComp \rightarrow strongConnComp)$ is an example of a self-loop.

Multiedges are two or more edges which have the same origin and destination pair of nodes. This can be useful in some graph representations. In the functions graph this is equivalent to some function being called twice inside another function.

Basic graph statistics are the **number of nodes** (n) and the **number of edges** (m). The functions graph has 118 nodes and 125 edges.

The **link density** is derived directly from the number of nodes and number of edges: it is the number of edges, divided by the maximum possible number of edges.

$$density = \frac{m}{n(n-1)/2} \tag{1}$$

For the functions graph, the link density is about 0.0181. Note that equation 1 is valid for undirected graphs only.

The average nodal degree is the average number of links per node. This is calculated as 2m/n (every edge is counted twice towards the total sum of degrees).

$$average \ degree = \frac{2m}{n} \tag{2}$$

The functions graph has 2.12 links per function on average.

A graph S is a **subgraph** of graph G, if the set of nodes (and edges) of S is subset of the set of nodes (and edges) of graph G.

A disconnected graph is a graph in which there are two nodes between which there exists no path of edges. In the functions graph there is no path between *rewire* and *subgraph*. So the functions graph is disconnected. Disconnected graphs consist of multiple connected components. The largest connected component (in number of nodes) is usually called the **giant component**. The giant component in Figure 1 has 80 functions. There are also one connected components of 6 functions, two 2-node components and 28 isolated nodes (functions that do not call or are not called within other functions).

In the context of **directed graphs**, the notion of strong and weak connectivity is important. A **strongly connected graph** is a graph in which there is a path from every node to every other node, where paths respect link directionality. In Figure 1, for example, there is a path from strongConnComp to tarjan, but no path in reverse. Therefore, the component (strongConnComp,tarjan) is not strongly connected. If, however, link directionality is disregarded, this subgraph is certainly connected. A **weakly connected graph** or subgraph is a graph which is connected if considered as undirected, but not connected if link directionality is taken into account. So the two-node subgraph (strongConnComp,tarjan) is definitely weakly connected.

2.2 Routines

2.2.1 getNodes.m

Returns the **list of nodes** for varying graph representations.

```
% Returns the list of nodes for varying graph representation types
% Inputs: graph structure (matrix or cell or struct) and type of structure (string)
          'type' can be: 'adjacency', 'edgelist', 'adjlist', 'incidence'
% Note 1: only the edge list allows/returns non-consecutive node indexing
\% Example representations of a directed 3-loop: 1->2->3->1
%
           'adj' - [0 1 0; 0 0 1; 1 0 0]
           'adjlist' - {1: [2], 2: [3], 3: [1]}
%
%
           'edgelist' - [1 2; 2 3; 3 1] or [1 2 1; 2 3 1; 3 1 1] (with edge weights)
%
           'inc' - [-1 \ 0 \ 1]
%
                       1 - 1 0
%
                       0 \quad 1 \quad -1
%
% GB: last updated, Jul 12 2014
```

Examples:

2.2.2 getEdges.m

Returns the **list of edges** for varying graph representations.

```
% using adjacency matrix representation
octave:46> getEdges([0 1 1; 1 0 1; 1 1 0], 'adjacency')
ans =
   1
        2
            1
        3
   1
            1
   2
       1
            1
   2
        3
            1
   3
        1
            1
   3
% using adjacency list representation
octave:47> adjL = {[2,3],[1,3],[1,2,4],[3,5,6],[4,6],[4,5]};
octave:48> getEdges(adjL, 'adjlist ')
ans =
   1
            1
   1
        3
            1
   2
        1
            1
   2
        3
            1
   3
       1
            1
   3
       2
            1
   3
        4
            1
        3
   4
            1
   4
        5
            1
   4
       6
            1
   5
        4
            1
   5
        6
            1
   6
        4
            1
   6
        5
```

Note that the column of 1s in the output shows the edge weight for every edge. If the graph is unweighted (as in this case), this column is unnecessary and is easy to remove. In fact, from the graph representations discussed in Section 1.1 only the *edge list* can carry edge weight information.

2.2.3 numNodes.m

Number of nodes/vertices in the network.

```
% Returns the number of nodes, given an adjacency list, or adjacency matrix % INPUTs: adjacency list: \{i:j_-1,j_-2...\} or adjacency matrix, ex: [0\ 1;\ 1\ 0] % OUTPUTs: number of nodes, integer % % GB: last update Sep 19, 2012 function n = numNodes(adjL)
```

```
n = length(adjL);
```

```
octave:2> N = randi (100);

octave:3> adj = randomGraph(N);

octave:3> % test whether the random graph does indeed have N nodes

octave:4> assert (numNodes(adj),N)

octave:4> % a graph (adjacency list) with 6 nodes

octave:4> adjL = {[2,3],[1,3],[1,2,4],[3,5,6],[4,6],[4,5]};

octave:5> numNodes(adjL)

ans = 6
```

2.2.4 numEdges.m

Number of edges/links in the network.

```
% Returns the total number of edges given the adjacency matrix
% INPUTs: adjacency matrix, nxn
% OUTPUTs: m - total number of edges/links
%
% Note: Valid for both directed and undirected, simple or general graph
% Other routines used: selfLoops.m, isSymmetric.m
% GB, last updated Sep 19, 2012
```

Examples:

2.2.5 linkDensity.m

The **density of links** of the graph. For an undirected graph the density is defined as $density = \frac{m}{n(n-1)/2}$, where n is the number of nodes and m is the number of edges. The directed graph version is the same but without the factor of 2, because the possible number of edges is twice as many.

See also 2.2.8.

```
% Computes the link density of a graph, defined as the number
% of edges divided by number_of_nodes(number_of_nodes-1)/2
% where the latter is the maximum possible number of edges.
%
% Inputs: adjacency matrix, nxn
% Outputs: link density, a float between 0 and 1
%
% Note 1: The graph has to be non-trivial (more than 1 node).
% Note 2: Routine works for both directed and undirected graphs.
%
% Other routines used: numNodes.m, numEdges.m, isDirected.m
% GB: last update Sep 19, 2012
```

2.2.6 selfLoops.m

Number of **self-loops**, i.e. number of edges that start and end at the same node.

```
% Counts the number of self-loops in the graph
%
% INPUT: adjacency matrix, nxn
% OUTPUT: integer, number of self-loops
%
% Note: in the adjacency matrix representation
% loops appear as non-zeros on the diagonal
% GB: last updated, Sep 20 2012
```

Examples:

```
octave:2> selfLoops([0 1; 0 0])
ans = 0

octave:3> % three self-loops
octave:3> adj = [1 0 0; 0 1 0; 0 0 1];
octave:4> selfLoops(adj)
ans = 3
```

2.2.7 multiEdges.m

An edge counts towards the **multi-edge** total if it shares origin and destination nodes with another edge.

```
% Counts the number of multiple edges in the graph.
%

NPUT: adjacency matrix, nxn
OUTPUT: integer, number of multiple edges
%

Examples: multiEdges([0 2; 2 0])=1, and multiEdges([0 0 1; 2 0 0; 0 1 0])=1
%

Note 1: The definition of number of multi-arcs/edges (node pairs
that have multiple edges across them) here is: mA = length(find(adj>1))
% (normalized by 2 depending on whether the graph is directed).
% Note 2: This creates a natural difference in counting for
% undirected and directed graphs.
%

Other routines used: isSymmetric.m
% GB: last updated, Sep 26 2014
```

```
octave:22> one_double_edge = [0 2; 0 0];
octave:23> multiEdges(one_double_edge)
ans = 1
octave:24> double_edge = [0 2; 2 0];
octave:25> multiEdges(double_edge)
ans = 1
octave:26> adj = [1 1 0; 1 0 0; 0 0 0];
octave:27> multiEdges(adj)
ans = 0
```

2.2.8 averageDegree.m

The average degree (number of links per node) across all nodes. Defined as: $\frac{2m}{n}$, where n is the number of nodes and m is the number of edges. Also, note that the link density (Section 2.2.5) is related to the average degree: $linkDensity = \frac{averageDegree}{n-1}$.

Examples:

```
octave:7> % undirected 3-node cycle
octave:7> adj = [0 1 1; 1 0 1; 1 1 0];
octave:8> averageDegree(adj)
ans = 2
octave:9> % undirected 3-node binary tree
octave:9> adj = [0 1 1; 1 0 0; 1 0 0];
octave:10> averageDegree(adj)
ans = 1.3333
```

2.2.9 numConnComp.m

Calculating the number of connected components in the graph by using the eigenvalues of the Laplacian.

```
% Calculate the number of connected components using the eigenvalues
% of the Laplacian - counting the number of zeros
%
% INPUTS: adjacency matrix, nxn
% OUTPUTs: positive integer - number of connected components
%
% Other routines used: graphSpectrum.m
% GB: last updated: September 22, 2012
```

```
octave:15> numConnComp(adj)
ans = 2
```

2.2.10 findConnComp.m

findConnCompI.m: Find the **connected component** to which a given node i belongs to. This function is called within findConnComp.m.

Example:

 $\mathbf{findConnComp.m} \text{.} \ \mathbf{Find\ the\ connected\ components\ in\ an\ undirected\ graph}. \ \ \mathbf{This\ function\ uses}\ findConnCompI.m.$

```
% Algorithm for finding connected components in a graph
% Note: Valid for undirected graphs only
%
% INPUTS: adj - adjacency matrix, nxn
% OUTPUTS: a list of the components comp{i}=[j1,j2,...jk]
%
% Other routines used: findConnCompI.m, degrees.m
% GB: last updated, September 22, 2012
```

${\bf 2.2.11 \quad giant Component.m}$

The **largest connected component** in a graph. Return the set of nodes in the largest component, as well as its adjacency matrix.

```
% Extract the giant component of a graph;
% The giant component is the largest connected component.
%
% INPUTS: adjacency matrix, nxn
% OUTPUTS: giant component matrix and node
% indices of the giant component
%
% Other routines used: findConnComp.m, subgraph.m
% GB: last updated: September 22, 2012
```

Example:

```
octave:23> adj = [0 1 0; 1 0 0; 0 0 1];
octave:24> [GC, I] = giantComponent(adj);
octave:25> GC
GC =

0 1
1 0

octave:26> I
I =
```

2.2.12 tarjan.m [8][9]

tarjan.m: Return the strongly connected components in a directed graph.

```
% Find the stronly connected components in a directed graph
% Source: Tarjan, "Depth-first search and linear graph algorithms",
% SIAM Journal on Computing 1 (2): 146-160, 1972
% Wikipedia description:
% http://en.wikipedia.org/wiki/Tarjan's_strongly_connected_components_algorithm
%
% Input: graph, set of nodes and edges, in adjacency list format,
% example: L{1}=[2], L{2]=[1] is a single (1,2) edge
% Outputs: set of strongly connected components, in cell array format
%
% Other routines used: strongConnComp.m
% GB: last updated, Sep 22, 2012
```

strongConnComp.m: Support function for *tarjan.m*.

```
% Support function for tarjan.m
% "Performs a single depth-first search of the graph, finding all
% successors from the node vi, and reporting all strongly connected
% components of that subgraph."
% See: http://en.wikipedia.org/wiki/Tarjan's_strongly_connected_components_algorithm
%
INPUTs: start node, vi;
graph structure (list), L
% tarjan.m variables to update: S, ind, v, GSCC
% OUTPUTs: updated tarjan.m variables: S, ind, v, GSCC%
```

```
% Note: Contains recursion.
% GB: last updated, Sep 22 2012
```

```
octave:55> \% same as \{1:[2], 2:[3], 3:[1]\} (a directed 3-cycle)
octave:55> L = \{[2], [3], [1]\}
  [1,1] = 2
  [1,2] = 3
  [1,3] = 1
octave:56 > comp = tarjan(L);
octave:57> comp
comp =
  [1,1] =
     1
         2
             3
octave:58>
octave:58> comp = tarjan(\{[2,3],[],[]\}) % undirected binary tree
comp =
{
  [1,1] = 2
  [1,2] = 3
  [1,3] = 1
```

2.2.13 graphComplement.m

A graph with the same nodes, but "flipped" edges: where the original graph has an edge, the **complement graph** doesn't, and where the original graph doesn't have an edge, the complement graph does.

```
% Returns the complement of a graph
% The complement graph has the same nodes, but edges where
% the original graph doesn't and vice versa.
%
% INPUTs: adj - original graph adjacency matrix, nxn
% OUTPUTs: complement graph adjacency matrix, nxn
%
% Note 1: Assumes no multiple edges.
% Note 2: To create a complement graph without self-loops,
% use adjC=ones(size(adj))-adj-eye(length(adj)); instead.
% GB: last updated, October 4, 2014
```

Example:

```
octave:16> g = [0 1 1; 1 0 0; 1 0 0];
octave:17> gc = graphComplement(g)
gc =
1 0 0
0 1 1
0 1 1
```

2.2.14 graphDual.m

The **graph dual** is the inverted nodes-edges graph.

```
% Finds the dual of a graph; a dual is the inverted nodes—edges graph.
% This is also called the line graph, adjoint graph or the edges adjacency.
%
INPUTs: adjacency (neighbor) list representation of the graph (see adj2adjL.m)
% OUTPUTs: adj (neighbor) list of the corresponding dual graph and cell array of edges
%
Note: This routine only works for undirected, simple graphs.
% GB: last updated, Sep 23, 2012
```

```
octave:3> % cycle3 in adjacency matrix format is [0 1 1; 1 0 1; 1 1 0]
octave:3 > \text{cycle3} = \{ [2,3]; [1,3]; [1,2] \};
octave:4> graphDual(cycle3)
ans =
  [1,1] =
     2
         3
  [1,2] =
     1
         3
  [1,3] =
         2
     1
}
octave:5>
octave:5> % undirected 3-node binary tree
octave:6> L = \{ [2,3]; [1]; [1] \};
octave:7> graphDual(L)
ans =
  [1,1] = 2
  [1,2] = 1
```

2.2.15 subgraph.m

Return the adjacency matrix of a **subgraph**, given a subset of nodes.

```
% This function outputs the adjacency matrix of a subgraph given % the supergraph and the node set of the subgraph. % % INPUTs: adj - supergraph adjacency matrix (nxn), S - vector of subgraph node indices % OUTPUTs: adj_sub - adjacency matrix of the subgraph (length(S) x length(S)) % GB: last update, September 23 2012
```

```
octave:10> subgraph(bowtie,[3, 4, 5])
ans =

0    1    0
1    0    1
0    1    0
1    0    1
0    1    0
```

2.2.16 leafNodes.m

Leaf nodes are nodes connected to only one other node.

Examples:

```
octave:30> % only 1 is not a leaf node, because it has degree 2
octave:30> adj = [0 1 1; 1 0 0; 1 0 0];
octave:31> leafNodes(adj)
ans =
    2    3

octave:32> % a cycle graph has no leaf nodes
octave:32> adj = [0 1 1; 1 0 1; 1 1 0];
octave:33> leafNodes(adj)
ans = [](1x0)
```

2.2.17 leafEdges.m

Leaf edges are edges with only one adjacent edge.

```
% Returns the leaf edges of the graph: edges with one adjacent edge only.

% Note 1: For a directed graph, leaf edges are those that "flow into" a leaf node.

% Note 2: There could be other definitions of leaves, for example:

% farthest away from a given root node.

% Note 3: Edges that are self-loops are not considered leaf edges.

% Note 4: Single floating disconnected edges are not considered to be leaf edges.

% Input: adjacency matrix, nxn

% Output: set of leaf edges: a (num edges x 2) matrix where every row

% contains the leaf edge nodal indices

% GB: last updated, Sep 23, 2012
```

```
octave:2> % a binary tree with two leaf edges/nodes
octave:2> adj = [0 1 1; 1 0 0; 1 0 0];
octave:3> leafEdges(adj)
ans =

1  2
  1  3

octave:4> % a cycle has no leaf edges
octave:4> adj = [0 1 1; 1 0 1; 1 1 0];
octave:5> leafEdges(adj)
ans = [](0x0)
```

2.2.18 minSpanTree.m

Given a general graph, return an undirected minimum spanning tree of the graph, using Prim's algorithm.

```
% Prim's minimal spanning tree algorithm
% Prim's alg idea:
% start at any node, find closest neighbor and mark edges
% for all remaining nodes, find closest to previous cluster, mark edge
% continue until no nodes remain
%
% INPUTS: graph defined by adjacency matrix, nxn
% OUTPUTS: matrix specifying minimum spanning tree (subgraph), nxn
%
% Other routines used: isConnected.m
% GB: Oct 7, 2012
```

Example:

```
octave:10> minSpanTree(bowtie)
ans =
 0
       0
          0
            0
   0
     0
 1
   0
     0
          0
            0
 1
       1
 0
   0
     1
       0
         1
            1
 0
   0
     0
          0
            0
       1
 0
   0
```

2.2.19 DFS.m

Depth first search. Given a graph, a *start* and an *end* node, return all possible paths from the *start* to the *end*, subject to a path length upper bound. Currently, no edge weights are implemented in the path length, but that should be straighforward to modify.

```
% Find all paths from a start to an end node,
% bounded by a constant, using depth-first-search.
%
% Source: Idea from 6.002x, Spring 2014.
% Note: Uses recursion.
%
% INPUTs: graph structure (a nxn adjacency matrix)
% s - start node
% e - end node
% upperBound = 5, some constant bounding the
% length of the path; typically size(adj,1)-1
% allPaths = {}, path = []; are by default empty,
```

```
% serving the recursion
%
% OUTPUTs: a list {} of all shortest paths from "s" to "e",
% shorter than "upperBound"
%
% Note: Uses also kneighbors.m, DFS.m.
% GB: last updated Oct 27 2014
```

3 Diagnostic routines

These are functions that return boolean values when querying some property of the graph. They are often used by other algorithms in this toolbox.

3.1 Routines

3.1.1 isSimple.m

A simple graph is undirected, without self-loops, no multiple edges and no edge weights.

```
% Checks whether a graph is simple (undirected, no self-loops,
% no multiple edges, no weighted edges)
%
% INPUTs: adj - adjacency matrix
% OUTPUTs: S - a Boolean variable; true (1) or false (0)
%
% Other routines used: selfLoops.m, multiEdges.m, isDirected.m
% GB: last updated, September 23, 2012
```

Examples:

```
octave:9> % undirected binary tree
octave:9> isSimple([0 1 1; 1 0 0; 1 0 0])
ans = 1
octave:10> % a weighted graph example
octave:10> isSimple([0 2 1; 2 0 0; 1 0 0])
ans = 0
octave:11> % directed graph example
octave:11> isSimple([0 1 1; 0 0 0; 0 0 0])
ans = 0
```

3.1.2 isDirected.m

This routine checks whether a graph is **directed** or not.

```
% Checks whether the graph is directed, using the matrix transpose function.
% INPUTS: adjacency matrix, nxn
% OUTPUTS: boolean variable, 0 or 1
%
% Note: one-liner alternative: S=not(isSymmetric(adj));
% GB: last updated, Sep 23, 2012

function S=isDirected(adj)

S = true;
if adj=transpose(adj); S = false; end
```

```
octave:3> isDirected([0 1 1; 1 0 0; 1 0 0])

ans = 0

octave:4> isDirected([0 1 1; 0 0 0; 0 0 0])

ans = 1
```

3.1.3 isSymmetric.m

Checks whether a matrix is **symmetric**. Note then when a graph is represented with an adjacency matrix, isSymmetric.m is the same as isDirected.m (3.1.2).

```
% Checks whether a matrix is symmetric (has to be square). % % INPUTS: adjacency matrix, nxn % OUTPUTS: boolean variable, \{0,1\} % GB: last update, Sep 23, 2012
```

Examples:

```
octave:7> isSymmetric([0 1 1; 1 0 0; 1 0 0])

ans = 1

octave:8> isSymmetric([0 1 1; 0 0 0; 0 0 0])

ans = 0
```

3.1.4 isConnected.m

Checks whether a graph is connected. A graph is **connected** if there is a path, via edges, from any node to any other node. There are many ways to check this. Some alternatives to *isConnected.m* are listed below. The idea behind the main routine is from Matgraph's *isconnected* function.

Alternative 1 to isConnected.m

If the algebraic connectivity is >0 then the graph is connected.

```
a = algebraic\_connectivity(adj);

S = false; if a > 0; S = true; end
```

Alternative 2 to isConnected.m

Uses the fact that multiplying the adjacency matrix to itself k times give the number of ways to get from i to j in k steps. If the end of the multiplication in the sum of all matrices there are 0 entries then the graph is disconnected. Computationally intensive, but can be sped up by the fact that in practice the diameter is very short compared to n, so it will terminate in order of log(n) steps.

```
function S=isconnected(el):
    S=false;
    adj=edgeL2adj(el);
    n=numnodes(adj); % number of nodes
    adjn=zeros(n);
    adji=adj;
    for i=1:n
        adjn=adjn+adji;
        adji=adji*adj;
    if length(find(adjn==0))==0
        S=true;
        return
    end
end
```

Alternative 3 to isConnected.m

Find all connected components, if their number is 1, the graph is connected. Use findConnComp.m 2.2.10.

Examples:

3.1.5 isWeighted.m

Checks whether a graph has weighted links. Uses the edge list representation.

```
% Check whether a graph is weighted, i.e edges have weights.
%
% INPUTS: edge list, m x 3, m: number of edges, [node 1, node 2, edge weight]
% OUTPUTS: Boolean variable, 0 or 1
%
% GB: last updated, Sep 23, 2012
```

```
octave:16> eL =[1 2 1; 1 3 1; 2 3 1];
```

```
octave:17> isWeighted(eL)
ans = 0
octave:18> % undirected double (weighted) edge
octave:18> isWeighted([1 2 2; 2 1 2])
ans = 1
```

3.1.6 is Regular.m

Checks whether a graph is regular. In a **regular graph** every node has the same number of links.

```
% Checks whether a graph is regular, i.e.
% whether every node has the same degree.
%
INPUTS: adjacency matrix, nxn
% OUTPUTS: Boolean, 0 or 1
%
% Note: Defined for unweighted graphs only.
% GB: last updated, Sep 23, 2012
```

Examples:

```
octave:3> % undirected binary tree
octave:3> adj = [0 1 1; 1 0 0; 1 0 0];
octave:4> isRegular(adj)
ans = 0
octave:5> % undirected 3-node cycle
octave:5> adj = [0 1 1; 1 0 1; 1 1 0];
octave:6> isRegular(adj)
ans = 1
octave:7> % same as above, but edges are weighted
octave:7> adj = [0 2 2; 2 0 2; 2 2 0];
octave:8> isRegular(adj)
ans = 1
octave:9> % a 4-node cycle
octave:9> isRegular([0 1 0 1; 1 0 1 0; 0 1 0 1; 1 0 1 0])
ans = 1
```

3.1.7 isComplete.m

A complete graph is a graph in which all nodes are connected to all other nodes.

```
% Check whether a graph is complete, i.e.
% whether every node is linked to every other node.
%
% INPUTS: adjacency matrix, nxn
% OUTPUTS: Boolean variable, true/false
%
% Note: Only defined for unweighted graphs.
% GB: last updated, Sep 23, 2012
```

```
octave:3> isComplete([0 1 1; 1 0 0; 1 0 0])

ans = 0

octave:4> isComplete([0 1 1; 1 0 1; 1 1 0])

ans = 1
```

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