# **NetworkX Reference**

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## CONTENTS

1	Over	view 1
	1.1	Who uses NetworkX?
	1.2	Goals
	1.3	The Python programming language
	1.4	Free software
	1.5	History
2	Intuo	duction 3
4	2.1	NetworkX Basics
	2.2	Nodes and Edges
3	Grap	oh types 9
	3.1	Which graph class should I use?
	3.2	Basic graph types
4	Algor	rithms 123
•	4.1	Approximation
	4.2	Assortativity
	4.3	Bipartite
	4.4	Blockmodeling
	4.5	Boundary
	4.6	Centrality
	4.7	Chordal
	4.8	Clique
	4.9	Clustering
	4.10	Coloring
	4.11	Communities
	4.12	Components
	4.13	Connectivity
	4.14	Cores
	4.15	Cycles
	4.16	Directed Acyclic Graphs
	4.17	Distance Measures
	4.17	Distance-Regular Graphs
	4.19	Dominance
	4.19	Dominating Sets
	4.20	Eulerian
	4.21	Flows
	4.23	Graphical degree sequence
	4.24	Hierarchy
	4.25	Isolates

		Isomorphism	
	4.27	Link Analysis	
	4.28	Link Prediction	
	4.29	Matching	
	4.30	Maximal independent set	
	4.31	Minimum Spanning Tree	
	4.32	Operators	
	4.33	Rich Club	
	4.34	Shortest Paths	
	4.35	Simple Paths	
	4.36	Swap	
	4.37	Traversal	
	4.38	Tree	
	4.39	Vitality	346
_	E	41	349
5	Func		
	5.1	Graph	
	5.2	Nodes	
	5.3	Edges	
	5.4	Attributes	
	5.5	Freezing graph structure	556
6	Gran	oh generators 3	359
U	6.1	Atlas	
	6.2	Classic	
	6.3	Expanders	
	6.4	Small	
	6.5	Random Graphs	
	6.6	Degree Sequence	
	6.7	Random Clustered	
	6.8	Directed	
	6.9	Geometric	
	6.10	Hybrid	
	6.11	Line Graph	
	6.12	Ego Graph	
	6.13	Stochastic	
	6.14	Intersection	
	6.15	Social Networks	
	0.15	Social Networks	,,,
7	Linea	ar algebra	397
	7.1	Graph Matrix	397
	7.2	Laplacian Matrix	399
	7.3	Spectrum	101
	7.4	Algebraic Connectivity	102
	7.5	Attribute Matrices	105
8			111
	8.1	· · · · · ·	111
	8.2		112
	8.3		113
	8.4	17	114
	8.5		118
	8.6	Pandas	121
0	Dead	ing and writing graphs	125
ソ	Kead	ing and writing graphs	125

	1 Adjacency List	425
9	2 Multiline Adjacency List	429
9	3 Edge List	432
9	4 GEXF	439
	5 GML	441
	6 Pickle	445
	7 GraphML	447
	8 JSON	448
	9 LEDA	453
	10 YAML	454
	11 SparseGraph6	456
	12 Pajek	461
	13 GIS Shapefile	463
10	rawing	465
	0.1 Matplotlib	465
	0.2 Graphviz AGraph (dot)	474
	0.3 Graphviz with pydot	477
	0.4 Graph Layout	479
11	xceptions	483
12	tilities	485
	2.1 Helper Functions	485
	2.2 Data Structures and Algorithms	486
	2.3 Random Sequence Generators	486
	2.4 Decorators	489
	2.5 Cuthill-Mckee Ordering	490
	2.6 Context Managers	492
13	icense	495
	iting	
14	iting	497
	redits	497 499
15		
15 ( 16 (	redits	499 501
15 (16 (Bib)	redits clossary ography	499 501 503
15 (16 (Bib)	redits clossary ography on Module Index	499 501

**CHAPTER** 

ONE

## **OVERVIEW**

NetworkX is a Python language software package for the creation, manipulation, and study of the structure, dynamics, and function of complex networks.

With NetworkX you can load and store networks in standard and nonstandard data formats, generate many types of random and classic networks, analyze network structure, build network models, design new network algorithms, draw networks, and much more.

## 1.1 Who uses NetworkX?

The potential audience for NetworkX includes mathematicians, physicists, biologists, computer scientists, and social scientists. Good reviews of the state-of-the-art in the science of complex networks are presented in Albert and Barabási [BA02], Newman [Newman03], and Dorogovtsev and Mendes [DM03]. See also the classic texts [Bollobas01], [Diestel97] and [West01] for graph theoretic results and terminology. For basic graph algorithms, we recommend the texts of Sedgewick, e.g. [Sedgewick01] and [Sedgewick02] and the survey of Brandes and Erlebach [BE05].

## 1.2 Goals

NetworkX is intended to provide

- tools for the study of the structure and dynamics of social, biological, and infrastructure networks,
- · a standard programming interface and graph implementation that is suitable for many applications,
- a rapid development environment for collaborative, multidisciplinary projects,
- an interface to existing numerical algorithms and code written in C, C++, and FORTRAN,
- the ability to painlessly slurp in large nonstandard data sets.

## 1.3 The Python programming language

Python is a powerful programming language that allows simple and flexible representations of networks, and clear and concise expressions of network algorithms (and other algorithms too). Python has a vibrant and growing ecosystem of packages that NetworkX uses to provide more features such as numerical linear algebra and drawing. In addition Python is also an excellent "glue" language for putting together pieces of software from other languages which allows reuse of legacy code and engineering of high-performance algorithms [Langtangen04].

Equally important, Python is free, well-supported, and a joy to use.

In order to make the most out of NetworkX you will want to know how to write basic programs in Python. Among the many guides to Python, we recommend the documentation at http://www.python.org and the text by Alex Martelli [Martelli03].

## 1.4 Free software

NetworkX is free software; you can redistribute it and/or modify it under the terms of the *BSD License*. We welcome contributions from the community. Information on NetworkX development is found at the NetworkX Developer Zone at Github https://github.com/networkx/networkx

## 1.5 History

NetworkX was born in May 2002. The original version was designed and written by Aric Hagberg, Dan Schult, and Pieter Swart in 2002 and 2003. The first public release was in April 2005.

Many people have contributed to the success of NetworkX. Some of the contributors are listed in the *credits*.

## 1.5.1 What Next

- A Brief Tour
- Installing
- Reference
- Examples

**CHAPTER** 

**TWO** 

## INTRODUCTION

The structure of NetworkX can be seen by the organization of its source code. The package provides classes for graph objects, generators to create standard graphs, IO routines for reading in existing datasets, algorithms to analyse the resulting networks and some basic drawing tools.

Most of the NetworkX API is provided by functions which take a graph object as an argument. Methods of the graph object are limited to basic manipulation and reporting. This provides modularity of code and documentation. It also makes it easier for newcomers to learn about the package in stages. The source code for each module is meant to be easy to read and reading this Python code is actually a good way to learn more about network algorithms, but we have put a lot of effort into making the documentation sufficient and friendly. If you have suggestions or questions please contact us by joining the NetworkX Google group.

Classes are named using CamelCase (capital letters at the start of each word). functions, methods and variable names are lower\_case\_underscore (lowercase with an underscore representing a space between words).

## 2.1 NetworkX Basics

After starting Python, import the networkx module with (the recommended way)

```
>>> import networkx as nx
```

To save repetition, in the documentation we assume that NetworkX has been imported this way.

If importing networkx fails, it means that Python cannot find the installed module. Check your installation and your PYTHONPATH.

The following basic graph types are provided as Python classes:

**Graph** This class implements an undirected graph. It ignores multiple edges between two nodes. It does allow self-loop edges between a node and itself.

**DiGraph** Directed graphs, that is, graphs with directed edges. Operations common to directed graphs, (a subclass of Graph).

**MultiGraph** A flexible graph class that allows multiple undirected edges between pairs of nodes. The additional flexibility leads to some degradation in performance, though usually not significant.

MultiDiGraph A directed version of a MultiGraph.

Empty graph-like objects are created with

```
>>> G=nx.Graph()
>>> G=nx.DiGraph()
>>> G=nx.MultiGraph()
>>> G=nx.MultiDiGraph()
```

All graph classes allow any *hashable* object as a node. Hashable objects include strings, tuples, integers, and more. Arbitrary edge attributes such as weights and labels can be associated with an edge.

The graph internal data structures are based on an adjacency list representation and implemented using Python *dictionary* datastructures. The graph adjaceny structure is implemented as a Python dictionary of dictionaries; the outer dictionary is keyed by nodes to values that are themselves dictionaries keyed by neighboring node to the edge attributes associated with that edge. This "dict-of-dicts" structure allows fast addition, deletion, and lookup of nodes and neighbors in large graphs. The underlying datastructure is accessed directly by methods (the programming interface "API") in the class definitions. All functions, on the other hand, manipulate graph-like objects solely via those API methods and not by acting directly on the datastructure. This design allows for possible replacement of the 'dicts-of-dicts'-based datastructure with an alternative datastructure that implements the same methods.

## **2.1.1 Graphs**

The first choice to be made when using NetworkX is what type of graph object to use. A graph (network) is a collection of nodes together with a collection of edges that are pairs of nodes. Attributes are often associated with nodes and/or edges. NetworkX graph objects come in different flavors depending on two main properties of the network:

- Directed: Are the edges **directed**? Does the order of the edge pairs (u,v) matter? A directed graph is specified by the "Di" prefix in the class name, e.g. DiGraph(). We make this distinction because many classical graph properties are defined differently for directed graphs.
- Multi-edges: Are multiple edges allowed between each pair of nodes? As you might imagine, multiple edges requires a different data structure, though tricky users could design edge data objects to support this functionality. We provide a standard data structure and interface for this type of graph using the prefix "Multi", e.g. MultiGraph().

The basic graph classes are named: Graph, DiGraph, MultiGraph, and MultiDiGraph

## 2.2 Nodes and Edges

The next choice you have to make when specifying a graph is what kinds of nodes and edges to use.

If the topology of the network is all you care about then using integers or strings as the nodes makes sense and you need not worry about edge data. If you have a data structure already in place to describe nodes you can simply use that structure as your nodes provided it is *hashable*. If it is not hashable you can use a unique identifier to represent the node and assign the data as a *node attribute*.

Edges often have data associated with them. Arbitrary data can associated with edges as an *edge attribute*. If the data is numeric and the intent is to represent a *weighted* graph then use the 'weight' keyword for the attribute. Some of the graph algorithms, such as Dijkstra's shortest path algorithm, use this attribute name to get the weight for each edge.

Other attributes can be assigned to an edge by using keyword/value pairs when adding edges. You can use any keyword except 'weight' to name your attribute and can then easily query the edge data by that attribute keyword.

Once you've decided how to encode the nodes and edges, and whether you have an undirected/directed graph with or without multiedges you are ready to build your network.

## 2.2.1 Graph Creation

NetworkX graph objects can be created in one of three ways:

- Graph generators standard algorithms to create network topologies.
- Importing data from pre-existing (usually file) sources.

Adding edges and nodes explicitly.

Explicit addition and removal of nodes/edges is the easiest to describe. Each graph object supplies methods to manipulate the graph. For example,

```
>>> import networkx as nx
>>> G=nx.Graph()
>>> G.add_edge(1,2) # default edge data=1
>>> G.add_edge(2,3,weight=0.9) # specify edge data
```

Edge attributes can be anything:

```
>>> import math
>>> G.add_edge('y','x',function=math.cos)
>>> G.add_node(math.cos) # any hashable can be a node
```

You can add many edges at one time:

```
>>> elist=[('a','b',5.0),('b','c',3.0),('a','c',1.0),('c','d',7.3)]
>>> G.add_weighted_edges_from(elist)
```

See the /tutorial/index for more examples.

Some basic graph operations such as union and intersection are described in the Operators module documentation.

Graph generators such as binomial\_graph and powerlaw\_graph are provided in the Graph generators subpackage.

For importing network data from formats such as GML, GraphML, edge list text files see the *Reading and writing graphs* subpackage.

## 2.2.2 Graph Reporting

Class methods are used for the basic reporting functions neighbors, edges and degree. Reporting of lists is often needed only to iterate through that list so we supply iterator versions of many property reporting methods. For example edges() and nodes() have corresponding methods edges\_iter() and nodes\_iter(). Using these methods when you can will save memory and often time as well.

The basic graph relationship of an edge can be obtained in two basic ways. One can look for neighbors of a node or one can look for edges incident to a node. We jokingly refer to people who focus on nodes/neighbors as node-centric and people who focus on edges as edge-centric. The designers of NetworkX tend to be node-centric and view edges as a relationship between nodes. You can see this by our avoidance of notation like G[u,v] in favor of G[u][v]. Most data structures for sparse graphs are essentially adjacency lists and so fit this perspective. In the end, of course, it doesn't really matter which way you examine the graph. G.edges() removes duplicate representations of each edge while G[u,v] is slightly faster but doesn't remove duplicates.

Any properties that are more complicated than edges, neighbors and degree are provided by functions. For example nx.triangles(G,n) gives the number of triangles which include node n as a vertex. These functions are grouped in the code and documentation under the term *algorithms*.

## 2.2.3 Algorithms

A number of graph algorithms are provided with NetworkX. These include shortest path, and breadth first search (see *traversal*), clustering and isomorphism algorithms and others. There are many that we have not developed yet too. If you implement a graph algorithm that might be useful for others please let us know through the NetworkX Google group or the Github Developer Zone.

As an example here is code to use Dijkstra's algorithm to find the shortest weighted path:

```
>>> G=nx.Graph()
>>> e=[('a','b',0.3),('b','c',0.9),('a','c',0.5),('c','d',1.2)]
>>> G.add_weighted_edges_from(e)
>>> print(nx.dijkstra_path(G,'a','d'))
['a', 'c', 'd']
```

## 2.2.4 Drawing

While NetworkX is not designed as a network layout tool, we provide a simple interface to drawing packages and some simple layout algorithms. We interface to the excellent Graphviz layout tools like dot and neato with the (suggested) pygraphviz package or the pydot interface. Drawing can be done using external programs or the Matplotlib Python package. Interactive GUI interfaces are possible though not provided. The drawing tools are provided in the module *drawing*.

The basic drawing functions essentially place the nodes on a scatterplot using the positions in a dictionary or computed with a layout function. The edges are then lines between those dots.

```
>>> G=nx.cubical_graph()
>>> nx.draw(G)  # default spring_layout
>>> nx.draw(G,pos=nx.spectral_layout(G), nodecolor='r',edge_color='b')
```

See the examples for more ideas.

## 2.2.5 Data Structure

NetworkX uses a "dictionary of dictionaries of dictionaries" as the basic network data structure. This allows fast lookup with reasonable storage for large sparse networks. The keys are nodes so G[u] returns an adjacency dictionary keyed by neighbor to the edge attribute dictionary. The expression G[u][v] returns the edge attribute dictionary itself. A dictionary of lists would have also been possible, but not allowed fast edge detection nor convenient storage of edge data

Advantages of dict-of-dicts-of-dicts data structure:

- Find edges and remove edges with two dictionary look-ups.
- Prefer to "lists" because of fast lookup with sparse storage.
- Prefer to "sets" since data can be attached to edge.
- G[u][v] returns the edge attribute dictionary.
- n in G tests if node n is in graph G.
- for n in G: iterates through the graph.
- for nbr in G[n]: iterates through neighbors.

As an example, here is a representation of an undirected graph with the edges ('A','B'), ('B','C')

```
>>> G=nx.Graph()
>>> G.add_edge('A','B')
>>> G.add_edge('B','C')
>>> print(G.adj)
{'A': {'B': {}}, 'C': {'B': {}}, 'B': {'A': {}, 'C': {}}}
```

The data structure gets morphed slightly for each base graph class. For DiGraph two dict-of-dicts structures are provided, one for successors and one for predecessors. For MultiGraph/MultiDiGraph we use a dict-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-of-dicts-

dicts-of-dicts <sup>1</sup> where the third dictionary is keyed by an edge key identifier to the fourth dictionary which contains the edge attributes for that edge between the two nodes.

Graphs use a dictionary of attributes for each edge. We use a dict-of-dicts-of-dicts data structure with the inner dictionary storing "name-value" relationships for that edge.

```
>>> G=nx.Graph()
>>> G.add_edge(1,2,color='red',weight=0.84,size=300)
>>> print(G[1][2]['size'])
300
```

<sup>&</sup>lt;sup>1</sup> "It's dictionaries all the way down."

## **GRAPH TYPES**

NetworkX provides data structures and methods for storing graphs.

All NetworkX graph classes allow (hashable) Python objects as nodes. and any Python object can be assigned as an edge attribute.

The choice of graph class depends on the structure of the graph you want to represent.

## 3.1 Which graph class should I use?

Graph Type	NetworkX Class
Undirected Simple	Graph
Directed Simple	DiGraph
With Self-loops	Graph, DiGraph
With Parallel edges	MultiGraph, MultiDiGraph

## 3.2 Basic graph types

## 3.2.1 Graph – Undirected graphs with self loops

## **Overview**

Graph (data=None, \*\*attr)

Base class for undirected graphs.

A Graph stores nodes and edges with optional data, or attributes.

Graphs hold undirected edges. Self loops are allowed but multiple (parallel) edges are not.

Nodes can be arbitrary (hashable) Python objects with optional key/value attributes.

Edges are represented as links between nodes with optional key/value attributes.

Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse matrix, or a PyGraphviz graph.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

#### See also:

```
DiGraph, MultiGraph, MultiDiGraph
```

## **Examples**

Create a graph object that tracks the order nodes are added.

Create a graph object that tracks the order nodes are added and for each node track the order that neighbors are added.

```
>>> class OrderedGraph(nx.Graph):
... node_dict_factory = OrderedDict
... adjlist_dict_factory = OrderedDict
>>> G = OrderedGraph()
>>> G.add_nodes_from((2,1))
>>> G.nodes()
[2, 1]
>>> G.add_edges_from(((2,2), (2,1), (1,1)))
>>> G.edges()
[(2, 2), (2, 1), (1, 1)]
```

Create a low memory graph class that effectively disallows edge attributes by using a single attribute dict for all edges. This reduces the memory used, but you lose edge attributes.

```
>>> class ThinGraph(nx.Graph):
... all_edge_dict = {'weight': 1}
... def single_edge_dict(self):
... return self.all_edge_dict
... edge_attr_dict_factory = single_edge_dict
>>> G = ThinGraph()
>>> G.add_edge(2,1)
>>> G.edges(data= True)
[(1, 2, {'weight': 1})]
>>> G.add_edge(2,2)
>>> G[2][1] is G[2][2]
True
```

## Adding and removing nodes and edges

Graphinit([data])	Initialize a graph with edges, name, graph attributes.
<pre>Graph.add_node(n[, attr_dict])</pre>	Add a single node n and update node attributes.
Graph.add_nodes_from(nodes, **attr)	Add multiple nodes.
Graph.remove_node(n)	Remove node n.
	Continued on next pag

Table 3.1 – continued from previous page

Graph.remove_nodes_from(nodes)	Remove multiple nodes.
Graph.add_edge(u, v[, attr_dict])	Add an edge between u and v.
<pre>Graph.add_edges_from(ebunch[, attr_dict])</pre>	Add all the edges in ebunch.
<pre>Graph.add_weighted_edges_from(ebunch[, weight])</pre>	Add all the edges in ebunch as weighted edges with specified weights
Graph.remove_edge(u,v)	Remove the edge between u and v.
Graph.remove_edges_from(ebunch)	Remove all edges specified in ebunch.
Graph.add_star(nodes, **attr)	Add a star.
Graph.add_path(nodes, **attr)	Add a path.
<pre>Graph.add_cycle(nodes, **attr)</pre>	Add a cycle.
Graph.clear()	Remove all nodes and edges from the graph.

## init

```
Graph.___init___(data=None, **attr)
```

Initialize a graph with edges, name, graph attributes.

## Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse matrix, or a PyGraphviz graph.

name : string, optional (default='')

An optional name for the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

### See also:

convert

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G = nx.Graph(name='my graph')
>>> e = [(1,2),(2,3),(3,4)] # list of edges
>>> G = nx.Graph(e)
```

Arbitrary graph attribute pairs (key=value) may be assigned

```
>>> G=nx.Graph(e, day="Friday")
>>> G.graph
{'day': 'Friday'}
```

#### add node

```
Graph.add_node (n, attr_dict=None, **attr)
```

Add a single node n and update node attributes.

Parameters n: node

A node can be any hashable Python object except None.

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of node attributes. Key/value pairs will update existing data associated with the node.

attr: keyword arguments, optional

Set or change attributes using key=value.

#### See also:

```
add_nodes_from
```

#### **Notes**

A hashable object is one that can be used as a key in a Python dictionary. This includes strings, numbers, tuples of strings and numbers, etc.

On many platforms hashable items also include mutables such as NetworkX Graphs, though one should be careful that the hash doesn't change on mutables.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_node(1)
>>> G.add_node('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_node(K3)
>>> G.number_of_nodes()
3
```

Use keywords set/change node attributes:

```
>>> G.add_node(1,size=10)
>>> G.add_node(3,weight=0.4,UTM=('13S',382871,3972649))
```

## add nodes from

```
Graph.add_nodes_from (nodes, **attr)
Add multiple nodes.
```

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). OR A container of (node, attribute dict) tuples. Node attributes are updated using the attribute dict.

attr: keyword arguments, optional (default= no attributes)

Update attributes for all nodes in nodes. Node attributes specified in nodes as a tuple take precedence over attributes specified generally.

## See also:

```
add_node
```

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_nodes_from('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_nodes_from(K3)
>>> sorted(G.nodes(),key=str)
[0, 1, 2, 'H', 'e', 'l', 'o']
```

Use keywords to update specific node attributes for every node.

```
>>> G.add_nodes_from([1,2], size=10)
>>> G.add_nodes_from([3,4], weight=0.4)
```

Use (node, attrdict) tuples to update attributes for specific nodes.

```
>>> G.add_nodes_from([(1,dict(size=11)), (2,{'color':'blue'})])
>>> G.node[1]['size']
11
>>> H = nx.Graph()
>>> H.add_nodes_from(G.nodes(data=True))
>>> H.node[1]['size']
11
```

#### remove node

```
Graph.remove_node(n)
```

Remove node n.

Removes the node n and all adjacent edges. Attempting to remove a non-existent node will raise an exception.

Parameters n: node

A node in the graph

## Raises NetworkXError

If n is not in the graph.

## See also:

```
remove nodes from
```

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.edges()
[(0, 1), (1, 2)]
>>> G.remove_node(1)
>>> G.edges()
[]
```

## remove\_nodes\_from

```
Graph.remove_nodes_from (nodes)
```

Remove multiple nodes.

## Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). If a node in the container is not in the graph it is silently ignored.

#### See also:

```
remove_node
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> e = G.nodes()
>>> e
[0, 1, 2]
>>> G.remove_nodes_from(e)
>>> G.nodes()
[]
```

## add edge

```
Graph.add_edge (u, v, attr_dict=None, **attr)
```

Add an edge between u and v.

The nodes u and v will be automatically added if they are not already in the graph.

Edge attributes can be specified with keywords or by providing a dictionary with key/value pairs. See examples below.

#### **Parameters u,v**: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

```
attr_dict : dictionary, optional (default= no attributes)
```

Dictionary of edge attributes. Key/value pairs will update existing data associated with the edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

#### See also:

```
add_edges_from add a collection of edges
```

#### **Notes**

Adding an edge that already exists updates the edge data.

Many NetworkX algorithms designed for weighted graphs use as the edge weight a numerical value assigned to a keyword which by default is 'weight'.

## **Examples**

The following all add the edge e=(1,2) to graph G:

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> e = (1,2)
>>> G.add_edge(1, 2) # explicit two-node form
>>> G.add_edge(*e) # single edge as tuple of two nodes
>>> G.add_edges_from([(1,2)]) # add edges from iterable container
```

Associate data to edges using keywords:

```
>>> G.add_edge(1, 2, weight=3)
>>> G.add_edge(1, 3, weight=7, capacity=15, length=342.7)
```

#### add edges from

```
Graph.add_edges_from(ebunch, attr_dict=None, **attr)
Add all the edges in ebunch.
```

Parameters ebunch: container of edges

Each edge given in the container will be added to the graph. The edges must be given as as 2-tuples (u,v) or 3-tuples (u,v,d) where d is a dictionary containing edge data.

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of edge attributes. Key/value pairs will update existing data associated with each edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

## See also:

```
add_edge add a single edge
add_weighted_edges_from convenient way to add weighted edges
```

#### **Notes**

Adding the same edge twice has no effect but any edge data will be updated when each duplicate edge is added. Edge attributes specified in edges take precedence over attributes specified generally.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edges_from([(0,1),(1,2)]) # using a list of edge tuples
>>> e = zip(range(0,3),range(1,4))
>>> G.add_edges_from(e) # Add the path graph 0-1-2-3

Associate data to edges
>>> G.add_edges_from([(1,2),(2,3)], weight=3)
>>> G.add_edges_from([(3,4),(1,4)], label='WN2898')
```

#### add weighted edges from

```
Graph.add_weighted_edges_from(ebunch, weight='weight', **attr)
```

Add all the edges in ebunch as weighted edges with specified weights.

## Parameters ebunch: container of edges

Each edge given in the list or container will be added to the graph. The edges must be given as 3-tuples (u,v,w) where w is a number.

```
weight : string, optional (default= 'weight')
```

The attribute name for the edge weights to be added.

attr: keyword arguments, optional (default= no attributes)

Edge attributes to add/update for all edges.

#### See also:

```
add_edge add a single edge
add_edges_from add multiple edges
```

## **Notes**

Adding the same edge twice for Graph/DiGraph simply updates the edge data. For MultiGraph/MultiDiGraph, duplicate edges are stored.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_weighted_edges_from([(0,1,3.0),(1,2,7.5)])
```

## remove edge

```
Graph.remove\_edge(u, v)
```

Remove the edge between u and v.

### Parameters u,v: nodes

Remove the edge between nodes u and v.

## Raises NetworkXError

If there is not an edge between u and v.

#### See also:

remove\_edges\_from remove a collection of edges

```
>>> G = nx.Graph() # or DiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.remove_edge(0,1)
>>> e = (1,2)
>>> G.remove_edge(*e) # unpacks e from an edge tuple
>>> e = (2,3,{'weight':7}) # an edge with attribute data
>>> G.remove_edge(*e[:2]) # select first part of edge tuple
```

## remove edges from

```
Graph.remove_edges_from(ebunch)
```

Remove all edges specified in ebunch.

## Parameters ebunch: list or container of edge tuples

Each edge given in the list or container will be removed from the graph. The edges can be:

- 2-tuples (u,v) edge between u and v.
- 3-tuples (u,v,k) where k is ignored.

## See also:

remove\_edge remove a single edge

#### **Notes**

Will fail silently if an edge in ebunch is not in the graph.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> ebunch=[(1,2),(2,3)]
>>> G.remove_edges_from(ebunch)
```

## add star

```
Graph.add_star(nodes, **attr)
```

Add a star.

The first node in nodes is the middle of the star. It is connected to all other nodes.

Parameters nodes: iterable container

A container of nodes.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in star.

## See also:

```
add_path, add_cycle
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_star([0,1,2,3])
>>> G.add_star([10,11,12],weight=2)
```

## add\_path

```
Graph.add_path (nodes, **attr)
Add a path.
```

Parameters nodes: iterable container

A container of nodes. A path will be constructed from the nodes (in order) and added to the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in path.

#### See also:

```
add_star, add_cycle
```

## **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.add_path([10,11,12], weight=7)
```

## add cycle

```
Graph.add_cycle (nodes, **attr)
Add a cycle.
```

#### Parameters nodes: iterable container

A container of nodes. A cycle will be constructed from the nodes (in order) and added to the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in cycle.

### See also:

```
add_path, add_star
```

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_cycle([0,1,2,3])
>>> G.add_cycle([10,11,12], weight=7)
```

#### clear

```
Graph.clear()
```

Remove all nodes and edges from the graph.

This also removes the name, and all graph, node, and edge attributes.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.clear()
>>> G.nodes()
[]
>>> G.edges()
[]
```

## Iterating over nodes and edges

Graph.nodes([data])	Return a list of the nodes in the graph.
Graph.nodes_iter([data])	Return an iterator over the nodes.
Graphiter()	Iterate over the nodes.
<pre>Graph.edges([nbunch, data, default])</pre>	Return a list of edges.
<pre>Graph.edges_iter([nbunch, data, default])</pre>	Return an iterator over the edges.
<pre>Graph.get_edge_data(u, v[, default])</pre>	Return the attribute dictionary associated with edge (u,v).
Graph.neighbors(n)	Return a list of the nodes connected to the node n.
Graph.neighbors_iter(n)	Return an iterator over all neighbors of node n.
Graphgetitem(n)	Return a dict of neighbors of node n.
Graph.adjacency_list()	Return an adjacency list representation of the graph.
Graph.adjacency_iter()	Return an iterator of (node, adjacency dict) tuples for all nodes.
Graph.nbunch_iter([nbunch])	Return an iterator of nodes contained in nbunch that are also in the graph.

## nodes

Graph.nodes (data=False)

Return a list of the nodes in the graph.

Parameters data: boolean, optional (default=False)

If False return a list of nodes. If True return a two-tuple of node and node data dictionary

Returns nlist: list

A list of nodes. If data=True a list of two-tuples containing (node, node data dictionary).

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.nodes()
[0, 1, 2]
>>> G.add_node(1, time='5pm')
```

```
>>> G.nodes(data=True)
[(0, {}), (1, {'time': '5pm'}), (2, {})]
```

## nodes\_iter

```
Graph.nodes_iter(data=False)
```

Return an iterator over the nodes.

Parameters data: boolean, optional (default=False)

If False the iterator returns nodes. If True return a two-tuple of node and node data dictionary

**Returns niter**: iterator

An iterator over nodes. If data=True the iterator gives two-tuples containing (node, node data, dictionary)

#### **Notes**

If the node data is not required it is simpler and equivalent to use the expression 'for n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
```

#### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])

>>> [d for n,d in G.nodes_iter(data=True)]
[{}, {}, {}]
```

\_\_iter\_\_

Graph.\_\_\_iter\_\_()

Iterate over the nodes. Use the expression 'for n in G'.

**Returns niter**: iterator

An iterator over all nodes in the graph.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
```

#### edges

```
Graph.edges (nbunch=None, data=False, default=None)
Return a list of edges.
```

Edges are returned as tuples with optional data in the order (node, neighbor, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,data) (True).

**Returns** edge\_list: list of edge tuples

Edges that are adjacent to any node in nbunch, or a list of all edges if nbunch is not specified.

### See also:

edges\_iter return an iterator over the edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> G.edges()
[(0, 1), (1, 2), (2, 3)]
>>> G.edges(data=True) # default edge data is {} (empty dictionary)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> G.edges([0,3])
[(0, 1), (3, 2)]
>>> G.edges(0)
[(0, 1)]
```

## edges\_iter

```
Graph.edges_iter(nbunch=None, data=False, default=None)
```

Return an iterator over the edges.

Edges are returned as tuples with optional data in the order (node, neighbor, data).

Parameters nbunch: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edge\_iter: iterator

An iterator of (u,v) or (u,v,d) tuples of edges.

## See also:

edges return a list of edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.Graph()  # or MultiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges_iter([0,3]))
[(0, 1), (3, 2)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

## get\_edge\_data

```
Graph.get_edge_data(u, v, default=None)
```

Return the attribute dictionary associated with edge (u,v).

Parameters u,v: nodes

default: any Python object (default=None)

Value to return if the edge (u,v) is not found.

Returns edge\_dict : dictionary

The edge attribute dictionary.

## **Notes**

It is faster to use G[u][v].

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G[0][1]
{}
```

Warning: Assigning G[u][v] corrupts the graph data structure. But it is safe to assign attributes to that dictionary,

```
>>> G[0][1]['weight'] = 7
>>> G[0][1]['weight']
7
>>> G[1][0]['weight']
7
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.get_edge_data(0,1) # default edge data is {}
{}
>>> e = (0,1)
>>> G.get_edge_data(*e) # tuple form
{}
>>> G.get_edge_data('a','b',default=0) # edge not in graph, return 0
0
```

### neighbors

## Graph.neighbors (n)

Return a list of the nodes connected to the node n.

Parameters n: node

A node in the graph

Returns nlist: list

A list of nodes that are adjacent to n.

## Raises NetworkXError

If the node n is not in the graph.

## **Notes**

It is usually more convenient (and faster) to access the adjacency dictionary as G[n]:

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=7)
>>> G['a']
{'b': {'weight': 7}}
```

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.neighbors(0)
[1]
```

### neighbors iter

## Graph.neighbors\_iter(n)

Return an iterator over all neighbors of node n.

## **Notes**

It is faster to use the idiom "in G[0]", e.g.

```
>>> G = nx.path_graph(4)
>>> [n for n in G[0]]
[1]
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [n for n in G.neighbors_iter(0)]
[1]
```

## getitem

```
Graph.___getitem___(n)
```

Return a dict of neighbors of node n. Use the expression 'G[n]'.

Parameters n: node

A node in the graph.

Returns adj\_dict : dictionary

The adjacency dictionary for nodes connected to n.

## **Notes**

G[n] is similar to G.neighbors(n) but the internal data dictionary is returned instead of a list.

Assigning G[n] will corrupt the internal graph data structure. Use G[n] for reading data only.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G[0]
{1: {}}
```

#### adjacency list

```
Graph.adjacency_list()
```

Return an adjacency list representation of the graph.

The output adjacency list is in the order of G.nodes(). For directed graphs, only outgoing adjacencies are included.

Returns adj list: lists of lists

The adjacency structure of the graph as a list of lists.

## See also:

```
adjacency_iter
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.adjacency_list() # in order given by G.nodes()
[[1], [0, 2], [1, 3], [2]]
```

## adjacency iter

```
Graph.adjacency_iter()
```

Return an iterator of (node, adjacency dict) tuples for all nodes.

This is the fastest way to look at every edge. For directed graphs, only outgoing adjacencies are included.

Returns adj\_iter: iterator

An iterator of (node, adjacency dictionary) for all nodes in the graph.

#### See also:

```
adjacency_list
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [(n,nbrdict) for n,nbrdict in G.adjacency_iter()]
[(0, {1: {}}), (1, {0: {}, 2: {}}), (2, {1: {}, 3: {}}), (3, {2: {}})]
```

## nbunch iter

```
Graph.nbunch_iter(nbunch=None)
```

Return an iterator of nodes contained in nbunch that are also in the graph.

The nodes in nbunch are checked for membership in the graph and if not are silently ignored.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

Returns niter: iterator

An iterator over nodes in nbunch that are also in the graph. If nbunch is None, iterate over all nodes in the graph.

#### Raises NetworkXError

If nbunch is not a node or or sequence of nodes. If a node in nbunch is not hashable.

#### See also:

```
Graph.__iter__
```

#### **Notes**

When nbunch is an iterator, the returned iterator yields values directly from nbunch, becoming exhausted when nbunch is exhausted.

To test whether nbunch is a single node, one can use "if nbunch in self:", even after processing with this routine.

If nbunch is not a node or a (possibly empty) sequence/iterator or None, a NetworkXError is raised. Also, if any object in nbunch is not hashable, a NetworkXError is raised.

## Information about graph structure

Graph.has_node(n)	Return True if the graph contains the node n.
Graphcontains(n)	Return True if n is a node, False otherwise.
$Graph.has\_edge(u, v)$	Return True if the edge (u,v) is in the graph.
Graph.order()	Return the number of nodes in the graph.
<pre>Graph.number_of_nodes()</pre>	Return the number of nodes in the graph.
Graphlen()	Return the number of nodes.
<pre>Graph.degree([nbunch, weight])</pre>	Return the degree of a node or nodes.
<pre>Graph.degree_iter([nbunch, weight])</pre>	Return an iterator for (node, degree).
<pre>Graph.size([weight])</pre>	Return the number of edges.
$Graph.number_of_edges([u, v])$	Return the number of edges between two nodes.
Graph.nodes_with_selfloops()	Return a list of nodes with self loops.
<pre>Graph.selfloop_edges([data, default])</pre>	Return a list of selfloop edges.
<pre>Graph.number_of_selfloops()</pre>	Return the number of selfloop edges.

## has node

```
Graph.has_node(n)
```

Return True if the graph contains the node n.

Parameters n: node

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.has_node(0)
True
```

It is more readable and simpler to use

```
>>> 0 in G
True
```

## contains

```
Graph.__contains__(n)
```

Return True if n is a node, False otherwise. Use the expression 'n in G'.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> 1 in G
True
```

## has\_edge

```
Graph.has_edge (u, v)
```

Return True if the edge (u,v) is in the graph.

Parameters u,v: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

Returns edge\_ind: bool

True if edge is in the graph, False otherwise.

## **Examples**

Can be called either using two nodes u,v or edge tuple (u,v)

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.has_edge(0,1)  # using two nodes
True
>>> e = (0,1)
>>> G.has_edge(*e)  # e is a 2-tuple (u,v)
True
>>> e = (0,1,{'weight':7})
>>> G.has_edge(*e[:2])  # e is a 3-tuple (u,v,data_dictionary)
True
```

The following syntax are all equivalent:

```
>>> G.has_edge(0,1)
True
>>> 1 in G[0] # though this gives KeyError if 0 not in G
True
```

#### order

```
Graph.order()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

## See also:

```
number_of_nodes, __len__
```

### number of nodes

```
Graph.number_of_nodes()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

## See also:

```
order, __len__
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> len(G)
3
```

### \_\_len\_\_

```
Graph.___len___()
```

Return the number of nodes. Use the expression 'len(G)'.

Returns nnodes: int

The number of nodes in the graph.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> len(G)
4
```

## degree

Graph.degree (nbunch=None, weight=None)

Return the degree of a node or nodes.

The node degree is the number of edges adjacent to that node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight: string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd: dictionary, or number

A dictionary with nodes as keys and degree as values or a number if a single node is specified.

#### **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.degree(0)
1
>>> G.degree([0,1])
{0: 1, 1: 2}
>>> list(G.degree([0,1]).values())
[1, 2]
```

## degree iter

```
Graph.degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, degree).

The node degree is the number of edges adjacent to the node.

Parameters nbunch: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight: string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, degree).

## See also:

degree

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> list(G.degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.degree_iter([0,1]))
[(0, 1), (1, 2)]
```

#### size

```
Graph.size(weight=None)
```

Return the number of edges.

**Parameters weight**: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

## Returns nedges: int

The number of edges or sum of edge weights in the graph.

#### See also:

```
number_of_edges
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.size()
3
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=2)
>>> G.add_edge('b','c',weight=4)
>>> G.size()
2
>>> G.size(weight='weight')
6.0
```

## number of edges

```
Graph.number of edges (u=None, v=None)
```

Return the number of edges between two nodes.

**Parameters u,v**: nodes, optional (default=all edges)

If u and v are specified, return the number of edges between u and v. Otherwise return the total number of all edges.

## Returns nedges: int

The number of edges in the graph. If nodes u and v are specified return the number of edges between those nodes.

## See also:

size

30

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.number_of_edges()
```

```
3
>>> G.number_of_edges(0,1)
1
>>> e = (0,1)
>>> G.number_of_edges(*e)
1
```

### nodes\_with\_selfloops

### Graph.nodes\_with\_selfloops()

Return a list of nodes with self loops.

A node with a self loop has an edge with both ends adjacent to that node.

Returns nodelist: list

A list of nodes with self loops.

#### See also:

```
selfloop edges, number of selfloops
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.nodes_with_selfloops()
[1]
```

# selfloop\_edges

Graph.selfloop\_edges (data=False, default=None)

Return a list of selfloop edges.

A selfloop edge has the same node at both ends.

Parameters data: string or bool, optional (default=False)

Return selfloop edges as two tuples (u,v) (data=False) or three-tuples (u,v,datadict) (data=True) or three-tuples (u,v,datavalue) (data='attrname')

default : value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edgelist: list of edge tuples

A list of all selfloop edges.

### See also:

```
nodes_with_selfloops, number_of_selfloops
```

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.selfloop_edges()
[(1, 1)]
>>> G.selfloop_edges(data=True)
[(1, 1, {})]
```

# number\_of\_selfloops

```
Graph.number_of_selfloops()
```

Return the number of selfloop edges.

A selfloop edge has the same node at both ends.

Returns nloops: int

The number of selfloops.

### See also:

```
nodes_with_selfloops, selfloop_edges
```

# **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.number_of_selfloops()
1
```

# Making copies and subgraphs

Graph.copy()	Return a copy of the graph.
Graph.to_undirected()	Return an undirected copy of the graph.
Graph.to_directed()	Return a directed representation of the graph.
Graph.subgraph(nbunch)	Return the subgraph induced on nodes in nbunch.

# сору

```
Graph.copy()
```

Return a copy of the graph.

Returns G: Graph

A copy of the graph.

# See also:

to\_directed return a directed copy of the graph.

#### **Notes**

This makes a complete copy of the graph including all of the node or edge attributes.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.copy()
```

### to\_undirected

### Graph.to\_undirected()

Return an undirected copy of the graph.

Returns G: Graph/MultiGraph

A deepcopy of the graph.

#### See also:

```
copy, add_edge, add_edges_from
```

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar G=DiGraph(D) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

# **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]
>>> G2 = H.to_undirected()
>>> G2.edges()
[(0, 1)]
```

### to\_directed

# Graph.to\_directed()

Return a directed representation of the graph.

# Returns G: DiGraph

A directed graph with the same name, same nodes, and with each edge (u,v,data) replaced by two directed edges (u,v,data) and (v,u,data).

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar D=DiGraph(G) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

### **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]

If already directed, return a (deep) copy
>>> G = nx.DiGraph() # or MultiDiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
```

#### subgraph

# Graph.subgraph (nbunch)

>>> H.edges()
[(0, 1)]

Return the subgraph induced on nodes in nbunch.

The induced subgraph of the graph contains the nodes in nbunch and the edges between those nodes.

**Parameters nbunch**: list, iterable

A container of nodes which will be iterated through once.

Returns G: Graph

A subgraph of the graph with the same edge attributes.

### **Notes**

The graph, edge or node attributes just point to the original graph. So changes to the node or edge structure will not be reflected in the original graph while changes to the attributes will.

To create a subgraph with its own copy of the edge/node attributes use: nx.Graph(G.subgraph(nbunch))

If edge attributes are containers, a deep copy can be obtained using: G.subgraph(nbunch).copy()

For an inplace reduction of a graph to a subgraph you can remove nodes: G.remove\_nodes\_from([ n in G if n not in set(nbunch)])

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.subgraph([0,1,2])
>>> H.edges()
[(0, 1), (1, 2)]
```

# 3.2.2 DiGraph - Directed graphs with self loops

#### Overview

```
DiGraph (data=None, **attr)
```

Base class for directed graphs.

A DiGraph stores nodes and edges with optional data, or attributes.

DiGraphs hold directed edges. Self loops are allowed but multiple (parallel) edges are not.

Nodes can be arbitrary (hashable) Python objects with optional key/value attributes.

Edges are represented as links between nodes with optional key/value attributes.

# Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse matrix, or a PyGraphviz graph.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

# See also:

```
Graph, MultiGraph, MultiDiGraph
```

# **Examples**

Create a graph object that tracks the order nodes are added.

```
>>> from collections import OrderedDict
>>> class OrderedNodeGraph(nx.Graph):
... node_dict_factory=OrderedDict
>>> G=OrderedNodeGraph()
>>> G.add_nodes_from((2,1))
>>> G.nodes()
[2, 1]
>>> G.add_edges_from(((2,2), (2,1), (1,1)))
>>> G.edges()
[(2, 1), (2, 2), (1, 1)]
```

Create a graph object that tracks the order nodes are added and for each node track the order that neighbors are added.

```
>>> class OrderedGraph(nx.Graph):
...    node_dict_factory = OrderedDict
...    adjlist_dict_factory = OrderedDict
>>> G = OrderedGraph()
>>> G.add_nodes_from((2,1))
>>> G.nodes()
[2, 1]
>>> G.add_edges_from(((2,2), (2,1), (1,1)))
>>> G.edges()
[(2, 2), (2, 1), (1, 1)]
```

Create a low memory graph class that effectively disallows edge attributes by using a single attribute dict for all edges. This reduces the memory used, but you lose edge attributes.

```
>>> class ThinGraph(nx.Graph):
... all_edge_dict = {'weight': 1}
... def single_edge_dict(self):
... return self.all_edge_dict
... edge_attr_dict_factory = single_edge_dict
>>> G = ThinGraph()
>>> G.add_edge(2,1)
>>> G.edges(data= True)
[(1, 2, {'weight': 1})]
>>> G.add_edge(2,2)
>>> G[2][1] is G[2][2]
True
```

# Adding and removing nodes and edges

DiGraphinit([data])	Initialize a graph with edges, name, graph attributes.
DiGraph.add_node(n[, attr_dict])	Add a single node n and update node attributes.
DiGraph.add_nodes_from(nodes, **attr)	Add multiple nodes.
DiGraph.remove_node(n)	Remove node n.
DiGraph.remove_nodes_from(nbunch)	Remove multiple nodes.
DiGraph.add_edge(u, v[, attr_dict])	Add an edge between u and v.
<pre>DiGraph.add_edges_from(ebunch[, attr_dict])</pre>	Add all the edges in ebunch.
<pre>DiGraph.add_weighted_edges_from(ebunch[, weight])</pre>	Add all the edges in ebunch as weighted edges with specified weighted
DiGraph.remove_edge $(u, v)$	Remove the edge between u and v.
DiGraph.remove_edges_from(ebunch)	Remove all edges specified in ebunch.
DiGraph.add_star(nodes, **attr)	Add a star.
DiGraph.add_path(nodes, **attr)	Add a path.
DiGraph.add_cycle(nodes, **attr)	Add a cycle.
DiGraph.clear()	Remove all nodes and edges from the graph.

#### init

```
DiGraph. init (data=None, **attr)
```

Initialize a graph with edges, name, graph attributes.

# Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse

```
matrix, or a PyGraphviz graph.
```

name : string, optional (default='')

An optional name for the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

#### See also:

convert

#### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G = nx.Graph(name='my graph')
>>> e = [(1,2),(2,3),(3,4)] # list of edges
>>> G = nx.Graph(e)
```

Arbitrary graph attribute pairs (key=value) may be assigned

```
>>> G=nx.Graph(e, day="Friday")
>>> G.graph
{'day': 'Friday'}
```

#### add node

```
DiGraph.add_node (n, attr_dict=None, **attr)
```

Add a single node n and update node attributes.

### Parameters n: node

A node can be any hashable Python object except None.

```
attr_dict : dictionary, optional (default= no attributes)
```

Dictionary of node attributes. Key/value pairs will update existing data associated with the node.

attr: keyword arguments, optional

Set or change attributes using key=value.

# See also:

```
add_nodes_from
```

#### Notes

A hashable object is one that can be used as a key in a Python dictionary. This includes strings, numbers, tuples of strings and numbers, etc.

On many platforms hashable items also include mutables such as NetworkX Graphs, though one should be careful that the hash doesn't change on mutables.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_node(1)
>>> G.add_node('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_node(K3)
>>> G.number_of_nodes()
```

Use keywords set/change node attributes:

```
>>> G.add_node(1,size=10)
>>> G.add_node(3,weight=0.4,UTM=('13S',382871,3972649))
```

#### add nodes from

```
DiGraph.add_nodes_from (nodes, **attr)
Add multiple nodes.
```

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). OR A container of (node, attribute dict) tuples. Node attributes are updated using the attribute dict.

attr: keyword arguments, optional (default= no attributes)

Update attributes for all nodes in nodes. Node attributes specified in nodes as a tuple take precedence over attributes specified generally.

#### See also:

add\_node

### **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_nodes_from('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_nodes_from(K3)
>>> sorted(G.nodes(),key=str)
[0, 1, 2, 'H', 'e', 'l', 'o']
```

Use keywords to update specific node attributes for every node.

```
>>> G.add_nodes_from([1,2], size=10)
>>> G.add_nodes_from([3,4], weight=0.4)
```

Use (node, attrdict) tuples to update attributes for specific nodes.

```
>>> G.add_nodes_from([(1,dict(size=11)), (2,{'color':'blue'})])
>>> G.node[1]['size']
11
>>> H = nx.Graph()
>>> H.add_nodes_from(G.nodes(data=True))
>>> H.node[1]['size']
11
```

#### remove node

```
{\tt DiGraph.remove\_node}\,(n)
```

Remove node n.

Removes the node n and all adjacent edges. Attempting to remove a non-existent node will raise an exception.

Parameters n: node

A node in the graph

#### Raises NetworkXError

If n is not in the graph.

# See also:

```
remove_nodes_from
```

# **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.edges()
[(0, 1), (1, 2)]
>>> G.remove_node(1)
>>> G.edges()
[]
```

# remove\_nodes\_from

```
DiGraph.remove_nodes_from(nbunch)
```

Remove multiple nodes.

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). If a node in the container is not in the graph it is silently ignored.

### See also:

```
remove_node
```

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> e = G.nodes()
>>> e
[0, 1, 2]
>>> G.remove_nodes_from(e)
>>> G.nodes()
[]
```

#### add edge

```
DiGraph.add_edge (u, v, attr\_dict=None, **attr)
Add an edge between u and v.
```

The nodes u and v will be automatically added if they are not already in the graph.

Edge attributes can be specified with keywords or by providing a dictionary with key/value pairs. See examples below.

#### Parameters u,v: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

```
attr_dict : dictionary, optional (default= no attributes)
```

Dictionary of edge attributes. Key/value pairs will update existing data associated with the edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

#### See also:

```
add_edges_from add a collection of edges
```

#### **Notes**

Adding an edge that already exists updates the edge data.

Many NetworkX algorithms designed for weighted graphs use as the edge weight a numerical value assigned to a keyword which by default is 'weight'.

#### **Examples**

The following all add the edge e=(1,2) to graph G:

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> e = (1,2)
>>> G.add_edge(1, 2) # explicit two-node form
>>> G.add_edge(*e) # single edge as tuple of two nodes
>>> G.add_edges_from([(1,2)]) # add edges from iterable container
```

Associate data to edges using keywords:

```
>>> G.add_edge(1, 2, weight=3)
>>> G.add_edge(1, 3, weight=7, capacity=15, length=342.7)
```

# add\_edges\_from

```
DiGraph.add_edges_from(ebunch, attr_dict=None, **attr)
Add all the edges in ebunch.
```

Parameters ebunch: container of edges

Each edge given in the container will be added to the graph. The edges must be given as as 2-tuples (u,v) or 3-tuples (u,v,d) where d is a dictionary containing edge data.

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of edge attributes. Key/value pairs will update existing data associated with each edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

### See also:

```
add_edge add a single edge
add_weighted_edges_from convenient way to add weighted edges
```

#### **Notes**

Adding the same edge twice has no effect but any edge data will be updated when each duplicate edge is added. Edge attributes specified in edges take precedence over attributes specified generally.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edges_from([(0,1),(1,2)]) # using a list of edge tuples
>>> e = zip(range(0,3),range(1,4))
>>> G.add_edges_from(e) # Add the path graph 0-1-2-3
Associate data to edges
```

```
>>> G.add_edges_from([(1,2),(2,3)], weight=3)
>>> G.add_edges_from([(3,4),(1,4)], label='WN2898')
```

# add\_weighted\_edges\_from

```
DiGraph.add_weighted_edges_from(ebunch, weight='weight', **attr)
```

Add all the edges in ebunch as weighted edges with specified weights.

Parameters ebunch: container of edges

Each edge given in the list or container will be added to the graph. The edges must be given as 3-tuples (u,v,w) where w is a number.

```
weight : string, optional (default= 'weight')
```

The attribute name for the edge weights to be added.

attr : keyword arguments, optional (default= no attributes)

Edge attributes to add/update for all edges.

# See also:

```
add_edge add a single edge
add_edges_from add multiple edges
```

#### **Notes**

Adding the same edge twice for Graph/DiGraph simply updates the edge data. For MultiGraph/MultiDiGraph, duplicate edges are stored.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_weighted_edges_from([(0,1,3.0),(1,2,7.5)])
```

### remove edge

```
DiGraph.remove_edge (u, v)
```

Remove the edge between u and v.

#### Parameters u,v: nodes

Remove the edge between nodes u and v.

#### Raises NetworkXError

If there is not an edge between u and v.

#### See also:

remove\_edges\_from remove a collection of edges

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.remove_edge(0,1)
>>> e = (1,2)
>>> G.remove_edge(*e) # unpacks e from an edge tuple
>>> e = (2,3,{'weight':7}) # an edge with attribute data
>>> G.remove_edge(*e[:2]) # select first part of edge tuple
```

# remove edges from

```
DiGraph.remove_edges_from(ebunch)
```

Remove all edges specified in ebunch.

# Parameters ebunch: list or container of edge tuples

Each edge given in the list or container will be removed from the graph. The edges can be:

- 2-tuples (u,v) edge between u and v.
- 3-tuples (u,v,k) where k is ignored.

# See also:

remove\_edge remove a single edge

#### **Notes**

Will fail silently if an edge in ebunch is not in the graph.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> ebunch=[(1,2),(2,3)]
>>> G.remove_edges_from(ebunch)
```

#### add star

```
DiGraph.add_star(nodes, **attr)
Add a star.
```

The first node in nodes is the middle of the star. It is connected to all other nodes.

Parameters nodes: iterable container

A container of nodes.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in star.

#### See also:

```
add_path, add_cycle
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_star([0,1,2,3])
>>> G.add_star([10,11,12],weight=2)
```

# add\_path

```
DiGraph.add_path (nodes, **attr)
Add a path.
```

Parameters nodes: iterable container

A container of nodes. A path will be constructed from the nodes (in order) and added to the graph.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to every edge in path.

#### See also:

```
add_star,add_cycle
```

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.add_path([10,11,12], weight=7)
```

# add\_cycle

```
DiGraph.add_cycle (nodes, **attr)
Add a cycle.
```

#### Parameters nodes: iterable container

A container of nodes. A cycle will be constructed from the nodes (in order) and added to the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in cycle.

#### See also:

```
add_path, add_star
```

# **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_cycle([0,1,2,3])
>>> G.add_cycle([10,11,12], weight=7)
```

### clear

```
DiGraph.clear()
```

Remove all nodes and edges from the graph.

This also removes the name, and all graph, node, and edge attributes.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.clear()
>>> G.nodes()
[]
>>> G.edges()
[]
```

# Iterating over nodes and edges

DiGraph.nodes([data])	Return a list of the nodes in the graph.	
DiGraph.nodes_iter([data])	Return an iterator over the nodes.	
		Continued on next page

Table 3.6 – continued from previous page

Iterate over the nodes.
Return a list of edges.
Return an iterator over the edges.
Return a list of edges.
Return an iterator over the edges.
Return a list of the incoming edges.
Return an iterator over the incoming edges.
Return the attribute dictionary associated with edge (u,v).
Return a list of successor nodes of n.
Return an iterator over successor nodes of n.
Return a dict of neighbors of node n.
Return a list of successor nodes of n.
Return an iterator over successor nodes of n.
Return a list of predecessor nodes of n.
Return an iterator over predecessor nodes of n.
Return an adjacency list representation of the graph.
Return an iterator of (node, adjacency dict) tuples for all nodes.
Return an iterator of nodes contained in nbunch that are also in the graph.

#### nodes

DiGraph.nodes (data=False)

Return a list of the nodes in the graph.

Parameters data: boolean, optional (default=False)

If False return a list of nodes. If True return a two-tuple of node and node data dictionary

Returns nlist: list

A list of nodes. If data=True a list of two-tuples containing (node, node data dictionary).

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.nodes()
[0, 1, 2]
>>> G.add_node(1, time='5pm')
>>> G.nodes(data=True)
[(0, {}), (1, {'time': '5pm'}), (2, {})]
```

### nodes\_iter

DiGraph.nodes\_iter(data=False)

Return an iterator over the nodes.

**Parameters data**: boolean, optional (default=False)

If False the iterator returns nodes. If True return a two-tuple of node and node data dictionary

Returns niter: iterator

An iterator over nodes. If data=True the iterator gives two-tuples containing (node, node data, dictionary)

#### **Notes**

If the node data is not required it is simpler and equivalent to use the expression 'for n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])

>>> [d for n,d in G.nodes_iter(data=True)]
[{}, {}, {}]
```

# \_\_iter\_\_

```
DiGraph.__iter__()
```

Iterate over the nodes. Use the expression 'for n in G'.

Returns niter: iterator

An iterator over all nodes in the graph.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
```

### edges

DiGraph.edges (nbunch=None, data=False, default=None)

Return a list of edges.

Edges are returned as tuples with optional data in the order (node, neighbor, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,data) (True).

**Returns** edge\_list: list of edge tuples

Edges that are adjacent to any node in nbunch, or a list of all edges if nbunch is not specified.

#### See also:

edges\_iter return an iterator over the edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

# **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> G.edges()
[(0, 1), (1, 2), (2, 3)]
>>> G.edges(data=True) # default edge data is {} (empty dictionary)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> G.edges([0,3])
[(0, 1), (3, 2)]
>>> G.edges(0)
[(0, 1)]
```

### edges\_iter

DiGraph.edges\_iter (nbunch=None, data=False, default=None)

Return an iterator over the edges.

Edges are returned as tuples with optional data in the order (node, neighbor, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edge\_iter: iterator

An iterator of (u,v) or (u,v,d) tuples of edges.

### See also:

edges return a list of edges

### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

```
>>> G = nx.DiGraph() # or MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges_iter([0,2]))
[(0, 1), (2, 3)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

### out edges

DiGraph.out\_edges (nbunch=None, data=False, default=None)
Return a list of edges.

Edges are returned as tuples with optional data in the order (node, neighbor, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,data) (True).

**Returns** edge\_list: list of edge tuples

Edges that are adjacent to any node in nbunch, or a list of all edges if nbunch is not specified.

### See also:

edges\_iter return an iterator over the edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> G.edges()
[(0, 1), (1, 2), (2, 3)]
>>> G.edges(data=True) # default edge data is {} (empty dictionary)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> G.edges([0,3])
```

```
[(0, 1), (3, 2)]

>>> G.edges(0)

[(0, 1)]
```

#### out edges iter

DiGraph.out\_edges\_iter(nbunch=None, data=False, default=None)

Return an iterator over the edges.

Edges are returned as tuples with optional data in the order (node, neighbor, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

**data**: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edge\_iter: iterator

An iterator of (u,v) or (u,v,d) tuples of edges.

### See also:

edges return a list of edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges_iter([0,2]))
[(0, 1), (2, 3)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

# in edges DiGraph.in\_edges (nbunch=None, data=False) Return a list of the incoming edges. See also: edges return a list of edges in\_edges\_iter DiGraph.in\_edges\_iter(nbunch=None, data=False) Return an iterator over the incoming edges. **Parameters nbunch**: iterable container, optional (default= all nodes) A container of nodes. The container will be iterated through once. data: bool, optional (default=False) If True, return edge attribute dict in 3-tuple (u,v,data). Returns in\_edge\_iter : iterator An iterator of (u,v) or (u,v,d) tuples of incoming edges. See also: edges\_iter return an iterator of edges get\_edge\_data DiGraph.get\_edge\_data(u, v, default=None) Return the attribute dictionary associated with edge (u,v). Parameters u.v: nodes default: any Python object (default=None) Value to return if the edge (u,v) is not found. Returns edge\_dict : dictionary The edge attribute dictionary. **Notes** It is faster to use G[u][v]. >>> G = nx.Graph()# or DiGraph, MultiGraph, MultiDiGraph, etc >>> G.add\_path([0,1,2,3])

Warning: Assigning G[u][v] corrupts the graph data structure. But it is safe to assign attributes to that dictionary,

>>> G[0][1]

{ }

```
>>> G[0][1]['weight'] = 7
>>> G[0][1]['weight']
7
>>> G[1][0]['weight']
7
```

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.get_edge_data(0,1) # default edge data is {}
{}
>>> e = (0,1)
>>> G.get_edge_data(*e) # tuple form
{}
>>> G.get_edge_data('a','b',default=0) # edge not in graph, return 0
0
```

#### neighbors

```
DiGraph.neighbors(n)
```

Return a list of successor nodes of n.

neighbors() and successors() are the same function.

### neighbors iter

```
DiGraph.neighbors_iter(n)
```

Return an iterator over successor nodes of n.

neighbors\_iter() and successors\_iter() are the same.

# \_\_getitem\_\_

```
DiGraph.__getitem__(n)
```

Return a dict of neighbors of node n. Use the expression 'G[n]'.

Parameters n: node

A node in the graph.

Returns adj\_dict : dictionary

The adjacency dictionary for nodes connected to n.

#### **Notes**

G[n] is similar to G.neighbors(n) but the internal data dictionary is returned instead of a list.

Assigning G[n] will corrupt the internal graph data structure. Use G[n] for reading data only.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G[0]
{1: {}}
```

#### successors

```
DiGraph.successors(n)
```

Return a list of successor nodes of n.

neighbors() and successors() are the same function.

#### successors iter

```
DiGraph.successors_iter(n)
```

Return an iterator over successor nodes of n.

neighbors\_iter() and successors\_iter() are the same.

# predecessors

```
DiGraph.predecessors(n)
```

Return a list of predecessor nodes of n.

# predecessors\_iter

```
DiGraph.predecessors_iter(n)
```

Return an iterator over predecessor nodes of n.

# adjacency\_list

```
DiGraph.adjacency_list()
```

Return an adjacency list representation of the graph.

The output adjacency list is in the order of G.nodes(). For directed graphs, only outgoing adjacencies are included.

```
Returns adj_list: lists of lists
```

The adjacency structure of the graph as a list of lists.

# See also:

```
adjacency_iter
```

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.adjacency_list() # in order given by G.nodes()
[[1], [0, 2], [1, 3], [2]]
```

#### adjacency iter

### DiGraph.adjacency\_iter()

Return an iterator of (node, adjacency dict) tuples for all nodes.

This is the fastest way to look at every edge. For directed graphs, only outgoing adjacencies are included.

## Returns adj\_iter: iterator

An iterator of (node, adjacency dictionary) for all nodes in the graph.

#### See also:

```
adjacency_list
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [(n,nbrdict) for n,nbrdict in G.adjacency_iter()]
[(0, {1: {}}), (1, {0: {}, 2: {}}), (2, {1: {}, 3: {}}), (3, {2: {}})]
```

### nbunch iter

# DiGraph.nbunch\_iter(nbunch=None)

Return an iterator of nodes contained in nbunch that are also in the graph.

The nodes in nbunch are checked for membership in the graph and if not are silently ignored.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

Returns niter: iterator

An iterator over nodes in nbunch that are also in the graph. If nbunch is None, iterate over all nodes in the graph.

# Raises NetworkXError

If nbunch is not a node or or sequence of nodes. If a node in nbunch is not hashable.

#### See also:

```
Graph.__iter__
```

#### **Notes**

When nbunch is an iterator, the returned iterator yields values directly from nbunch, becoming exhausted when nbunch is exhausted.

To test whether nbunch is a single node, one can use "if nbunch in self:", even after processing with this routine.

If nbunch is not a node or a (possibly empty) sequence/iterator or None, a NetworkXError is raised. Also, if any object in nbunch is not hashable, a NetworkXError is raised.

# Information about graph structure

DiGraph.has_node(n)	Return True if the graph contains the node n.
DiGraphcontains(n)	Return True if n is a node, False otherwise.
DiGraph.has_edge(u, v)	Return True if the edge (u,v) is in the graph.
DiGraph.order()	Return the number of nodes in the graph.
DiGraph.number_of_nodes()	Return the number of nodes in the graph.
DiGraphlen()	Return the number of nodes.
DiGraph.degree([nbunch, weight])	Return the degree of a node or nodes.
DiGraph.degree_iter([nbunch, weight])	Return an iterator for (node, degree).
DiGraph.in_degree([nbunch, weight])	Return the in-degree of a node or nodes.
DiGraph.in_degree_iter([nbunch, weight])	Return an iterator for (node, in-degree).
DiGraph.out_degree([nbunch, weight])	Return the out-degree of a node or nodes.
<pre>DiGraph.out_degree_iter([nbunch, weight])</pre>	Return an iterator for (node, out-degree).
DiGraph.size([weight])	Return the number of edges.
DiGraph.number_of_edges([u, v])	Return the number of edges between two nodes.
DiGraph.nodes_with_selfloops()	Return a list of nodes with self loops.
DiGraph.selfloop_edges([data, default])	Return a list of selfloop edges.
DiGraph.number_of_selfloops()	Return the number of selfloop edges.

# has\_node

```
DiGraph.has_node(n)
```

Return True if the graph contains the node n.

Parameters n: node

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.has_node(0)
True
```

It is more readable and simpler to use

```
>>> 0 in G
True
```

### \_\_contains\_\_

```
DiGraph.__contains__(n)
```

Return True if n is a node, False otherwise. Use the expression 'n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> 1 in G
True
```

#### has edge

```
DiGraph.has_edge (u, v)
```

Return True if the edge (u,v) is in the graph.

Parameters u,v: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

Returns edge\_ind: bool

True if edge is in the graph, False otherwise.

### **Examples**

Can be called either using two nodes u,v or edge tuple (u,v)

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.has_edge(0,1)  # using two nodes
True
>>> e = (0,1)
>>> G.has_edge(*e)  # e is a 2-tuple (u,v)
True
>>> e = (0,1,{'weight':7})
>>> G.has_edge(*e[:2])  # e is a 3-tuple (u,v,data_dictionary)
True
```

The following syntax are all equivalent:

```
>>> G.has_edge(0,1)
True
>>> 1 in G[0] # though this gives KeyError if 0 not in G
True
```

#### order

```
DiGraph.order()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

# See also:

```
number_of_nodes, __len__
```

### number of nodes

```
DiGraph.number_of_nodes()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

#### See also:

```
order, __len__
```

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> len(G)
3
```

# len

```
DiGraph.__len__()
```

Return the number of nodes. Use the expression 'len(G)'.

Returns nnodes: int

The number of nodes in the graph.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> len(G)
4
```

# degree

```
DiGraph.degree (nbunch=None, weight=None)
```

Return the degree of a node or nodes.

The node degree is the number of edges adjacent to that node.

Parameters nbunch: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

**Returns nd**: dictionary, or number

A dictionary with nodes as keys and degree as values or a number if a single node is specified.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.degree(0)
1
>>> G.degree([0,1])
{0: 1, 1: 2}
>>> list(G.degree([0,1]).values())
[1, 2]
```

#### degree iter

```
DiGraph.degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, degree).

The node degree is the number of edges adjacent to the node.

Parameters nbunch: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, degree).

# See also:

```
degree, in_degree, out_degree, in_degree_iter, out_degree_iter
```

# **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> list(G.degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.degree_iter([0,1]))
[(0, 1), (1, 2)]
```

### in\_degree

```
DiGraph.in_degree (nbunch=None, weight=None)
```

Return the in-degree of a node or nodes.

The node in-degree is the number of edges pointing in to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd: dictionary, or number

A dictionary with nodes as keys and in-degree as values or a number if a single node is specified.

#### See also:

```
degree, out_degree, in_degree_iter
```

### **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.in_degree(0)
0
>>> G.in_degree([0,1])
{0: 0, 1: 1}
>>> list(G.in_degree([0,1]).values())
[0, 1]
```

### in degree iter

```
DiGraph.in_degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, in-degree).

The node in-degree is the number of edges pointing in to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight: string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd iter: an iterator

The iterator returns two-tuples of (node, in-degree).

# See also:

```
degree, in_degree, out_degree_iter
```

# **Examples**

```
>>> G = nx.DiGraph()
>>> G.add_path([0,1,2,3])
>>> list(G.in_degree_iter(0)) # node 0 with degree 0
[(0, 0)]
>>> list(G.in_degree_iter([0,1]))
[(0, 0), (1, 1)]
```

#### out degree

```
DiGraph.out_degree (nbunch=None, weight=None)
```

Return the out-degree of a node or nodes.

The node out-degree is the number of edges pointing out of the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

**Returns nd**: dictionary, or number

A dictionary with nodes as keys and out-degree as values or a number if a single node is specified.

#### **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.out_degree(0)
1
>>> G.out_degree([0,1])
{0: 1, 1: 1}
>>> list(G.out_degree([0,1]).values())
[1, 1]
```

# out degree iter

```
DiGraph.out_degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, out-degree).

The node out-degree is the number of edges pointing out of the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, out-degree).

### See also:

```
degree, in_degree, out_degree, in_degree_iter
```

```
>>> G = nx.DiGraph()
>>> G.add_path([0,1,2,3])
>>> list(G.out_degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.out_degree_iter([0,1]))
[(0, 1), (1, 1)]
```

#### size

DiGraph.size(weight=None)

Return the number of edges.

Parameters weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

Returns nedges: int

The number of edges or sum of edge weights in the graph.

#### See also:

```
number_of_edges
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.size()
3

>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=2)
>>> G.add_edge('b','c',weight=4)
>>> G.size()
2
>>> G.size(weight='weight')
6.0
```

#### number of edges

```
DiGraph.number_of_edges(u=None, v=None)
```

Return the number of edges between two nodes.

**Parameters u,v**: nodes, optional (default=all edges)

If u and v are specified, return the number of edges between u and v. Otherwise return the total number of all edges.

Returns nedges: int

The number of edges in the graph. If nodes u and v are specified return the number of edges between those nodes.

#### See also:

size

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.number_of_edges()
3
>>> G.number_of_edges(0,1)
1
>>> e = (0,1)
>>> G.number_of_edges(*e)
```

# nodes\_with\_selfloops

```
DiGraph.nodes_with_selfloops()
```

Return a list of nodes with self loops.

A node with a self loop has an edge with both ends adjacent to that node.

Returns nodelist: list

A list of nodes with self loops.

#### See also:

```
selfloop_edges, number_of_selfloops
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.nodes_with_selfloops()
[1]
```

### selfloop\_edges

```
DiGraph.selfloop_edges(data=False, default=None)
```

Return a list of selfloop edges.

A selfloop edge has the same node at both ends.

Parameters data: string or bool, optional (default=False)

Return selfloop edges as two tuples (u,v) (data=False) or three-tuples (u,v,datadict) (data=True) or three-tuples (u,v,datavalue) (data='attrname')

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edgelist: list of edge tuples

A list of all selfloop edges.

### See also:

```
nodes_with_selfloops, number_of_selfloops
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.selfloop_edges()
[(1, 1)]
>>> G.selfloop_edges(data=True)
[(1, 1, {})]
```

# number\_of\_selfloops

```
DiGraph.number_of_selfloops()
```

Return the number of selfloop edges.

A selfloop edge has the same node at both ends.

**Returns nloops**: int

The number of selfloops.

#### See also:

```
nodes_with_selfloops, selfloop_edges
```

# **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.number_of_selfloops()
1
```

# Making copies and subgraphs

DiGraph.copy()	Return a copy of the graph.
DiGraph.to_undirected([reciprocal])	Return an undirected representation of the digraph.
DiGraph.to_directed()	Return a directed copy of the graph.
DiGraph.subgraph(nbunch)	Return the subgraph induced on nodes in nbunch.
DiGraph.reverse([copy])	Return the reverse of the graph.

# сору

```
DiGraph.copy()
```

Return a copy of the graph.

Returns G: Graph

A copy of the graph.

#### See also:

to\_directed return a directed copy of the graph.

#### **Notes**

This makes a complete copy of the graph including all of the node or edge attributes.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.copy()
```

# to undirected

```
DiGraph.to_undirected(reciprocal=False)
```

Return an undirected representation of the digraph.

Parameters reciprocal: bool (optional)

If True only keep edges that appear in both directions in the original digraph.

Returns G: Graph

An undirected graph with the same name and nodes and with edge (u,v,data) if either (u,v,data) or (v,u,data) is in the digraph. If both edges exist in digraph and their edge data is different, only one edge is created with an arbitrary choice of which edge data to use. You must check and correct for this manually if desired.

#### **Notes**

If edges in both directions (u,v) and (v,u) exist in the graph, attributes for the new undirected edge will be a combination of the attributes of the directed edges. The edge data is updated in the (arbitrary) order that the edges are encountered. For more customized control of the edge attributes use add\_edge().

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar G=DiGraph(D) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

#### to directed

```
DiGraph.to directed()
```

Return a directed copy of the graph.

Returns G: DiGraph

A deepcopy of the graph.

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar D=DiGraph(G) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

### **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]

If already directed, return a (deep) copy
>>> G = nx.DiGraph() # or MultiDiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
```

### subgraph

```
DiGraph.subgraph(nbunch)
```

>>> H.edges()
[(0, 1)]

Return the subgraph induced on nodes in nbunch.

The induced subgraph of the graph contains the nodes in nbunch and the edges between those nodes.

**Parameters nbunch**: list, iterable

A container of nodes which will be iterated through once.

Returns G: Graph

A subgraph of the graph with the same edge attributes.

### **Notes**

The graph, edge or node attributes just point to the original graph. So changes to the node or edge structure will not be reflected in the original graph while changes to the attributes will.

To create a subgraph with its own copy of the edge/node attributes use: nx.Graph(G.subgraph(nbunch))

If edge attributes are containers, a deep copy can be obtained using: G.subgraph(nbunch).copy()

For an inplace reduction of a graph to a subgraph you can remove nodes: G.remove\_nodes\_from([ n in G if n not in set(nbunch)])

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.subgraph([0,1,2])
>>> H.edges()
[(0, 1), (1, 2)]
```

#### reverse

```
DiGraph.reverse(copy=True)
```

Return the reverse of the graph.

The reverse is a graph with the same nodes and edges but with the directions of the edges reversed.

**Parameters copy**: bool optional (default=True)

If True, return a new DiGraph holding the reversed edges. If False, reverse the reverse graph is created using the original graph (this changes the original graph).

# 3.2.3 MultiGraph - Undirected graphs with self loops and parallel edges

#### Overview

```
MultiGraph (data=None, **attr)
```

An undirected graph class that can store multiedges.

Multiedges are multiple edges between two nodes. Each edge can hold optional data or attributes.

A MultiGraph holds undirected edges. Self loops are allowed.

Nodes can be arbitrary (hashable) Python objects with optional key/value attributes.

Edges are represented as links between nodes with optional key/value attributes.

Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse matrix, or a PyGraphviz graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

### See also:

```
Graph, DiGraph, MultiDiGraph
```

# **Examples**

Create a multigraph object that tracks the order nodes are added.

```
>>> from collections import OrderedDict
>>> class OrderedGraph(nx.MultiGraph):
... node_dict_factory = OrderedDict
>>> G = OrderedGraph()
```

```
>>> G.add_nodes_from( (2,1) )
>>> G.nodes()
[2, 1]
>>> G.add_edges_from( ((2,2), (2,1), (2,1), (1,1)) )
>>> G.edges()
[(2, 1), (2, 1), (2, 2), (1, 1)]
```

Create a multgraph object that tracks the order nodes are added and for each node track the order that neighbors are added and for each neighbor tracks the order that multiedges are added.

# Adding and removing nodes and edges

MultiGraphinit([data])	
MultiGraph.add_node(n[, attr_dict])	Add a single node n and update node attributes.
MultiGraph.add_nodes_from(nodes, **attr)	Add multiple nodes.
MultiGraph.remove_node(n)	Remove node n.
MultiGraph.remove_nodes_from(nodes)	Remove multiple nodes.
MultiGraph.add_edge(u, v[, key, attr_dict])	Add an edge between u and v.
MultiGraph.add_edges_from(ebunch[, attr_dict])	Add all the edges in ebunch.
MultiGraph.add_weighted_edges_from(ebunch[,])	Add all the edges in ebunch as weighted edges with specified weig
MultiGraph.remove_edge(u,v[,key])	Remove an edge between u and v.
MultiGraph.remove_edges_from(ebunch)	Remove all edges specified in ebunch.
MultiGraph.add_star(nodes, **attr)	Add a star.
MultiGraph.add_path(nodes, **attr)	Add a path.
MultiGraph.add_cycle(nodes, **attr)	Add a cycle.
MultiGraph.clear()	Remove all nodes and edges from the graph.

```
__init__
MultiGraph.__init__ (data=None, **attr)

add_node

MultiGraph.add_node (n, attr_dict=None, **attr)
    Add a single node n and update node attributes.

Parameters n: node
```

A node can be any hashable Python object except None.

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of node attributes. Key/value pairs will update existing data associated with the node.

attr: keyword arguments, optional

Set or change attributes using key=value.

#### See also:

```
add_nodes_from
```

#### **Notes**

A hashable object is one that can be used as a key in a Python dictionary. This includes strings, numbers, tuples of strings and numbers, etc.

On many platforms hashable items also include mutables such as NetworkX Graphs, though one should be careful that the hash doesn't change on mutables.

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_node(1)
>>> G.add_node('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_node(K3)
>>> G.number_of_nodes()
3
```

Use keywords set/change node attributes:

```
>>> G.add_node(1,size=10)
>>> G.add_node(3,weight=0.4,UTM=('13S',382871,3972649))
```

#### add nodes from

```
MultiGraph.add_nodes_from (nodes, **attr)
Add multiple nodes.
```

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). OR A container of (node, attribute dict) tuples. Node attributes are updated using the attribute dict.

attr : keyword arguments, optional (default= no attributes)

Update attributes for all nodes in nodes. Node attributes specified in nodes as a tuple take precedence over attributes specified generally.

## See also:

```
add_node
```

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_nodes_from('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_nodes_from(K3)
>>> sorted(G.nodes(),key=str)
[0, 1, 2, 'H', 'e', 'l', 'o']

Use keywords to update specific node attributes for every node.
>>> G.add_nodes_from([1,2], size=10)
>>> G.add_nodes_from([3,4], weight=0.4)

Use (node, attrdict) tuples to update attributes for specific nodes.
>>> G.add_nodes_from([(1,dict(size=11)), (2,{'color':'blue'})])
>>> G.node[1]['size']
11
>>> H = nx.Graph()
```

## remove\_node

11

```
MultiGraph.remove_node(n)
```

>>> H.node[1]['size']

Remove node n.

Removes the node n and all adjacent edges. Attempting to remove a non-existent node will raise an exception.

Parameters n: node

A node in the graph

>>> H.add\_nodes\_from(G.nodes(data=True))

## Raises NetworkXError

If n is not in the graph.

## See also:

```
remove_nodes_from
```

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.edges()
[(0, 1), (1, 2)]
>>> G.remove_node(1)
>>> G.edges()
[]
```

## remove nodes from

```
MultiGraph.remove_nodes_from(nodes)
```

Remove multiple nodes.

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). If a node in the container is not in the graph it is silently ignored.

#### See also:

```
remove node
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> e = G.nodes()
>>> e
[0, 1, 2]
>>> G.remove_nodes_from(e)
>>> G.nodes()
[]
```

## add\_edge

```
MultiGraph.add_edge (u, v, key=None, attr_dict=None, **attr)
Add an edge between u and v.
```

The nodes u and v will be automatically added if they are not already in the graph.

Edge attributes can be specified with keywords or by providing a dictionary with key/value pairs. See examples below.

#### Parameters u,v: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

key: hashable identifier, optional (default=lowest unused integer)

Used to distinguish multiedges between a pair of nodes.

```
attr_dict : dictionary, optional (default= no attributes)
```

Dictionary of edge attributes. Key/value pairs will update existing data associated with the edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

#### See also:

```
add_edges_from add a collection of edges
```

#### **Notes**

To replace/update edge data, use the optional key argument to identify a unique edge. Otherwise a new edge will be created.

NetworkX algorithms designed for weighted graphs cannot use multigraphs directly because it is not clear how to handle multiedge weights. Convert to Graph using edge attribute 'weight' to enable weighted graph algorithms.

The following all add the edge e=(1,2) to graph G:

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> e = (1,2)
>>> G.add_edge(1, 2) # explicit two-node form
>>> G.add_edge(*e) # single edge as tuple of two nodes
>>> G.add_edges_from([(1,2)]) # add edges from iterable container
```

Associate data to edges using keywords:

```
>>> G.add_edge(1, 2, weight=3)
>>> G.add_edge(1, 2, key=0, weight=4) # update data for key=0
>>> G.add_edge(1, 3, weight=7, capacity=15, length=342.7)
```

#### add edges from

```
MultiGraph.add_edges_from(ebunch, attr_dict=None, **attr)
Add all the edges in ebunch.
```

Parameters ebunch: container of edges

Each edge given in the container will be added to the graph. The edges can be:

- 2-tuples (u,v) or
- 3-tuples (u,v,d) for an edge attribute dict d, or
- 4-tuples (u,v,k,d) for an edge identified by key k

attr dict: dictionary, optional (default= no attributes)

Dictionary of edge attributes. Key/value pairs will update existing data associated with each edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

### See also:

```
add_edge add a single edge
add_weighted_edges_from convenient way to add weighted edges
```

#### **Notes**

Adding the same edge twice has no effect but any edge data will be updated when each duplicate edge is added. Edge attributes specified in edges take precedence over attributes specified generally.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edges_from([(0,1),(1,2)]) # using a list of edge tuples
>>> e = zip(range(0,3),range(1,4))
>>> G.add_edges_from(e) # Add the path graph 0-1-2-3
```

### Associate data to edges

```
>>> G.add_edges_from([(1,2),(2,3)], weight=3)
>>> G.add_edges_from([(3,4),(1,4)], label='WN2898')
```

## add\_weighted\_edges\_from

MultiGraph.add\_weighted\_edges\_from(ebunch, weight='weight', \*\*attr)

Add all the edges in ebunch as weighted edges with specified weights.

## Parameters ebunch: container of edges

Each edge given in the list or container will be added to the graph. The edges must be given as 3-tuples (u,v,w) where w is a number.

weight: string, optional (default= 'weight')

The attribute name for the edge weights to be added.

attr: keyword arguments, optional (default= no attributes)

Edge attributes to add/update for all edges.

#### See also:

```
add_edge add a single edge
add_edges_from add multiple edges
```

#### **Notes**

Adding the same edge twice for Graph/DiGraph simply updates the edge data. For MultiGraph/MultiDiGraph, duplicate edges are stored.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_weighted_edges_from([(0,1,3.0),(1,2,7.5)])
```

## remove\_edge

```
MultiGraph.remove_edge(u, v, key=None)
```

Remove an edge between u and v.

### Parameters u,v: nodes

Remove an edge between nodes u and v.

key: hashable identifier, optional (default=None)

Used to distinguish multiple edges between a pair of nodes. If None remove a single (abritrary) edge between u and v.

## Raises NetworkXError

If there is not an edge between u and v, or if there is no edge with the specified key.

See also:

remove\_edges\_from remove a collection of edges

## **Examples**

```
>>> G = nx.MultiGraph()
>>> G.add_path([0,1,2,3])
>>> G.remove_edge(0,1)
>>> e = (1,2)
>>> G.remove_edge(*e) # unpacks e from an edge tuple
For multiple edges
                         # or MultiDiGraph, etc
>>> G = nx.MultiGraph()
>>> G.add_edges_from([(1,2),(1,2),(1,2)])
>>> G.remove_edge(1,2) # remove a single (arbitrary) edge
For edges with keys
>>> G = nx.MultiGraph()
                         # or MultiDiGraph, etc
>>> G.add_edge(1,2,key='first')
>>> G.add_edge(1,2,key='second')
>>> G.remove_edge(1,2,key='second')
```

#### remove edges from

```
MultiGraph.remove_edges_from(ebunch)
```

Remove all edges specified in ebunch.

## Parameters ebunch: list or container of edge tuples

Each edge given in the list or container will be removed from the graph. The edges can be:

- 2-tuples (u,v) All edges between u and v are removed.
- 3-tuples (u,v,key) The edge identified by key is removed.
- 4-tuples (u,v,key,data) where data is ignored.

## See also:

remove\_edge remove a single edge

### **Notes**

Will fail silently if an edge in ebunch is not in the graph.

## **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> ebunch=[(1,2),(2,3)]
>>> G.remove_edges_from(ebunch)
```

Removing multiple copies of edges

```
>>> G = nx.MultiGraph()
>>> G.add_edges_from([(1,2),(1,2)])
>>> G.remove_edges_from([(1,2),(1,2)])
>>> G.edges()
[(1, 2)]
>>> G.remove_edges_from([(1,2),(1,2)]) # silently ignore extra copy
>>> G.edges() # now empty graph
[]
```

## add\_star

```
MultiGraph.add_star (nodes, **attr)
```

Add a star.

The first node in nodes is the middle of the star. It is connected to all other nodes.

Parameters nodes: iterable container

A container of nodes.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to every edge in star.

## See also:

```
add_path, add_cycle
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_star([0,1,2,3])
>>> G.add_star([10,11,12],weight=2)
```

## add\_path

```
MultiGraph.add_path (nodes, **attr)
Add a path.
```

Parameters nodes: iterable container

A container of nodes. A path will be constructed from the nodes (in order) and added to the graph.

**attr**: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in path.

## See also:

```
add_star,add_cycle
```

#### **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.add_path([10,11,12], weight=7)
```

## add\_cycle

```
MultiGraph.add_cycle (nodes, **attr)
Add a cycle.
```

## Parameters nodes: iterable container

A container of nodes. A cycle will be constructed from the nodes (in order) and added to the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in cycle.

## See also:

```
add_path, add_star
```

## **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_cycle([0,1,2,3])
>>> G.add_cycle([10,11,12], weight=7)
```

#### clear

```
MultiGraph.clear()
```

Remove all nodes and edges from the graph.

This also removes the name, and all graph, node, and edge attributes.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.clear()
>>> G.nodes()
[]
>>> G.edges()
[]
```

## Iterating over nodes and edges

MultiGraph.nodes([data])	Return a list of the nodes in the graph.
MultiGraph.nodes_iter([data])	Return an iterator over the nodes.
MultiGraphiter()	Iterate over the nodes.
MultiGraph.edges([nbunch, data, keys, default])	Return a list of edges.
MultiGraph.edges_iter([nbunch, data, keys,])	Return an iterator over the edges.
MultiGraph.get_edge_data(u, v[, key, default])	Return the attribute dictionary associated with edge (u,v).
MultiGraph.neighbors(n)	Return a list of the nodes connected to the node n.
MultiGraph.neighbors_iter(n)	Return an iterator over all neighbors of node n.
	Continued on next page

Table 3.10 – continued from previous page

MultiGraphgetitem(n)	Return a dict of neighbors of node n.
MultiGraph.adjacency_list()	Return an adjacency list representation of the graph.
MultiGraph.adjacency_iter()	Return an iterator of (node, adjacency dict) tuples for all nodes.
MultiGraph.nbunch_iter([nbunch])	Return an iterator of nodes contained in nbunch that are also in the graph.

#### nodes

MultiGraph.nodes (data=False)

Return a list of the nodes in the graph.

Parameters data: boolean, optional (default=False)

If False return a list of nodes. If True return a two-tuple of node and node data dictionary

Returns nlist: list

A list of nodes. If data=True a list of two-tuples containing (node, node data dictionary).

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.nodes()
[0, 1, 2]
>>> G.add_node(1, time='5pm')
>>> G.nodes(data=True)
[(0, {}), (1, {'time': '5pm'}), (2, {})]
```

## nodes iter

MultiGraph.nodes\_iter(data=False)

Return an iterator over the nodes.

Parameters data: boolean, optional (default=False)

If False the iterator returns nodes. If True return a two-tuple of node and node data dictionary

Returns niter: iterator

An iterator over nodes. If data=True the iterator gives two-tuples containing (node, node data, dictionary)

### **Notes**

If the node data is not required it is simpler and equivalent to use the expression 'for n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
```

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])

>>> [d for n,d in G.nodes_iter(data=True)]
[{}, {}, {}]

__iter__

MultiGraph.__iter__()
Iterate over the nodes. Use the expression 'for n in G'.
```

Returns niter: iterator

An iterator over all nodes in the graph.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
```

## edges

MultiGraph.edges (nbunch=None, data=False, keys=False, default=None)
Return a list of edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,data) (True).

**keys**: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,key) (True).

**Returns** edge\_list: list of edge tuples

Edges that are adjacent to any node in nbunch, or a list of all edges if nbunch is not specified.

## See also:

edges\_iter return an iterator over the edges

### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

```
>>> G = nx.MultiGraph()
                         # or MultiDiGraph
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> G.edges()
[(0, 1), (1, 2), (2, 3)]
>>> G.edges(data=True) # default edge data is {} (empty dictionary)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> G.edges(keys=True) # default keys are integers
[(0, 1, 0), (1, 2, 0), (2, 3, 0)]
>>> G.edges(data=True, keys=True) # default keys are integers
[(0, 1, 0, {}), (1, 2, 0, {}), (2, 3, 0, {'weight': 5})]
>>> list(G.edges(data='weight',default=1,keys=True))
[(0, 1, 0, 1), (1, 2, 0, 1), (2, 3, 0, 5)]
>>> G.edges([0,3])
[(0, 1), (3, 2)]
>>> G.edges(0)
[(0, 1)]
```

### edges\_iter

MultiGraph.edges\_iter (nbunch=None, data=False, keys=False, default=None)
Return an iterator over the edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

**data**: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

keys: bool, optional (default=False)

If True, return edge keys with each edge.

Returns edge iter: iterator

An iterator of (u,v), (u,v,d) or (u,v,key,d) tuples of edges.

## See also:

edges return a list of edges

### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

```
>>> G = nx.MultiGraph()
                          # or MultiDiGraph
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges(keys=True)) # default keys are integers
[(0, 1, 0), (1, 2, 0), (2, 3, 0)]
>>> list(G.edges(data=True, keys=True))  # default keys are integers
[(0, 1, 0, {}), (1, 2, 0, {}), (2, 3, 0, {'weight': 5})]
>>> list(G.edges(data='weight',default=1,keys=True))
[(0, 1, 0, 1), (1, 2, 0, 1), (2, 3, 0, 5)]
>>> list(G.edges_iter([0,3]))
[(0, 1), (3, 2)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

#### get edge data

MultiGraph.get\_edge\_data (u, v, key=None, default=None)Return the attribute dictionary associated with edge (u,v).

Parameters u,v: nodes

## default: any Python object (default=None)

Value to return if the edge (u,v) is not found.

key: hashable identifier, optional (default=None)

Return data only for the edge with specified key.

## Returns edge\_dict: dictionary

The edge attribute dictionary.

#### **Notes**

It is faster to use G[u][v][key].

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_edge(0,1,key='a',weight=7)
>>> G[0][1]['a'] # key='a'
{'weight': 7}
```

Warning: Assigning G[u][v][key] corrupts the graph data structure. But it is safe to assign attributes to that dictionary,

```
>>> G[0][1]['a']['weight'] = 10
>>> G[0][1]['a']['weight']
10
>>> G[1][0]['a']['weight']
10
```

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.get_edge_data(0,1)
{0: {}}
>>> e = (0,1)
>>> G.get_edge_data(*e) # tuple form
{0: {}}
>>> G.get_edge_data('a','b',default=0) # edge not in graph, return 0
0
```

#### neighbors

MultiGraph.neighbors(n)

Return a list of the nodes connected to the node n.

Parameters n : node

A node in the graph

Returns nlist: list

A list of nodes that are adjacent to n.

Raises NetworkXError

If the node n is not in the graph.

### **Notes**

It is usually more convenient (and faster) to access the adjacency dictionary as G[n]:

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=7)
>>> G['a']
{'b': {'weight': 7}}
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.neighbors(0)
[1]
```

## neighbors iter

```
MultiGraph.neighbors_iter(n)
```

Return an iterator over all neighbors of node n.

#### **Notes**

It is faster to use the idiom "in G[0]", e.g.

```
>>> G = nx.path_graph(4)
>>> [n for n in G[0]]
[1]
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [n for n in G.neighbors_iter(0)]
[1]
```

## \_\_getitem\_\_

```
MultiGraph.__getitem__(n)
```

Return a dict of neighbors of node n. Use the expression 'G[n]'.

Parameters n: node

A node in the graph.

Returns adj\_dict: dictionary

The adjacency dictionary for nodes connected to n.

## **Notes**

G[n] is similar to G.neighbors(n) but the internal data dictionary is returned instead of a list.

Assigning G[n] will corrupt the internal graph data structure. Use G[n] for reading data only.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G[0]
{1: {}}
```

## adjacency\_list

```
MultiGraph.adjacency_list()
```

Return an adjacency list representation of the graph.

The output adjacency list is in the order of G.nodes(). For directed graphs, only outgoing adjacencies are included.

Returns adj\_list : lists of lists

The adjacency structure of the graph as a list of lists.

## See also:

```
adjacency_iter
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.adjacency_list() # in order given by G.nodes()
[[1], [0, 2], [1, 3], [2]]
```

### adjacency iter

```
MultiGraph.adjacency_iter()
```

Return an iterator of (node, adjacency dict) tuples for all nodes.

This is the fastest way to look at every edge. For directed graphs, only outgoing adjacencies are included.

## Returns adj\_iter: iterator

An iterator of (node, adjacency dictionary) for all nodes in the graph.

#### See also:

```
adjacency list
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [(n,nbrdict) for n,nbrdict in G.adjacency_iter()]
[(0, {1: {}}), (1, {0: {}, 2: {}}), (2, {1: {}, 3: {}}), (3, {2: {}})]
```

#### nbunch iter

```
MultiGraph.nbunch_iter(nbunch=None)
```

Return an iterator of nodes contained in nbunch that are also in the graph.

The nodes in nbunch are checked for membership in the graph and if not are silently ignored.

Parameters nbunch: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

Returns niter: iterator

An iterator over nodes in nbunch that are also in the graph. If nbunch is None, iterate over all nodes in the graph.

## Raises NetworkXError

If nbunch is not a node or or sequence of nodes. If a node in nbunch is not hashable.

### See also:

```
Graph.__iter__
```

#### **Notes**

When nbunch is an iterator, the returned iterator yields values directly from nbunch, becoming exhausted when nbunch is exhausted.

To test whether nbunch is a single node, one can use "if nbunch in self:", even after processing with this routine.

If nbunch is not a node or a (possibly empty) sequence/iterator or None, a NetworkXError is raised. Also, if any object in nbunch is not hashable, a NetworkXError is raised.

## Information about graph structure

MultiGraph.has_node(n)	Return True if the graph contains the node n.
MultiGraphcontains(n)	Return True if n is a node, False otherwise.
MultiGraph.has_edge(u,v[,key])	Return True if the graph has an edge between nodes u and v.
MultiGraph.order()	Return the number of nodes in the graph.
MultiGraph.number_of_nodes()	Return the number of nodes in the graph.
MultiGraphlen()	Return the number of nodes.
MultiGraph.degree([nbunch, weight])	Return the degree of a node or nodes.
MultiGraph.degree_iter([nbunch, weight])	Return an iterator for (node, degree).
MultiGraph.size([weight])	Return the number of edges.
MultiGraph.number_of_edges([u, v])	Return the number of edges between two nodes.
MultiGraph.nodes_with_selfloops()	Return a list of nodes with self loops.
MultiGraph.selfloop_edges([data, keys, default])	Return a list of selfloop edges.
MultiGraph.number_of_selfloops()	Return the number of selfloop edges.

## has\_node

```
MultiGraph.has_node(n)
```

Return True if the graph contains the node n.

Parameters n: node

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.has_node(0)
True
```

It is more readable and simpler to use

```
>>> 0 in G
True
```

## \_\_contains\_\_

```
MultiGraph.__contains__(n)
```

Return True if n is a node, False otherwise. Use the expression 'n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> 1 in G
True
```

#### has edge

```
MultiGraph.has_edge (u, v, key=None)
```

Return True if the graph has an edge between nodes u and v.

Parameters u,v: nodes

Nodes can be, for example, strings or numbers.

**key**: hashable identifier, optional (default=None)

If specified return True only if the edge with key is found.

Returns edge ind: bool

True if edge is in the graph, False otherwise.

## **Examples**

Can be called either using two nodes u,v, an edge tuple (u,v), or an edge tuple (u,v,key).

```
>>> G = nx.MultiGraph()  # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.has_edge(0,1)  # using two nodes
True
>>> e = (0,1)
>>> G.has_edge(*e)  # e is a 2-tuple (u,v)
True
>>> G.add_edge(0,1,key='a')
>>> G.has_edge(0,1,key='a')  # specify key
True
>>> e=(0,1,'a')
>>> G.has_edge(*e)  # e is a 3-tuple (u,v,'a')
True
```

### The following syntax are equivalent:

```
>>> G.has_edge(0,1)
True
>>> 1 in G[0] # though this gives KeyError if 0 not in G
True
```

## order

```
MultiGraph.order()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

### See also:

```
number_of_nodes, __len__
```

## number of nodes

```
MultiGraph.number_of_nodes()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

### See also:

```
order, __len__
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> len(G)
3
```

## \_\_len\_\_

```
MultiGraph.__len__()
```

Return the number of nodes. Use the expression 'len(G)'.

Returns nnodes: int

The number of nodes in the graph.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> len(G)
4
```

### degree

```
MultiGraph.degree (nbunch=None, weight=None)
```

Return the degree of a node or nodes.

The node degree is the number of edges adjacent to that node.

Parameters nbunch: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

## Returns nd: dictionary, or number

A dictionary with nodes as keys and degree as values or a number if a single node is specified.

#### **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.degree(0)
1
>>> G.degree([0,1])
{0: 1, 1: 2}
>>> list(G.degree([0,1]).values())
[1, 2]
```

## degree iter

```
MultiGraph.degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, degree).

The node degree is the number of edges adjacent to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight: string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, degree).

### See also:

degree

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> list(G.degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.degree_iter([0,1]))
[(0, 1), (1, 2)]
```

#### size

```
MultiGraph.size(weight=None)
```

Return the number of edges.

**Parameters weight**: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

## Returns nedges: int

The number of edges or sum of edge weights in the graph.

#### See also:

```
number_of_edges
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.size()
3
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=2)
>>> G.add_edge('b','c',weight=4)
>>> G.size()
2
>>> G.size(weight='weight')
6.0
```

## number of edges

```
MultiGraph.number_of_edges(u=None, v=None)
```

Return the number of edges between two nodes.

**Parameters u,v**: nodes, optional (default=all edges)

If u and v are specified, return the number of edges between u and v. Otherwise return the total number of all edges.

## Returns nedges: int

The number of edges in the graph. If nodes u and v are specified return the number of edges between those nodes.

## See also:

size

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.number_of_edges()
3
>>> G.number_of_edges(0,1)
1
>>> e = (0,1)
>>> G.number_of_edges(*e)
```

## nodes\_with\_selfloops

```
MultiGraph.nodes_with_selfloops()
```

Return a list of nodes with self loops.

A node with a self loop has an edge with both ends adjacent to that node.

Returns nodelist: list

A list of nodes with self loops.

#### See also:

```
selfloop_edges, number_of_selfloops
```

#### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.nodes_with_selfloops()
[1]
```

## selfloop\_edges

MultiGraph.selfloop\_edges(data=False, keys=False, default=None)

Return a list of selfloop edges.

A selfloop edge has the same node at both ends.

Parameters data: bool, optional (default=False)

Return selfloop edges as two tuples (u,v) (data=False) or three-tuples (u,v,datadict) (data=True) or three-tuples (u,v,datavalue) (data='attrname')

default : value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

**keys**: bool, optional (default=False)

If True, return edge keys with each edge.

**Returns edgelist**: list of edge tuples

A list of all selfloop edges.

#### See also:

```
nodes_with_selfloops, number_of_selfloops
```

# **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.selfloop_edges()
[(1, 1)]
```

```
>>> G.selfloop_edges(data=True)
[(1, 1, {})]
>>> G.selfloop_edges(keys=True)
[(1, 1, 0)]
>>> G.selfloop_edges(keys=True, data=True)
[(1, 1, 0, {})]
```

## number of selfloops

```
MultiGraph.number_of_selfloops()
```

Return the number of selfloop edges.

A selfloop edge has the same node at both ends.

Returns nloops: int

The number of selfloops.

## See also:

```
nodes_with_selfloops, selfloop_edges
```

## **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.number_of_selfloops()
1
```

## Making copies and subgraphs

MultiGraph.copy()	Return a copy of the graph.
MultiGraph.to_undirected()	Return an undirected copy of the graph.
MultiGraph.to_directed()	Return a directed representation of the graph.
MultiGraph.subgraph(nbunch)	Return the subgraph induced on nodes in nbunch.

## сору

```
MultiGraph.copy()
```

Return a copy of the graph.

**Returns G**: Graph

A copy of the graph.

## See also:

to\_directed return a directed copy of the graph.

## **Notes**

This makes a complete copy of the graph including all of the node or edge attributes.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.copy()
```

## to\_undirected

```
MultiGraph.to_undirected()
```

Return an undirected copy of the graph.

Returns G: Graph/MultiGraph

A deepcopy of the graph.

## See also:

```
copy, add_edge, add_edges_from
```

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar G=DiGraph(D) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

## **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]
>>> G2 = H.to_undirected()
>>> G2.edges()
[(0, 1)]
```

## to\_directed

```
MultiGraph.to_directed()
```

Return a directed representation of the graph.

### Returns G: MultiDiGraph

A directed graph with the same name, same nodes, and with each edge (u,v,data) replaced by two directed edges (u,v,data) and (v,u,data).

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar D=DiGraph(G) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

## **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]

If already directed, return a (deep) copy
>>> G = nx.DiGraph() # or MultiDiGraph, etc
>>> G.add_path([0,1])
```

# subgraph

```
MultiGraph.subgraph(nbunch)
```

>>> H = G.to\_directed()

>>> H.edges()
[(0, 1)]

Return the subgraph induced on nodes in nbunch.

The induced subgraph of the graph contains the nodes in nbunch and the edges between those nodes.

**Parameters nbunch**: list, iterable

A container of nodes which will be iterated through once.

Returns G: Graph

A subgraph of the graph with the same edge attributes.

### **Notes**

The graph, edge or node attributes just point to the original graph. So changes to the node or edge structure will not be reflected in the original graph while changes to the attributes will.

To create a subgraph with its own copy of the edge/node attributes use: nx.Graph(G.subgraph(nbunch))

If edge attributes are containers, a deep copy can be obtained using: G.subgraph(nbunch).copy()

For an inplace reduction of a graph to a subgraph you can remove nodes: G.remove\_nodes\_from([ n in G if n not in set(nbunch)])

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.subgraph([0,1,2])
>>> H.edges()
[(0, 1), (1, 2)]
```

# 3.2.4 MultiDiGraph - Directed graphs with self loops and parallel edges

#### Overview

```
MultiDiGraph (data=None, **attr)
```

A directed graph class that can store multiedges.

Multiedges are multiple edges between two nodes. Each edge can hold optional data or attributes.

A MultiDiGraph holds directed edges. Self loops are allowed.

Nodes can be arbitrary (hashable) Python objects with optional key/value attributes.

Edges are represented as links between nodes with optional key/value attributes.

## Parameters data: input graph

Data to initialize graph. If data=None (default) an empty graph is created. The data can be an edge list, or any NetworkX graph object. If the corresponding optional Python packages are installed the data can also be a NumPy matrix or 2d ndarray, a SciPy sparse matrix, or a PyGraphviz graph.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to graph as key=value pairs.

## See also:

```
Graph, DiGraph, MultiGraph
```

## **Examples**

Create a multigraph object that tracks the order nodes are added.

```
>>> from collections import OrderedDict
>>> class OrderedGraph(nx.MultiDiGraph):
... node_dict_factory = OrderedDict
>>> G = OrderedGraph()
>>> G.add_nodes_from((2,1))
>>> G.nodes()
[2, 1]
>>> G.add_edges_from(((2,2), (2,1), (2,1), (1,1)))
>>> G.edges()
[(2, 1), (2, 1), (2, 2), (1, 1)]
```

Create a multdigraph object that tracks the order nodes are added and for each node track the order that neighbors are added and for each neighbor tracks the order that multiedges are added.

## **Adding and Removing Nodes and Edges**

MultiDiGraphinit([data])	
MultiDiGraph.add_node(n[, attr_dict])	Add a single node n and update node attributes.
MultiDiGraph.add_nodes_from(nodes, **attr)	Add multiple nodes.
MultiDiGraph.remove_node(n)	Remove node n.
MultiDiGraph.remove_nodes_from(nbunch)	Remove multiple nodes.
$\texttt{MultiDiGraph.add\_edge}(u, v[, key, attr\_dict])$	Add an edge between u and v.
<pre>MultiDiGraph.add_edges_from(ebunch[, attr_dict])</pre>	Add all the edges in ebunch.
MultiDiGraph.add_weighted_edges_from(ebunch)	Add all the edges in ebunch as weighted edges with specified weight
MultiDiGraph.remove_edge(u,v[,key])	Remove an edge between u and v.
MultiDiGraph.remove_edges_from(ebunch)	Remove all edges specified in ebunch.
MultiDiGraph.add_star(nodes, **attr)	Add a star.
MultiDiGraph.add_path(nodes, **attr)	Add a path.
MultiDiGraph.add_cycle(nodes, **attr)	Add a cycle.
MultiDiGraph.clear()	Remove all nodes and edges from the graph.

## \_\_init\_\_

MultiDiGraph. init (data=None, \*\*attr)

## add\_node

MultiDiGraph.add\_node (n, attr\_dict=None, \*\*attr)

Add a single node n and update node attributes.

## Parameters n: node

A node can be any hashable Python object except None.

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of node attributes. Key/value pairs will update existing data associated with the node.

attr: keyword arguments, optional

Set or change attributes using key=value.

## See also:

 $add\_nodes\_from$ 

#### **Notes**

A hashable object is one that can be used as a key in a Python dictionary. This includes strings, numbers, tuples of strings and numbers, etc.

On many platforms hashable items also include mutables such as NetworkX Graphs, though one should be careful that the hash doesn't change on mutables.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_node(1)
>>> G.add_node('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_node(K3)
>>> G.number_of_nodes()
3
```

Use keywords set/change node attributes:

```
>>> G.add_node(1,size=10)
>>> G.add_node(3,weight=0.4,UTM=('13S',382871,3972649))
```

#### add nodes from

```
MultiDiGraph.add_nodes_from(nodes, **attr)
Add multiple nodes.
```

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). OR A container of (node, attribute dict) tuples. Node attributes are updated using the attribute dict.

attr: keyword arguments, optional (default= no attributes)

Update attributes for all nodes in nodes. Node attributes specified in nodes as a tuple take precedence over attributes specified generally.

## See also:

```
add node
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_nodes_from('Hello')
>>> K3 = nx.Graph([(0,1),(1,2),(2,0)])
>>> G.add_nodes_from(K3)
>>> sorted(G.nodes(),key=str)
[0, 1, 2, 'H', 'e', 'l', 'o']
```

Use keywords to update specific node attributes for every node.

```
>>> G.add_nodes_from([1,2], size=10)
>>> G.add_nodes_from([3,4], weight=0.4)
```

Use (node, attrdict) tuples to update attributes for specific nodes.

```
>>> G.add_nodes_from([(1,dict(size=11)), (2,{'color':'blue'})])
>>> G.node[1]['size']
11
>>> H = nx.Graph()
>>> H.add_nodes_from(G.nodes(data=True))
>>> H.node[1]['size']
11
```

## remove\_node

```
MultiDiGraph.remove_node(n)
```

Remove node n.

Removes the node n and all adjacent edges. Attempting to remove a non-existent node will raise an exception.

Parameters n: node

A node in the graph

## Raises NetworkXError

If n is not in the graph.

## See also:

```
remove_nodes_from
```

## **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.edges()
[(0, 1), (1, 2)]
>>> G.remove_node(1)
>>> G.edges()
[]
```

### remove\_nodes\_from

```
MultiDiGraph.remove_nodes_from(nbunch)
```

Remove multiple nodes.

Parameters nodes: iterable container

A container of nodes (list, dict, set, etc.). If a node in the container is not in the graph it is silently ignored.

## See also:

```
remove_node
```

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> e = G.nodes()
>>> e
[0, 1, 2]
>>> G.remove_nodes_from(e)
>>> G.nodes()
[]
```

## add\_edge

```
MultiDiGraph.add_edge (u, v, key=None, attr_dict=None, **attr)
Add an edge between u and v.
```

The nodes u and v will be automatically added if they are not already in the graph.

Edge attributes can be specified with keywords or by providing a dictionary with key/value pairs. See examples below.

### Parameters u,v: nodes

Nodes can be, for example, strings or numbers. Nodes must be hashable (and not None) Python objects.

key: hashable identifier, optional (default=lowest unused integer)

Used to distinguish multiedges between a pair of nodes.

```
attr_dict : dictionary, optional (default= no attributes)
```

Dictionary of edge attributes. Key/value pairs will update existing data associated with the edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

## See also:

```
add_edges_from add a collection of edges
```

### Notes

To replace/update edge data, use the optional key argument to identify a unique edge. Otherwise a new edge will be created.

NetworkX algorithms designed for weighted graphs cannot use multigraphs directly because it is not clear how to handle multiedge weights. Convert to Graph using edge attribute 'weight' to enable weighted graph algorithms.

## **Examples**

The following all add the edge e=(1,2) to graph G:

```
>>> G = nx.MultiDiGraph()
>>> e = (1,2)
>>> G.add_edge(1, 2)  # explicit two-node form
>>> G.add_edge(*e)  # single edge as tuple of two nodes
>>> G.add_edges_from([(1,2)]) # add edges from iterable container
```

Associate data to edges using keywords:

```
>>> G.add_edge(1, 2, weight=3)
>>> G.add_edge(1, 2, key=0, weight=4) # update data for key=0
>>> G.add_edge(1, 3, weight=7, capacity=15, length=342.7)
```

### add edges from

```
MultiDiGraph.add_edges_from(ebunch, attr_dict=None, **attr)
Add all the edges in ebunch.
```

Parameters ebunch: container of edges

Each edge given in the container will be added to the graph. The edges can be:

- 2-tuples (u,v) or
- 3-tuples (u,v,d) for an edge attribute dict d, or
- 4-tuples (u,v,k,d) for an edge identified by key k

attr\_dict : dictionary, optional (default= no attributes)

Dictionary of edge attributes. Key/value pairs will update existing data associated with each edge.

attr: keyword arguments, optional

Edge data (or labels or objects) can be assigned using keyword arguments.

#### See also:

```
add_edge add a single edge
add_weighted_edges_from convenient way to add weighted edges
```

### **Notes**

Adding the same edge twice has no effect but any edge data will be updated when each duplicate edge is added. Edge attributes specified in edges take precedence over attributes specified generally.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edges_from([(0,1),(1,2)]) # using a list of edge tuples
>>> e = zip(range(0,3),range(1,4))
>>> G.add_edges_from(e) # Add the path graph 0-1-2-3
```

Associate data to edges

```
>>> G.add_edges_from([(1,2),(2,3)], weight=3)
>>> G.add_edges_from([(3,4),(1,4)], label='WN2898')
```

#### add weighted edges from

MultiDiGraph.add\_weighted\_edges\_from(ebunch, weight='weight', \*\*attr)

Add all the edges in ebunch as weighted edges with specified weights.

## Parameters ebunch: container of edges

Each edge given in the list or container will be added to the graph. The edges must be given as 3-tuples (u,v,w) where w is a number.

```
weight: string, optional (default= 'weight')
```

The attribute name for the edge weights to be added.

attr: keyword arguments, optional (default= no attributes)

Edge attributes to add/update for all edges.

#### See also:

```
add_edge add a single edge
add_edges_from add multiple edges
```

#### **Notes**

Adding the same edge twice for Graph/DiGraph simply updates the edge data. For MultiGraph/MultiDiGraph, duplicate edges are stored.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_weighted_edges_from([(0,1,3.0),(1,2,7.5)])
```

### remove edge

```
MultiDiGraph.remove edge (u, v, key=None)
```

Remove an edge between u and v.

## Parameters u,v: nodes

Remove an edge between nodes u and v.

**key**: hashable identifier, optional (default=None)

Used to distinguish multiple edges between a pair of nodes. If None remove a single (abritrary) edge between u and v.

## Raises NetworkXError

If there is not an edge between u and v, or if there is no edge with the specified key.

## See also:

remove\_edges\_from remove a collection of edges

## **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2,3])
>>> G.remove_edge(0,1)
>>> e = (1,2)
>>> G.remove_edge(*e) # unpacks e from an edge tuple

For multiple edges
>>> G = nx.MultiDiGraph()
>>> G.add_edges_from([(1,2),(1,2),(1,2)])
>>> G.remove_edge(1,2) # remove a single (arbitrary) edge

For edges with keys
>>> G = nx.MultiDiGraph()
>>> G.add_edge(1,2,key='first')
>>> G.add_edge(1,2,key='second')
>>> G.remove_edge(1,2,key='second')
```

#### remove edges from

```
MultiDiGraph.remove_edges_from(ebunch)
```

Remove all edges specified in ebunch.

## Parameters ebunch: list or container of edge tuples

Each edge given in the list or container will be removed from the graph. The edges can be:

- 2-tuples (u,v) All edges between u and v are removed.
- 3-tuples (u,v,key) The edge identified by key is removed.
- 4-tuples (u,v,key,data) where data is ignored.

## See also:

remove\_edge remove a single edge

### **Notes**

Will fail silently if an edge in ebunch is not in the graph.

## **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> ebunch=[(1,2),(2,3)]
>>> G.remove_edges_from(ebunch)
```

Removing multiple copies of edges

```
>>> G = nx.MultiGraph()
>>> G.add_edges_from([(1,2),(1,2)])
>>> G.remove_edges_from([(1,2),(1,2)])
>>> G.edges()
[(1, 2)]
>>> G.remove_edges_from([(1,2),(1,2)]) # silently ignore extra copy
>>> G.edges() # now empty graph
[]
```

## add\_star

```
MultiDiGraph.add_star(nodes, **attr)
Add a star.
```

The first node in nodes is the middle of the star. It is connected to all other nodes.

Parameters nodes: iterable container

A container of nodes.

attr : keyword arguments, optional (default= no attributes)

Attributes to add to every edge in star.

#### See also:

```
add_path, add_cycle
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_star([0,1,2,3])
>>> G.add_star([10,11,12],weight=2)
```

## add\_path

```
MultiDiGraph.add_path (nodes, **attr)
Add a path.
```

Parameters nodes: iterable container

A container of nodes. A path will be constructed from the nodes (in order) and added to the graph.

**attr**: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in path.

## See also:

```
add_star,add_cycle
```

#### **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.add_path([10,11,12], weight=7)
```

## add\_cycle

```
MultiDiGraph.add_cycle (nodes, **attr) Add a cycle.
```

## Parameters nodes: iterable container

A container of nodes. A cycle will be constructed from the nodes (in order) and added to the graph.

attr: keyword arguments, optional (default= no attributes)

Attributes to add to every edge in cycle.

## See also:

```
add_path, add_star
```

## **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_cycle([0,1,2,3])
>>> G.add_cycle([10,11,12], weight=7)
```

#### clear

```
MultiDiGraph.clear()
```

Remove all nodes and edges from the graph.

This also removes the name, and all graph, node, and edge attributes.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.clear()
>>> G.nodes()
[]
>>> G.edges()
[]
```

## Iterating over nodes and edges

MultiDiGraph.nodes([data])	Return a list of the nodes in the graph.
MultiDiGraph.nodes_iter([data])	Return an iterator over the nodes.
MultiDiGraphiter()	Iterate over the nodes.
MultiDiGraph.edges([nbunch, data, keys, default])	Return a list of edges.
MultiDiGraph.edges_iter([nbunch, data,])	Return an iterator over the edges.
MultiDiGraph.out_edges([nbunch, keys, data])	Return a list of the outgoing edges.
MultiDiGraph.out_edges_iter([nbunch, data,])	Return an iterator over the edges.
MultiDiGraph.in_edges([nbunch, keys, data])	Return a list of the incoming edges.
	Continued on next

Table 3.14 – continued from previous page

<pre>MultiDiGraph.in_edges_iter([nbunch, data, keys])</pre>	Return an iterator over the incoming edges.
MultiDiGraph.get_edge_data(u, v[, key, default])	Return the attribute dictionary associated with edge (u,v).
MultiDiGraph.neighbors(n)	Return a list of successor nodes of n.
MultiDiGraph.neighbors_iter(n)	Return an iterator over successor nodes of n.
MultiDiGraphgetitem(n)	Return a dict of neighbors of node n.
MultiDiGraph.successors(n)	Return a list of successor nodes of n.
MultiDiGraph.successors_iter(n)	Return an iterator over successor nodes of n.
MultiDiGraph.predecessors(n)	Return a list of predecessor nodes of n.
MultiDiGraph.predecessors_iter(n)	Return an iterator over predecessor nodes of n.
MultiDiGraph.adjacency_list()	Return an adjacency list representation of the graph.
MultiDiGraph.adjacency_iter()	Return an iterator of (node, adjacency dict) tuples for all nodes.
MultiDiGraph.nbunch_iter([nbunch])	Return an iterator of nodes contained in nbunch that are also in the gra-

### nodes

MultiDiGraph.nodes (data=False)

Return a list of the nodes in the graph.

**Parameters data**: boolean, optional (default=False)

If False return a list of nodes. If True return a two-tuple of node and node data dictionary

Returns nlist: list

A list of nodes. If data=True a list of two-tuples containing (node, node data dictionary).

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> G.nodes()
[0, 1, 2]
>>> G.add_node(1, time='5pm')
>>> G.nodes(data=True)
[(0, {}), (1, {'time': '5pm'}), (2, {})]
```

### nodes iter

MultiDiGraph.nodes\_iter(data=False)

Return an iterator over the nodes.

**Parameters data**: boolean, optional (default=False)

If False the iterator returns nodes. If True return a two-tuple of node and node data dictionary

Returns niter: iterator

An iterator over nodes. If data=True the iterator gives two-tuples containing (node, node data, dictionary)

#### **Notes**

If the node data is not required it is simpler and equivalent to use the expression 'for n in G'.

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
```

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])

>>> [d for n,d in G.nodes_iter(data=True)]
[{}, {}, {}]
_iter__
```

MultiDiGraph.\_\_iter\_\_()

Iterate over the nodes. Use the expression 'for n in G'.

Returns niter: iterator

An iterator over all nodes in the graph.

## **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
```

## edges

MultiDiGraph.edges (nbunch=None, data=False, keys=False, default=None)
Return a list of edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,data) (True).

**keys**: bool, optional (default=False)

Return two tuples (u,v) (False) or three-tuples (u,v,key) (True).

**Returns** edge\_list: list of edge tuples

Edges that are adjacent to any node in nbunch, or a list of all edges if nbunch is not specified.

#### See also:

edges\_iter return an iterator over the edges

### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> G.edges()
[(0, 1), (1, 2), (2, 3)]
>>> G.edges(data=True) # default edge data is {} (empty dictionary)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> G.edges(keys=True) # default keys are integers
[(0, 1, 0), (1, 2, 0), (2, 3, 0)]
>>> G.edges(data=True, keys=True) # default keys are integers
[(0, 1, 0, {}), (1, 2, 0, {}), (2, 3, 0, {'weight': 5})]
>>> list(G.edges(data='weight',default=1,keys=True))
[(0, 1, 0, 1), (1, 2, 0, 1), (2, 3, 0, 5)]
>>> G.edges([0,3])
[(0, 1), (3, 2)]
>>> G.edges(0)
[(0, 1)]
```

### edges\_iter

MultiDiGraph.edges\_iter (nbunch=None, data=False, keys=False, default=None)
Return an iterator over the edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

**data**: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**keys**: bool, optional (default=False)

If True, return edge keys with each edge.

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edge iter: iterator

An iterator of (u,v), (u,v,d) or (u,v,key,d) tuples of edges.

See also:

edges return a list of edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, \{\}), (1, 2, \{\}), (2, 3, \{'weight': 5\})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges(keys=True)) # default keys are integers
[(0, 1, 0), (1, 2, 0), (2, 3, 0)]
>>> list(G.edges(data=True, keys=True)) # default keys are integers
[(0, 1, 0, {}), (1, 2, 0, {}), (2, 3, 0, {'weight': 5})]
>>> list(G.edges(data='weight',default=1,keys=True))
[(0, 1, 0, 1), (1, 2, 0, 1), (2, 3, 0, 5)]
>>> list(G.edges_iter([0,2]))
[(0, 1), (2, 3)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

### out\_edges

MultiDiGraph.out\_edges (nbunch=None, keys=False, data=False)

Return a list of the outgoing edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

data: bool, optional (default=False)

If True, return edge attribute dict with each edge.

**keys**: bool, optional (default=False)

If True, return edge keys with each edge.

Returns out\_edges: list

An listr of (u,v), (u,v,d) or (u,v,key,d) tuples of edges.

### See also:

in edges return a list of incoming edges

### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs edges() is the same as out\_edges().

### out edges iter

MultiDiGraph.out\_edges\_iter (nbunch=None, data=False, keys=False, default=None) Return an iterator over the edges.

Edges are returned as tuples with optional data and keys in the order (node, neighbor, key, data).

**Parameters nbunch**: iterable container, optional (default= all nodes)

A container of nodes. The container will be iterated through once.

**data**: string or bool, optional (default=False)

The edge attribute returned in 3-tuple (u,v,ddict[data]). If True, return edge attribute dict in 3-tuple (u,v,ddict). If False, return 2-tuple (u,v).

**keys**: bool, optional (default=False)

If True, return edge keys with each edge.

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

Returns edge\_iter: iterator

An iterator of (u,v), (u,v,d) or (u,v,key,d) tuples of edges.

### See also:

edges return a list of edges

#### **Notes**

Nodes in nbunch that are not in the graph will be (quietly) ignored. For directed graphs this returns the out-edges.

### **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2])
>>> G.add_edge(2,3,weight=5)
>>> [e for e in G.edges_iter()]
[(0, 1), (1, 2), (2, 3)]
>>> list(G.edges_iter(data=True)) # default data is {} (empty dict)
[(0, 1, {}), (1, 2, {}), (2, 3, {'weight': 5})]
>>> list(G.edges_iter(data='weight', default=1))
[(0, 1, 1), (1, 2, 1), (2, 3, 5)]
>>> list(G.edges(keys=True)) # default keys are integers
[(0, 1, 0), (1, 2, 0), (2, 3, 0)]
>>> list(G.edges(data=True, keys=True)) # default keys are integers
[(0, 1, 0, {}), (1, 2, 0, {}), (2, 3, 0, {'weight': 5})]
>>> list(G.edges(data='weight',default=1,keys=True))
[(0, 1, 0, 1), (1, 2, 0, 1), (2, 3, 0, 5)]
>>> list(G.edges_iter([0,2]))
[(0, 1), (2, 3)]
>>> list(G.edges_iter(0))
[(0, 1)]
```

in edges

# MultiDiGraph.in\_edges (nbunch=None, keys=False, data=False) Return a list of the incoming edges. **Parameters nbunch**: iterable container, optional (default= all nodes) A container of nodes. The container will be iterated through once. data: bool, optional (default=False) If True, return edge attribute dict with each edge. keys: bool, optional (default=False) If True, return edge keys with each edge. Returns in edges: list A list of (u,v), (u,v,d) or (u,v,key,d) tuples of edges. See also: out\_edges return a list of outgoing edges in edges iter MultiDiGraph.in\_edges\_iter(nbunch=None, data=False, keys=False) Return an iterator over the incoming edges. **Parameters nbunch**: iterable container, optional (default= all nodes) A container of nodes. The container will be iterated through once. data: bool, optional (default=False) If True, return edge attribute dict with each edge. **keys**: bool, optional (default=False) If True, return edge keys with each edge. Returns in\_edge\_iter : iterator An iterator of (u,v), (u,v,d) or (u,v,key,d) tuples of edges. See also: edges\_iter return an iterator of edges get edge data MultiDiGraph.get\_edge\_data(u, v, key=None, default=None) Return the attribute dictionary associated with edge (u,v). Parameters u,v: nodes **default:** any Python object (default=None) Value to return if the edge (u,v) is not found.

key: hashable identifier, optional (default=None)

Return data only for the edge with specified key.

# Returns edge\_dict : dictionary

The edge attribute dictionary.

#### **Notes**

It is faster to use G[u][v][key].

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_edge(0,1,key='a',weight=7)
>>> G[0][1]['a'] # key='a'
{'weight': 7}
```

Warning: Assigning G[u][v][key] corrupts the graph data structure. But it is safe to assign attributes to that dictionary,

```
>>> G[0][1]['a']['weight'] = 10
>>> G[0][1]['a']['weight']
10
>>> G[1][0]['a']['weight']
10
```

# **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.get_edge_data(0,1)
{0: {}}
>>> e = (0,1)
>>> G.get_edge_data(*e) # tuple form
{0: {}}
>>> G.get_edge_data('a','b',default=0) # edge not in graph, return 0
0
```

# neighbors

```
MultiDiGraph.neighbors(n)
```

Return a list of successor nodes of n.

neighbors() and successors() are the same function.

### neighbors iter

```
MultiDiGraph.neighbors_iter(n)
```

Return an iterator over successor nodes of n.

neighbors\_iter() and successors\_iter() are the same.

```
getitem
MultiDiGraph.__getitem__(n)
     Return a dict of neighbors of node n. Use the expression 'G[n]'.
          Parameters n: node
                  A node in the graph.
          Returns adj_dict : dictionary
                  The adjacency dictionary for nodes connected to n.
     Notes
     G[n] is similar to G.neighbors(n) but the internal data dictionary is returned instead of a list.
     Assigning G[n] will corrupt the internal graph data structure. Use G[n] for reading data only.
     Examples
     >>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
     >>> G.add_path([0,1,2,3])
     >>> G[0]
      {1: {}}
successors
MultiDiGraph.successors(n)
     Return a list of successor nodes of n.
     neighbors() and successors() are the same function.
successors_iter
MultiDiGraph.successors_iter(n)
     Return an iterator over successor nodes of n.
     neighbors iter() and successors iter() are the same.
predecessors
MultiDiGraph.predecessors(n)
     Return a list of predecessor nodes of n.
predecessors_iter
```

MultiDiGraph.predecessors\_iter(n)

Return an iterator over predecessor nodes of n.

### adjacency list

```
MultiDiGraph.adjacency_list()
```

Return an adjacency list representation of the graph.

The output adjacency list is in the order of G.nodes(). For directed graphs, only outgoing adjacencies are included.

Returns adj list: lists of lists

The adjacency structure of the graph as a list of lists.

### See also:

```
adjacency_iter
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.adjacency_list() # in order given by G.nodes()
[[1], [0, 2], [1, 3], [2]]
```

### adjacency iter

```
MultiDiGraph.adjacency_iter()
```

Return an iterator of (node, adjacency dict) tuples for all nodes.

This is the fastest way to look at every edge. For directed graphs, only outgoing adjacencies are included.

Returns adj\_iter: iterator

An iterator of (node, adjacency dictionary) for all nodes in the graph.

#### See also:

```
adjacency_list
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> [(n,nbrdict) for n,nbrdict in G.adjacency_iter()]
[(0, {1: {}}), (1, {0: {}, 2: {}}), (2, {1: {}, 3: {}}), (3, {2: {}})]
```

### nbunch iter

```
MultiDiGraph.nbunch_iter(nbunch=None)
```

Return an iterator of nodes contained in nbunch that are also in the graph.

The nodes in nbunch are checked for membership in the graph and if not are silently ignored.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

Returns niter: iterator

An iterator over nodes in nbunch that are also in the graph. If nbunch is None, iterate over all nodes in the graph.

#### Raises NetworkXError

If nbunch is not a node or or sequence of nodes. If a node in nbunch is not hashable.

### See also:

```
Graph.__iter__
```

### **Notes**

When nbunch is an iterator, the returned iterator yields values directly from nbunch, becoming exhausted when nbunch is exhausted.

To test whether nbunch is a single node, one can use "if nbunch in self:", even after processing with this routine.

If nbunch is not a node or a (possibly empty) sequence/iterator or None, a NetworkXError is raised. Also, if any object in nbunch is not hashable, a NetworkXError is raised.

# Information about graph structure

MultiDiGraph.has_node(n)	Return True if the graph contains the node n.
MultiDiGraphcontains(n)	Return True if n is a node, False otherwise.
MultiDiGraph.has_edge(u, v[, key])	Return True if the graph has an edge between nodes u and v.
MultiDiGraph.order()	Return the number of nodes in the graph.
MultiDiGraph.number_of_nodes()	Return the number of nodes in the graph.
MultiDiGraphlen()	Return the number of nodes.
MultiDiGraph.degree([nbunch, weight])	Return the degree of a node or nodes.
MultiDiGraph.degree_iter([nbunch, weight])	Return an iterator for (node, degree).
MultiDiGraph.in_degree([nbunch, weight])	Return the in-degree of a node or nodes.
<pre>MultiDiGraph.in_degree_iter([nbunch, weight])</pre>	Return an iterator for (node, in-degree).
MultiDiGraph.out_degree([nbunch, weight])	Return the out-degree of a node or nodes.
<pre>MultiDiGraph.out_degree_iter([nbunch, weight])</pre>	Return an iterator for (node, out-degree).
MultiDiGraph.size([weight])	Return the number of edges.
$ exttt{MultiDiGraph.number\_of\_edges}([u,v])$	Return the number of edges between two nodes.
MultiDiGraph.nodes_with_selfloops()	Return a list of nodes with self loops.
MultiDiGraph.selfloop_edges([data, keys,])	Return a list of selfloop edges.
MultiDiGraph.number_of_selfloops()	Return the number of selfloop edges.

# has\_node

```
MultiDiGraph.has_node(n)
```

Return True if the graph contains the node n.

Parameters n: node

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
```

```
>>> G.has_node(0)
True
```

It is more readable and simpler to use

```
>>> 0 in G
True
```

### contains

```
MultiDiGraph.__contains__(n)
```

Return True if n is a node, False otherwise. Use the expression 'n in G'.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> 1 in G
True
```

### has\_edge

```
MultiDiGraph.has edge (u, v, key=None)
```

Return True if the graph has an edge between nodes u and v.

Parameters u,v: nodes

Nodes can be, for example, strings or numbers.

key: hashable identifier, optional (default=None)

If specified return True only if the edge with key is found.

Returns edge\_ind: bool

True if edge is in the graph, False otherwise.

# **Examples**

Can be called either using two nodes u,v, an edge tuple (u,v), or an edge tuple (u,v,key).

```
>>> G = nx.MultiGraph()  # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.has_edge(0,1)  # using two nodes
True
>>> e = (0,1)
>>> G.has_edge(*e)  # e is a 2-tuple (u,v)
True
>>> G.add_edge(0,1,key='a')
>>> G.has_edge(0,1,key='a')  # specify key
True
>>> e=(0,1,'a')
>>> G.has_edge(*e)  # e is a 3-tuple (u,v,'a')
True
```

The following syntax are equivalent:

```
>>> G.has_edge(0,1)
True
>>> 1 in G[0] # though this gives KeyError if 0 not in G
True
```

### order

```
MultiDiGraph.order()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

### See also:

```
number_of_nodes, __len__
```

### number\_of\_nodes

```
MultiDiGraph.number_of_nodes()
```

Return the number of nodes in the graph.

Returns nnodes: int

The number of nodes in the graph.

### See also:

```
order, __len__
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2])
>>> len(G)
3
```

# \_\_len\_\_

```
MultiDiGraph.__len__()
```

Return the number of nodes. Use the expression 'len(G)'.

Returns nnodes: int

The number of nodes in the graph.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> len(G)
```

#### degree

```
MultiDiGraph.degree (nbunch=None, weight=None)
```

Return the degree of a node or nodes.

The node degree is the number of edges adjacent to that node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

**Returns nd**: dictionary, or number

A dictionary with nodes as keys and degree as values or a number if a single node is specified.

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.degree(0)
1
>>> G.degree([0,1])
{0: 1, 1: 2}
>>> list(G.degree([0,1]).values())
[1, 2]
```

# degree\_iter

```
MultiDiGraph.degree_iter (nbunch=None, weight=None)
```

Return an iterator for (node, degree).

The node degree is the number of edges adjacent to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight: string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, degree).

## See also:

degree

#### **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2,3])
>>> list(G.degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.degree_iter([0,1]))
[(0, 1), (1, 2)]
```

### in degree

MultiDiGraph.in\_degree (nbunch=None, weight=None)

Return the in-degree of a node or nodes.

The node in-degree is the number of edges pointing in to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd: dictionary, or number

A dictionary with nodes as keys and in-degree as values or a number if a single node is specified.

### See also:

```
degree, out degree, in degree iter
```

### **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.in_degree(0)
0
>>> G.in_degree([0,1])
{0: 0, 1: 1}
>>> list(G.in_degree([0,1]).values())
[0, 1]
```

# in\_degree\_iter

```
MultiDiGraph.in_degree_iter(nbunch=None, weight=None)
```

Return an iterator for (node, in-degree).

The node in-degree is the number of edges pointing in to the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

weight : string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, in-degree).

### See also:

```
degree, in_degree, out_degree_iter
```

### **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2,3])
>>> list(G.in_degree_iter(0)) # node 0 with degree 0
[(0, 0)]
>>> list(G.in_degree_iter([0,1]))
[(0, 0), (1, 1)]
```

### out degree

MultiDiGraph.out\_degree (nbunch=None, weight=None)

Return the out-degree of a node or nodes.

The node out-degree is the number of edges pointing out of the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

**Returns nd**: dictionary, or number

A dictionary with nodes as keys and out-degree as values or a number if a single node is specified.

### **Examples**

```
>>> G = nx.DiGraph() # or MultiDiGraph
>>> G.add_path([0,1,2,3])
>>> G.out_degree(0)
1
>>> G.out_degree([0,1])
{0: 1, 1: 1}
>>> list(G.out_degree([0,1]).values())
[1, 1]
```

# out\_degree\_iter

```
MultiDiGraph.out_degree_iter (nbunch=None, weight=None)
Return an iterator for (node, out-degree).
```

The node out-degree is the number of edges pointing out of the node.

**Parameters nbunch**: iterable container, optional (default=all nodes)

A container of nodes. The container will be iterated through once.

```
weight : string or None, optional (default=None)
```

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights.

Returns nd\_iter: an iterator

The iterator returns two-tuples of (node, out-degree).

### See also:

```
degree, in_degree, out_degree, in_degree_iter
```

# **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1,2,3])
>>> list(G.out_degree_iter(0)) # node 0 with degree 1
[(0, 1)]
>>> list(G.out_degree_iter([0,1]))
[(0, 1), (1, 1)]
```

#### size

MultiDiGraph.size(weight=None)

Return the number of edges.

Parameters weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

Returns nedges: int

The number of edges or sum of edge weights in the graph.

### See also:

```
number_of_edges
```

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.size()
3
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge('a','b',weight=2)
>>> G.add_edge('b','c',weight=4)
>>> G.size()
2
```

```
>>> G.size(weight='weight')
6.0
```

### number\_of\_edges

```
MultiDiGraph.number of edges (u=None, v=None)
```

Return the number of edges between two nodes.

**Parameters u,v**: nodes, optional (default=all edges)

If u and v are specified, return the number of edges between u and v. Otherwise return the total number of all edges.

# Returns nedges: int

The number of edges in the graph. If nodes u and v are specified return the number of edges between those nodes.

### See also:

size

### **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> G.number_of_edges()
3
>>> G.number_of_edges(0,1)
1
>>> e = (0,1)
>>> G.number_of_edges(*e)
```

### nodes with selfloops

```
MultiDiGraph.nodes_with_selfloops()
```

Return a list of nodes with self loops.

A node with a self loop has an edge with both ends adjacent to that node.

### Returns nodelist: list

A list of nodes with self loops.

### See also:

```
selfloop_edges, number_of_selfloops
```

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.nodes_with_selfloops()
[1]
```

### selfloop edges

```
MultiDiGraph.selfloop_edges (data=False, keys=False, default=None) Return a list of selfloop edges.
```

A selfloop edge has the same node at both ends.

Parameters data: bool, optional (default=False)

Return selfloop edges as two tuples (u,v) (data=False) or three-tuples (u,v,datadict) (data=True) or three-tuples (u,v,datavalue) (data='attrname')

**default**: value, optional (default=None)

Value used for edges that dont have the requested attribute. Only relevant if data is not True or False.

**keys**: bool, optional (default=False)

If True, return edge keys with each edge.

**Returns edgelist**: list of edge tuples

A list of all selfloop edges.

### See also:

```
nodes_with_selfloops, number_of_selfloops
```

### **Examples**

```
>>> G = nx.MultiGraph() # or MultiDiGraph
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.selfloop_edges()
[(1, 1)]
>>> G.selfloop_edges(data=True)
[(1, 1, {})]
>>> G.selfloop_edges(keys=True)
[(1, 1, 0)]
>>> G.selfloop_edges(keys=True, data=True)
[(1, 1, 0, {})]
```

# number\_of\_selfloops

```
MultiDiGraph.number_of_selfloops()
```

Return the number of selfloop edges.

A selfloop edge has the same node at both ends.

**Returns nloops**: int

The number of selfloops.

## See also:

```
nodes_with_selfloops, selfloop_edges
```

### **Examples**

```
>>> G=nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_edge(1,1)
>>> G.add_edge(1,2)
>>> G.number_of_selfloops()
1
```

# Making copies and subgraphs

MultiDiGraph.copy()	Return a copy of the graph.
MultiDiGraph.to_undirected([reciprocal])	Return an undirected representation of the digraph.
MultiDiGraph.to_directed()	Return a directed copy of the graph.
MultiDiGraph.subgraph(nbunch)	Return the subgraph induced on nodes in nbunch.
MultiDiGraph.reverse([copy])	Return the reverse of the graph.

### сору

```
MultiDiGraph.copy()
```

Return a copy of the graph.

Returns G: Graph

A copy of the graph.

### See also:

to\_directed return a directed copy of the graph.

### Notes

This makes a complete copy of the graph including all of the node or edge attributes.

# **Examples**

```
>>> G = nx.Graph() # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.copy()
```

# to\_undirected

MultiDiGraph.to\_undirected(reciprocal=False)

Return an undirected representation of the digraph.

Parameters reciprocal: bool (optional)

If True only keep edges that appear in both directions in the original digraph.

Returns G: MultiGraph

An undirected graph with the same name and nodes and with edge (u,v,data) if either (u,v,data) or (v,u,data) is in the digraph. If both edges exist in digraph and their edge data is different, only one edge is created with an arbitrary choice of which edge data to use. You must check and correct for this manually if desired.

#### **Notes**

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar D=DiGraph(G) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

### to directed

```
MultiDiGraph.to_directed()

Return a directed copy of the graph.
```

Returns G: MultiDiGraph

A deepcopy of the graph.

#### **Notes**

If edges in both directions (u,v) and (v,u) exist in the graph, attributes for the new undirected edge will be a combination of the attributes of the directed edges. The edge data is updated in the (arbitrary) order that the edges are encountered. For more customized control of the edge attributes use add\_edge().

This returns a "deepcopy" of the edge, node, and graph attributes which attempts to completely copy all of the data and references.

This is in contrast to the similar G=DiGraph(D) which returns a shallow copy of the data.

See the Python copy module for more information on shallow and deep copies, http://docs.python.org/library/copy.html.

### **Examples**

```
>>> G = nx.Graph() # or MultiGraph, etc
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1), (1, 0)]
```

If already directed, return a (deep) copy

```
>>> G = nx.MultiDiGraph()
>>> G.add_path([0,1])
>>> H = G.to_directed()
>>> H.edges()
[(0, 1)]
```

#### subgraph

```
MultiDiGraph.subgraph(nbunch)
```

Return the subgraph induced on nodes in nbunch.

The induced subgraph of the graph contains the nodes in nbunch and the edges between those nodes.

Parameters nbunch: list, iterable

A container of nodes which will be iterated through once.

Returns G: Graph

A subgraph of the graph with the same edge attributes.

#### **Notes**

The graph, edge or node attributes just point to the original graph. So changes to the node or edge structure will not be reflected in the original graph while changes to the attributes will.

To create a subgraph with its own copy of the edge/node attributes use: nx.Graph(G.subgraph(nbunch))

If edge attributes are containers, a deep copy can be obtained using: G.subgraph(nbunch).copy()

For an inplace reduction of a graph to a subgraph you can remove nodes: G.remove\_nodes\_from([ n in G if n not in set(nbunch)])

# **Examples**

```
>>> G = nx.Graph()  # or DiGraph, MultiGraph, MultiDiGraph, etc
>>> G.add_path([0,1,2,3])
>>> H = G.subgraph([0,1,2])
>>> H.edges()
[(0, 1), (1, 2)]
```

### reverse

```
MultiDiGraph.reverse(copy=True)
```

Return the reverse of the graph.

The reverse is a graph with the same nodes and edges but with the directions of the edges reversed.

Parameters copy: bool optional (default=True)

If True, return a new DiGraph holding the reversed edges. If False, reverse the reverse graph is created using the original graph (this changes the original graph).

**CHAPTER** 

# **FOUR**

# **ALGORITHMS**

# 4.1 Approximation

# **4.1.1 Clique**

Cliques.

max_clique(G)	Find the Maximum Clique
clique_removal(G)	Repeatedly remove cliques from the graph.

### max\_clique

 $\max\_{\texttt{clique}}(G)$ 

Find the Maximum Clique

Finds the  $O(|V|/(log|V|)^2)$  apx of maximum clique/independent set in the worst case.

Parameters G: NetworkX graph

Undirected graph

Returns clique: set

The apx-maximum clique of the graph

### **Notes**

A clique in an undirected graph G = (V, E) is a subset of the vertex set  $C \subseteq V$ , such that for every two vertices in C, there exists an edge connecting the two. This is equivalent to saying that the subgraph induced by C is complete (in some cases, the term clique may also refer to the subgraph).

A maximum clique is a clique of the largest possible size in a given graph. The clique number  $\omega(G)$  of a graph G is the number of vertices in a maximum clique in G. The intersection number of G is the smallest number of cliques that together cover all edges of G.

http://en.wikipedia.org/wiki/Maximum\_clique

### References

[R161]

### clique removal

### clique removal(G)

Repeatedly remove cliques from the graph.

Results in a  $O(|V|/(\log |V|)^2)$  approximation of maximum clique & independent set. Returns the largest independent set found, along with found maximal cliques.

Parameters G: NetworkX graph

Undirected graph

Returns max\_ind\_cliques: (set, list) tuple

Maximal independent set and list of maximal cliques (sets) in the graph.

### References

[R160]

# 4.1.2 Clustering

average\_clustering(G[, trials]) Estimates the average clustering coefficient of G.

# average\_clustering

### average\_clustering(G, trials=1000)

Estimates the average clustering coefficient of G.

The local clustering of each node in G is the fraction of triangles that actually exist over all possible triangles in its neighborhood. The average clustering coefficient of a graph G is the mean of local clusterings.

This function finds an approximate average clustering coefficient for G by repeating n times (defined in trials) the following experiment: choose a node at random, choose two of its neighbors at random, and check if they are connected. The approximate coefficient is the fraction of triangles found over the number of trials [R162].

Parameters G: NetworkX graph

trials: integer

Number of trials to perform (default 1000).

Returns c: float

Approximated average clustering coefficient.

### References

[R162]

# 4.1.3 Dominating Set

A dominating set for a graph G = (V, E) is a subset D of V such that every vertex not in D is joined to at least one member of D by some edge. The domination number gamma(G) is the number of vertices in a smallest dominating set for G. Given a graph G = (V, E) find a minimum weight dominating set V'.

http://en.wikipedia.org/wiki/Dominating\_set

An edge dominating set for a graph G = (V, E) is a subset D of E such that every edge not in D is adjacent to at least one edge in D.

http://en.wikipedia.org/wiki/Edge\_dominating\_set

$min\_weighted\_dominating\_set(G[, weight])$	Return minimum weight vertex dominating set.
$min\_edge\_dominating\_set(G)$	Return minimum cardinality edge dominating set.

# min\_weighted\_dominating\_set

### min\_weighted\_dominating\_set (G, weight=None)

Return minimum weight vertex dominating set.

**Parameters G**: NetworkX graph

Undirected graph

**weight**: None or string, optional (default = None)

If None, every edge has weight/distance/weight 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

Returns min\_weight\_dominating\_set : set

Returns a set of vertices whose weight sum is no more than log w(V) \* OPT

### **Notes**

This algorithm computes an approximate minimum weighted dominating set for the graph G. The upper-bound on the size of the solution is  $\log w(V) * OPT$ . Runtime of the algorithm is O(|E|).

### References

[R163]

# min\_edge\_dominating\_set

# $min\_edge\_dominating\_set(G)$

Return minimum cardinality edge dominating set.

**Parameters G**: NetworkX graph

Undirected graph

Returns min\_edge\_dominating\_set : set

Returns a set of dominating edges whose size is no more than 2 \* OPT.

### **Notes**

The algorithm computes an approximate solution to the edge dominating set problem. The result is no more than 2 \* OPT in terms of size of the set. Runtime of the algorithm is O(|E|).

4.1. Approximation 125

# 4.1.4 Independent Set

Independent Set

Independent set or stable set is a set of vertices in a graph, no two of which are adjacent. That is, it is a set I of vertices such that for every two vertices in I, there is no edge connecting the two. Equivalently, each edge in the graph has at most one endpoint in I. The size of an independent set is the number of vertices it contains.

A maximum independent set is a largest independent set for a given graph G and its size is denoted  $\alpha(G)$ . The problem of finding such a set is called the maximum independent set problem and is an NP-hard optimization problem. As such, it is unlikely that there exists an efficient algorithm for finding a maximum independent set of a graph.

http://en.wikipedia.org/wiki/Independent\_set\_(graph\_theory)

Independent set algorithm is based on the following paper:

 $O(|V|/(log|V|)^2)$  apx of maximum clique/independent set.

Boppana, R., & Halldórsson, M. M. (1992). Approximating maximum independent sets by excluding subgraphs. BIT Numerical Mathematics, 32(2), 180–196. Springer. doi:10.1007/BF01994876

maximum\_independent\_set(G) Return an approximate maximum independent set.

# maximum\_independent\_set

### maximum independent set(G)

Return an approximate maximum independent set.

**Parameters G** : NetworkX graph

Undirected graph

Returns iset: Set

The apx-maximum independent set

### **Notes**

Finds the  $O(|V|/(log|V|)^2)$  apx of independent set in the worst case.

### References

[R164]

# 4.1.5 Matching

Given a graph G = (V,E), a matching M in G is a set of pairwise non-adjacent edges; that is, no two edges share a common vertex.

http://en.wikipedia.org/wiki/Matching\_(graph\_theory)

min\_maximal\_matching(G) Returns the minimum maximal matching of G.

# min\_maximal\_matching

## $min_maximal_matching(G)$

Returns the minimum maximal matching of G. That is, out of all maximal matchings of the graph G, the smallest is returned.

**Parameters G**: NetworkX graph

Undirected graph

Returns min\_maximal\_matching: set

Returns a set of edges such that no two edges share a common endpoint and every edge not in the set shares some common endpoint in the set. Cardinality will be 2\*OPT in the worst case.

### **Notes**

The algorithm computes an approximate solution fo the minimum maximal cardinality matching problem. The solution is no more than 2 \* OPT in size. Runtime is O(|E|).

### References

[R165]

# 4.1.6 Ramsey

Ramsey numbers.

ramsey\_R2(G) Approximately computes the Ramsey number R(2; s, t) for graph.

# ramsey\_R2

# ${\tt ramsey\_R2}\;(G)$

Approximately computes the Ramsey number R(2; s, t) for graph.

Parameters G: NetworkX graph

Undirected graph

**Returns** max\_pair: (set, set) tuple

Maximum clique, Maximum independent set.

## 4.1.7 Vertex Cover

Given an undirected graph G=(V,E) and a function w assigning nonnegative weights to its vertices, find a minimum weight subset of V such that each edge in E is incident to at least one vertex in the subset.

http://en.wikipedia.org/wiki/Vertex\_cover

min\_weighted\_vertex\_cover(G[, weight]) 2-OPT Local Ratio for Minimum Weighted Vertex Cover

4.1. Approximation 127

### min weighted vertex cover

### min\_weighted\_vertex\_cover(G, weight=None)

2-OPT Local Ratio for Minimum Weighted Vertex Cover

Find an approximate minimum weighted vertex cover of a graph.

Parameters G: NetworkX graph

Undirected graph

**weight**: None or string, optional (default = None)

If None, every edge has weight/distance/cost 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

Returns min\_weighted\_cover: set

Returns a set of vertices whose weight sum is no more than 2 \* OPT.

### **Notes**

Local-Ratio algorithm for computing an approximate vertex cover. Algorithm greedily reduces the costs over edges and iteratively builds a cover. Worst-case runtime is O(|E|).

#### References

[R166]

# 4.2 Assortativity

# 4.2.1 Assortativity

$\texttt{degree\_assortativity\_coefficient}(G[, x, y,])$	Compute degree assortativity of graph.
attribute_assortativity_coefficient(G, attribute)	Compute assortativity for node attributes.
<pre>numeric_assortativity_coefficient(G, attribute)</pre>	Compute assortativity for numerical node attributes.
degree_pearson_correlation_coefficient(G[,])	Compute degree assortativity of graph.

# degree\_assortativity\_coefficient

```
\textbf{degree\_assortativity\_coefficient} \ (G, x = 'out', y = 'in', weight = None, nodes = None)
```

Compute degree assortativity of graph.

Assortativity measures the similarity of connections in the graph with respect to the node degree.

Parameters G: NetworkX graph

x: string ('in','out')

The degree type for source node (directed graphs only).

y: string ('in','out')

The degree type for target node (directed graphs only).

# weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

### nodes: list or iterable (optional)

Compute degree assortativity only for nodes in container. The default is all nodes.

### Returns r: float

Assortativity of graph by degree.

### See also:

```
attribute_assortativity_coefficient, numeric_assortativity_coefficient, neighbor_connectivity, degree_mixing_dict, degree_mixing_matrix
```

#### **Notes**

This computes Eq. (21) in Ref. [R170], where e is the joint probability distribution (mixing matrix) of the degrees. If G is directed than the matrix e is the joint probability of the user-specified degree type for the source and target.

#### References

```
[R170], [R171]
```

# **Examples**

```
>>> G=nx.path_graph(4)
>>> r=nx.degree_assortativity_coefficient(G)
>>> print("%3.1f"%r)
-0.5
```

# attribute\_assortativity\_coefficient

# attribute\_assortativity\_coefficient(G, attribute, nodes=None)

Compute assortativity for node attributes.

Assortativity measures the similarity of connections in the graph with respect to the given attribute.

Parameters G: NetworkX graph

attribute: string

Node attribute key

nodes: list or iterable (optional)

Compute attribute assortativity for nodes in container. The default is all nodes.

Returns r: float

Assortativity of graph for given attribute

4.2. Assortativity 129

### **Notes**

This computes Eq. (2) in Ref. [R167], trace(M)-sum(M)/(1-sum(M), where M is the joint probability distribution (mixing matrix) of the specified attribute.

### References

[R167]

# **Examples**

```
>>> G=nx.Graph()
>>> G.add_nodes_from([0,1],color='red')
>>> G.add_nodes_from([2,3],color='blue')
>>> G.add_edges_from([(0,1),(2,3)])
>>> print(nx.attribute_assortativity_coefficient(G,'color'))
1.0
```

## numeric assortativity coefficient

```
numeric_assortativity_coefficient(G, attribute, nodes=None)
```

Compute assortativity for numerical node attributes.

Assortativity measures the similarity of connections in the graph with respect to the given numeric attribute.

```
Parameters G: NetworkX graph
```

attribute: string

Node attribute key

nodes: list or iterable (optional)

Compute numeric assortativity only for attributes of nodes in container. The default is all nodes.

Returns r: float

Assortativity of graph for given attribute

### **Notes**

This computes Eq. (21) in Ref. [R175], for the mixing matrix of of the specified attribute.

#### References

[R175]

# **Examples**

```
>>> G=nx.Graph()
>>> G.add_nodes_from([0,1],size=2)
>>> G.add_nodes_from([2,3],size=3)
>>> G.add_edges_from([(0,1),(2,3)])
>>> print(nx.numeric_assortativity_coefficient(G,'size'))
1 0
```

## degree pearson correlation coefficient

```
degree_pearson_correlation_coefficient (G, x='out', y='in', weight=None, nodes=None) Compute degree assortativity of graph.
```

Assortativity measures the similarity of connections in the graph with respect to the node degree.

This is the same as degree\_assortativity\_coefficient but uses the potentially faster scipy.stats.pearsonr function.

# Parameters G: NetworkX graph

```
x: string ('in','out')
```

The degree type for source node (directed graphs only).

```
y: string ('in','out')
```

The degree type for target node (directed graphs only).

# weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

### nodes: list or iterable (optional)

Compute pearson correlation of degrees only for specified nodes. The default is all nodes.

### Returns r: float

Assortativity of graph by degree.

### **Notes**

This calls scipy.stats.pearsonr.

### References

```
[R172], [R173]
```

# **Examples**

```
>>> G=nx.path_graph(4)
>>> r=nx.degree_pearson_correlation_coefficient(G)
>>> print("%3.1f"%r)
-0.5
```

4.2. Assortativity 131

# 4.2.2 Average neighbor degree

average\_neighbor\_degree(G[, source, target, ...]) Returns the average degree of the neighborhood of each node.

# average\_neighbor\_degree

 $\textbf{average\_neighbor\_degree} \ (\textit{G}, \textit{source='out'}, \textit{target='out'}, \textit{nodes=None}, \textit{weight=None})$ 

Returns the average degree of the neighborhood of each node.

The average degree of a node i is

$$k_{nn,i} = \frac{1}{|N(i)|} \sum_{j \in N(i)} k_j$$

where N(i) are the neighbors of node i and  $k_j$  is the degree of node j which belongs to N(i). For weighted graphs, an analogous measure can be defined [R169],

$$k_{nn,i}^w = \frac{1}{s_i} \sum_{j \in N(i)} w_{ij} k_j$$

where  $s_i$  is the weighted degree of node i,  $w_{ij}$  is the weight of the edge that links i and j and N(i) are the neighbors of node i.

Parameters G: NetworkX graph

source: string ("in"l"out")

Directed graphs only. Use "in"- or "out"-degree for source node.

target : string ("in"|"out")

Directed graphs only. Use "in"- or "out"-degree for target node.

nodes: list or iterable, optional

Compute neighbor degree for specified nodes. The default is all nodes in the graph.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

Returns d: dict

A dictionary keyed by node with average neighbors degree value.

### See also:

average\_degree\_connectivity

# **Notes**

For directed graphs you can also specify in-degree or out-degree by passing keyword arguments.

# References

[R169]

### **Examples**

```
>>> G=nx.path_graph(4)
>>> G.edge[0][1]['weight'] = 5
>>> G.edge[2][3]['weight'] = 3

>>> nx.average_neighbor_degree(G)
{0: 2.0, 1: 1.5, 2: 1.5, 3: 2.0}
>>> nx.average_neighbor_degree(G, weight='weight')
{0: 2.0, 1: 1.166666666666666667, 2: 1.25, 3: 2.0}

>>> G=nx.DiGraph()
>>> G=nx.DiGraph()
>>> nx.average_neighbor_degree(G, source='in', target='in')
{0: 1.0, 1: 1.0, 2: 1.0, 3: 0.0}

>>> nx.average_neighbor_degree(G, source='out', target='out')
{0: 1.0, 1: 1.0, 2: 0.0, 3: 0.0}
```

# 4.2.3 Average degree connectivity

```
average_degree_connectivity(G[, source, ...]) Compute the average degree connectivity of graph.

k_nearest_neighbors(G[, source, target, ...]) Compute the average degree connectivity of graph.
```

### average\_degree\_connectivity

**average\_degree\_connectivity** (*G*, source='in+out', target='in+out', nodes=None, weight=None) Compute the average degree connectivity of graph.

The average degree connectivity is the average nearest neighbor degree of nodes with degree k. For weighted graphs, an analogous measure can be computed using the weighted average neighbors degree defined in [R168], for a node i, as:

$$k_{nn,i}^w = \frac{1}{s_i} \sum_{j \in N(i)} w_{ij} k_j$$

where  $s_i$  is the weighted degree of node i,  $w_{ij}$  is the weight of the edge that links i and j, and N(i) are the neighbors of node i.

**Parameters G**: NetworkX graph

source : "in"|"out"|"in+out" (default:"in+out")

Directed graphs only. Use "in"- or "out"-degree for source node.

target: "in"|"out"|"in+out" (default:"in+out"

Directed graphs only. Use "in"- or "out"-degree for target node.

nodes: list or iterable (optional)

Compute neighbor connectivity for these nodes. The default is all nodes.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

4.2. Assortativity 133

### Returns d: dict

A dictionary keyed by degree k with the value of average connectivity.

### See also:

```
neighbors_average_degree
```

#### **Notes**

This algorithm is sometimes called "k nearest neighbors' and is also available as  $k_n earest_n eighbors$ .

### References

[R168]

# **Examples**

```
>>> G=nx.path_graph(4)
>>> G.edge[1][2]['weight'] = 3
>>> nx.k_nearest_neighbors(G)
{1: 2.0, 2: 1.5}
>>> nx.k_nearest_neighbors(G, weight='weight')
{1: 2.0, 2: 1.75}
```

### k nearest neighbors

**k\_nearest\_neighbors** (*G*, source='in+out', target='in+out', nodes=None, weight=None) Compute the average degree connectivity of graph.

The average degree connectivity is the average nearest neighbor degree of nodes with degree k. For weighted graphs, an analogous measure can be computed using the weighted average neighbors degree defined in [R174], for a node i, as:

$$k_{nn,i}^w = \frac{1}{s_i} \sum_{j \in N(i)} w_{ij} k_j$$

where  $s_i$  is the weighted degree of node i,  $w_{ij}$  is the weight of the edge that links i and j, and N(i) are the neighbors of node i.

### **Parameters G**: NetworkX graph

```
source: "in"|"out"|"in+out" (default:"in+out")
```

Directed graphs only. Use "in"- or "out"-degree for source node.

target: "in"|"out"|"in+out" (default:"in+out"

Directed graphs only. Use "in"- or "out"-degree for target node.

### nodes: list or iterable (optional)

Compute neighbor connectivity for these nodes. The default is all nodes.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

#### Returns d: dict

A dictionary keyed by degree k with the value of average connectivity.

### See also:

```
neighbors_average_degree
```

### **Notes**

This algorithm is sometimes called "k nearest neighbors' and is also available as  $k_n earest_n eighbors$ .

## References

[R174]

### **Examples**

```
>>> G=nx.path_graph(4)
>>> G.edge[1][2]['weight'] = 3
>>> nx.k_nearest_neighbors(G)
{1: 2.0, 2: 1.5}
>>> nx.k_nearest_neighbors(G, weight='weight')
{1: 2.0, 2: 1.75}
```

# **4.2.4 Mixing**

attribute_mixing_matrix(G, attribute[,])	Return mixing matrix for attribute.
$degree\_mixing\_matrix(G[, x, y, weight,])$	Return mixing matrix for attribute.
$degree\_mixing\_dict(G[, x, y, weight, nodes,])$	Return dictionary representation of mixing matrix for degree.
attribute_mixing_dict(G, attribute[, nodes,])	Return dictionary representation of mixing matrix for attribute.

# attribute\_mixing\_matrix

```
Parameters G: graph
```

NetworkX graph object.

attribute: string

Node attribute key.

nodes: list or iterable (optional)

Use only nodes in container to build the matrix. The default is all nodes.

mapping: dictionary, optional

4.2. Assortativity 135

Mapping from node attribute to integer index in matrix. If not specified, an arbitrary ordering will be used.

normalized: bool (default=False)

Return counts if False or probabilities if True.

**Returns** m: numpy array

Counts or joint probability of occurrence of attribute pairs.

# degree\_mixing\_matrix

 $degree_mixing_matrix$  (G, x='out', y='in', weight=None, nodes=None, normalized=True)

Return mixing matrix for attribute.

Parameters G: graph

NetworkX graph object.

x: string ('in','out')

The degree type for source node (directed graphs only).

y: string ('in','out')

The degree type for target node (directed graphs only).

nodes: list or iterable (optional)

Build the matrix using only nodes in container. The default is all nodes.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

normalized: bool (default=False)

Return counts if False or probabilities if True.

**Returns** m: numpy array

Counts, or joint probability, of occurrence of node degree.

# degree\_mixing\_dict

 $degree_mixing_dict(G, x='out', y='in', weight=None, nodes=None, normalized=False)$ Return dictionary representation of mixing matrix for degree.

 $Parameters \ G: {\tt graph}$ 

NetworkX graph object.

x: string ('in','out')

The degree type for source node (directed graphs only).

y: string ('in','out')

The degree type for target node (directed graphs only).

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

```
normalized: bool (default=False)
```

Return counts if False or probabilities if True.

# Returns d: dictionary

Counts or joint probability of occurrence of degree pairs.

# attribute mixing dict

```
attribute_mixing_dict (G, attribute, nodes=None, normalized=False)
Return dictionary representation of mixing matrix for attribute.
```

## Parameters G: graph

NetworkX graph object.

attribute: string

Node attribute key.

# nodes: list or iterable (optional)

Unse nodes in container to build the dict. The default is all nodes.

normalized: bool (default=False)

Return counts if False or probabilities if True.

# Returns d: dictionary

Counts or joint probability of occurrence of attribute pairs.

# **Examples**

```
>>> G=nx.Graph()
>>> G.add_nodes_from([0,1],color='red')
>>> G.add_nodes_from([2,3],color='blue')
>>> G.add_edge(1,3)
>>> d=nx.attribute_mixing_dict(G,'color')
>>> print(d['red']['blue'])
1
>>> print(d['blue']['red']) # d symmetric for undirected graphs
1
```

# 4.3 Bipartite

This module provides functions and operations for bipartite graphs. Bipartite graphs B=(U,V,E) have two node sets U,V and edges in E that only connect nodes from opposite sets. It is common in the literature to use an spatial analogy referring to the two node sets as top and bottom nodes.

The bipartite algorithms are not imported into the networkx namespace at the top level so the easiest way to use them is with:

```
>>> import networkx as nx
>>> from networkx.algorithms import bipartite
```

4.3. Bipartite 137

NetworkX does not have a custom bipartite graph class but the Graph() or DiGraph() classes can be used to represent bipartite graphs. However, you have to keep track of which set each node belongs to, and make sure that there is no edge between nodes of the same set. The convention used in NetworkX is to use a node attribute named "bipartite" with values 0 or 1 to identify the sets each node belongs to.

For example:

```
>>> B = nx.Graph()
>>> B.add_nodes_from([1,2,3,4], bipartite=0) # Add the node attribute "bipartite"
>>> B.add_nodes_from(['a','b','c'], bipartite=1)
>>> B.add_edges_from([(1,'a'), (1,'b'), (2,'b'), (2,'c'), (3,'c'), (4,'a')])
```

Many algorithms of the bipartite module of NetworkX require, as an argument, a container with all the nodes that belong to one set, in addition to the bipartite graph B. If B is connected, you can find the node sets using a two-coloring algorithm:

```
>>> nx.is_connected(B)
True
>>> bottom_nodes, top_nodes = bipartite.sets(B)
```

list(top\_nodes) [1, 2, 3, 4] list(bottom\_nodes) ['a', 'c', 'b']

However, if the input graph is not connected, there are more than one possible colorations. Thus, the following result is correct:

```
>>> B.remove_edge(2,'c')
>>> nx.is_connected(B)
False
>>> bottom_nodes, top_nodes = bipartite.sets(B)
```

list(top\_nodes) [1, 2, 4, 'c'] list(bottom\_nodes) ['a', 3, 'b']

Using the "bipartite" node attribute, you can easily get the two node sets:

```
>>> top_nodes = set(n for n,d in B.nodes(data=True) if d['bipartite']==0)
>>> bottom_nodes = set(B) - top_nodes
```

```
list(top_nodes) [1, 2, 3, 4] list(bottom_nodes) ['a', 'c', 'b']
```

So you can easily use the bipartite algorithms that require, as an argument, a container with all nodes that belong to one node set:

```
>>> print(round(bipartite.density(B, bottom_nodes),2))
0.42
>>> G = bipartite.projected_graph(B, top_nodes)
>>> G.edges()
[(1, 2), (1, 4)]
```

All bipartite graph generators in NetworkX build bipartite graphs with the "bipartite" node attribute. Thus, you can use the same approach:

```
>>> RB = bipartite.random_graph(5, 7, 0.2)
>>> RB_top = set(n for n,d in RB.nodes(data=True) if d['bipartite']==0)
>>> RB_bottom = set(RB) - RB_top
>>> list(RB_top)
[0, 1, 2, 3, 4]
>>> list(RB_bottom)
[5, 6, 7, 8, 9, 10, 11]
```

For other bipartite graph generators see the bipartite section of *Graph generators*.

# 4.3.1 Basic functions

is_bipartite(G)	Returns True if graph G is bipartite, False if not.
is_bipartite_node_set(G, nodes)	Returns True if nodes and G/nodes are a bipartition of G.
sets(G)	Returns bipartite node sets of graph G.
color(G)	Returns a two-coloring of the graph.
density(B, nodes)	Return density of bipartite graph B.
degrees(B, nodes[, weight])	Return the degrees of the two node sets in the bipartite graph B.
biadjacency_matrix	

# is\_bipartite

# $is\_bipartite(G)$

Returns True if graph G is bipartite, False if not.

Parameters G: NetworkX graph

## See also:

```
color, is_bipartite_node_set
```

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4)
>>> print(bipartite.is_bipartite(G))
True
```

# is\_bipartite\_node\_set

# is\_bipartite\_node\_set(G, nodes)

Returns True if nodes and G/nodes are a bipartition of G.

Parameters G: NetworkX graph

nodes: list or container

Check if nodes are a one of a bipartite set.

## **Notes**

For connected graphs the bipartite sets are unique. This function handles disconnected graphs.

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4)
>>> X = set([1,3])
>>> bipartite.is_bipartite_node_set(G,X)
True
```

#### sets

#### sets(G)

Returns bipartite node sets of graph G.

Raises an exception if the graph is not bipartite.

**Parameters** G: NetworkX graph **Returns** (X,Y): two-tuple of sets

One set of nodes for each part of the bipartite graph.

## See also:

color

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4)
>>> X, Y = bipartite.sets(G)
>>> list(X)
[0, 2]
>>> list(Y)
[1, 3]
```

#### color

## $\mathtt{color}(G)$

Returns a two-coloring of the graph.

Raises an exception if the graph is not bipartite.

Parameters G: NetworkX graph

Returns color: dictionary

A dictionary keyed by node with a 1 or 0 as data for each node color.

Raises NetworkXError if the graph is not two-colorable.

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4)
>>> c = bipartite.color(G)
>>> print(c)
{0: 1, 1: 0, 2: 1, 3: 0}

You can use this to set a node attribute indicating the biparite set:
>>> nx.set_node_attributes(G, 'bipartite', c)
>>> print(G.node[0]['bipartite'])
1
>>> print(G.node[1]['bipartite'])
```

# density

```
density (B, nodes)
Return density of bipartite graph B.

Parameters G: NetworkX graph
nodes: list or container
Nodes in one set of the bipartite graph.

Returns d: float
The bipartite density

See also:
color
```

#### **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.complete_bipartite_graph(3,2)
>>> X=set([0,1,2])
>>> bipartite.density(G,X)
1.0
>>> Y=set([3,4])
>>> bipartite.density(G,Y)
1.0
```

## degrees

```
degrees (B, nodes, weight=None)
```

Return the degrees of the two node sets in the bipartite graph B.

**Parameters G**: NetworkX graph

nodes: list or container

Nodes in one set of the bipartite graph.

**weight**: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1. The degree is the sum of the edge weights adjacent to the node.

Returns (degX,degY): tuple of dictionaries

The degrees of the two bipartite sets as dictionaries keyed by node.

#### See also:

```
color, density
```

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.complete_bipartite_graph(3,2)
>>> Y=set([3,4])
>>> degX,degY=bipartite.degrees(G,Y)
>>> degX
{0: 2, 1: 2, 2: 2}
```

# 4.3.2 Matching

Provides functions for computing a maximum cardinality matching in a bipartite graph.

If you don't care about the particular implementation of the maximum matching algorithm, simply use the maximum\_matching(). If you do care, you can import one of the named maximum matching algorithms directly.

For example, to find a maximum matching in the complete bipartite graph with two vertices on the left and three vertices on the right:

```
>>> import networkx as nx
>>> G = nx.complete_bipartite_graph(2, 3)
>>> left, right = nx.bipartite.sets(G)
>>> list(left)
[0, 1]
>>> list(right)
[2, 3, 4]
>>> nx.bipartite.maximum_matching(G)
{0: 2, 1: 3, 2: 0, 3: 1}
```

The dictionary returned by maximum\_matching() includes a mapping for vertices in both the left and right vertex sets.

eppstein_matching(G)	Returns the maximum cardinality matching of the bipartite graph $G$ .
$hopcroft\_karp\_matching(G)$	Returns the maximum cardinality matching of the bipartite graph $G$ .
to_vertex_cover(G, matching)	Returns the minimum vertex cover corresponding to the given maximum matching of the bipar

## eppstein matching

# ${\tt eppstein\_matching}\,(G)$

Returns the maximum cardinality matching of the bipartite graph G.

**Parameters G**: NetworkX graph

Undirected bipartite graph

Returns matches: dictionary

The matching is returned as a dictionary, matches, such that matches [v] == w if node v is matched to node w. Unmatched nodes do not occur as a key in mate.

# See also:

hopcroft\_karp\_matching

#### **Notes**

This function is implemented with David Eppstein's version of the algorithm Hopcroft-Karp algorithm (see hopcroft\_karp\_matching()), which originally appeared in the Python Algorithms and Data Structures

library (PADS).

## hopcroft\_karp\_matching

## $hopcroft_karp_matching(G)$

Returns the maximum cardinality matching of the bipartite graph G.

Parameters G: NetworkX graph

Undirected bipartite graph

Returns matches: dictionary

The matching is returned as a dictionary, matches, such that matches [v] == w if node v is matched to node w. Unmatched nodes do not occur as a key in mate.

#### See also:

eppstein\_matching

#### **Notes**

This function is implemented with the Hopcroft–Karp matching algorithm for bipartite graphs.

#### References

[R185]

# to\_vertex\_cover

## to\_vertex\_cover(G, matching)

Returns the minimum vertex cover corresponding to the given maximum matching of the bipartite graph G.

## **Parameters G**: NetworkX graph

Undirected bipartite graph

matching: dictionary

A dictionary whose keys are vertices in G and whose values are the distinct neighbors comprising the maximum matching for G, as returned by, for example, maximum\_matching(). The dictionary *must* represent the maximum matching.

# Returns vertex\_cover: set

The minimum vertex cover in G.

## **Notes**

This function is implemented using the procedure guaranteed by Konig's theorem, which proves an equivalence between a maximum matching and a minimum vertex cover in bipartite graphs.

Since a minimum vertex cover is the complement of a maximum independent set for any graph, one can compute the maximum independent set of a bipartite graph this way:

```
>>> import networkx as nx
>>> G = nx.complete_bipartite_graph(2, 3)
>>> matching = nx.bipartite.maximum_matching(G)
>>> vertex_cover = nx.bipartite.to_vertex_cover(G, matching)
>>> independent_set = set(G) - vertex_cover
>>> print(list(independent_set))
[2, 3, 4]
```

# 4.3.3 Projections

One-mode (unipartite) projections of bipartite graphs.

<pre>projected_graph(B, nodes[, multigraph])</pre>	Returns the projection of B onto one of its node sets.
<pre>weighted_projected_graph(B, nodes[, ratio])</pre>	Returns a weighted projection of B onto one of its node sets.
collaboration_weighted_projected_graph(B, nodes)	Newman's weighted projection of B onto one of its node sets.
overlap_weighted_projected_graph( $B, nodes[,]$ )	Overlap weighted projection of B onto one of its node sets.
$generic\_weighted\_projected\_graph(B, nodes[,])$	Weighted projection of B with a user-specified weight function.

# projected\_graph

```
projected_graph (B, nodes, multigraph=False)
```

Returns the projection of B onto one of its node sets.

Returns the graph G that is the projection of the bipartite graph B onto the specified nodes. They retain their attributes and are connected in G if they have a common neighbor in B.

#### **Parameters B**: NetworkX graph

The input graph should be bipartite.

nodes: list or iterable

Nodes to project onto (the "bottom" nodes).

#### multigraph: bool (default=False)

If True return a multigraph where the multiple edges represent multiple shared neighbors. They edge key in the multigraph is assigned to the label of the neighbor.

Returns Graph: NetworkX graph or multigraph

A graph that is the projection onto the given nodes.

#### See also:

```
is_bipartite, is_bipartite_node_set, sets, weighted_projected_graph,
collaboration_weighted_projected_graph, overlap_weighted_projected_graph,
generic_weighted_projected_graph
```

#### **Notes**

No attempt is made to verify that the input graph B is bipartite. Returns a simple graph that is the projection of the bipartite graph B onto the set of nodes given in list nodes. If multigraph=True then a multigraph is returned with an edge for every shared neighbor.

Directed graphs are allowed as input. The output will also then be a directed graph with edges if there is a directed path between the nodes.

The graph and node properties are (shallow) copied to the projected graph.

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> B = nx.path_graph(4)
>>> G = bipartite.projected_graph(B, [1,3])
>>> print(G.nodes())
[1, 3]
>>> print(G.edges())
[(1, 3)]
```

If nodes a, and b are connected through both nodes 1 and 2 then building a multigraph results in two edges in the projection onto [a, b]:

```
>>> B = nx.Graph()
>>> B.add_edges_from([('a', 1), ('b', 1), ('a', 2), ('b', 2)])
>>> G = bipartite.projected_graph(B, ['a', 'b'], multigraph=True)
>>> print([sorted((u,v)) for u,v in G.edges()])
[['a', 'b'], ['a', 'b']]
```

## weighted projected graph

#### weighted\_projected\_graph (B, nodes, ratio=False)

Returns a weighted projection of B onto one of its node sets.

The weighted projected graph is the projection of the bipartite network B onto the specified nodes with weights representing the number of shared neighbors or the ratio between actual shared neighbors and possible shared neighbors if ratio=True [R188]. The nodes retain their attributes and are connected in the resulting graph if they have an edge to a common node in the original graph.

# Parameters B: NetworkX graph

The input graph should be bipartite.

nodes: list or iterable

Nodes to project onto (the "bottom" nodes).

```
ratio: Bool (default=False)
```

If True, edge weight is the ratio between actual shared neighbors and possible shared neighbors. If False, edges weight is the number of shared neighbors.

#### **Returns Graph**: NetworkX graph

A graph that is the projection onto the given nodes.

# See also:

```
is_bipartite, is_bipartite_node_set, sets, collaboration_weighted_projected_graph,
overlap_weighted_projected_graph,
projected_graph
```

#### **Notes**

No attempt is made to verify that the input graph B is bipartite. The graph and node properties are (shallow) copied to the projected graph.

#### References

[R188]

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> B = nx.path_graph(4)
>>> G = bipartite.weighted_projected_graph(B, [1,3])
>>> print(G.nodes())
[1, 3]
>>> print(G.edges(data=True))
[(1, 3, {'weight': 1})]
>>> G = bipartite.weighted_projected_graph(B, [1,3], ratio=True)
>>> print(G.edges(data=True))
[(1, 3, {'weight': 0.5})]
```

## collaboration weighted projected graph

## collaboration\_weighted\_projected\_graph(B, nodes)

Newman's weighted projection of B onto one of its node sets.

The collaboration weighted projection is the projection of the bipartite network B onto the specified nodes with weights assigned using Newman's collaboration model [R186]:

$$w_{v,u} = \sum_{k} \frac{\delta_v^w \delta_w^k}{k_w - 1}$$

where v and u are nodes from the same bipartite node set, and w is a node of the opposite node set. The value  $k_w$  is the degree of node w in the bipartite network and  $\delta_v^w$  is 1 if node v is linked to node w in the original bipartite graph or 0 otherwise.

The nodes retain their attributes and are connected in the resulting graph if have an edge to a common node in the original bipartite graph.

Parameters B: NetworkX graph

The input graph should be bipartite.

nodes: list or iterable

Nodes to project onto (the "bottom" nodes).

**Returns Graph**: NetworkX graph

A graph that is the projection onto the given nodes.

## See also:

```
is_bipartite, is_bipartite_node_set, sets, weighted_projected_graph,
overlap_weighted_projected_graph,
projected_graph
```

# **Notes**

No attempt is made to verify that the input graph B is bipartite. The graph and node properties are (shallow) copied to the projected graph.

#### References

[R186]

#### **Examples**

```
>>> from networkx.algorithms import bipartite
>>> B = nx.path_graph(5)
>>> B.add_edge(1,5)
>>> G = bipartite.collaboration_weighted_projected_graph(B, [0, 2, 4, 5])
>>> print(G.nodes())
[0, 2, 4, 5]
>>> for edge in G.edges(data=True): print(edge)
...
(0, 2, {'weight': 0.5})
(0, 5, {'weight': 0.5})
(2, 4, {'weight': 1.0})
(2, 5, {'weight': 0.5})
```

# overlap\_weighted\_projected\_graph

# overlap\_weighted\_projected\_graph(B, nodes, jaccard=True)

Overlap weighted projection of B onto one of its node sets.

The overlap weighted projection is the projection of the bipartite network B onto the specified nodes with weights representing the Jaccard index between the neighborhoods of the two nodes in the original bipartite network [R187]:

$$w_{v,u} = \frac{|N(u) \cap N(v)|}{|N(u) \cup N(v)|}$$

or if the parameter 'jaccard' is False, the fraction of common neighbors by minimum of both nodes degree in the original bipartite graph [R187]:

$$w_{v,u} = \frac{|N(u) \cap N(v)|}{\min(|N(u)|, |N(v)|)}$$

The nodes retain their attributes and are connected in the resulting graph if have an edge to a common node in the original bipartite graph.

Parameters B: NetworkX graph

The input graph should be bipartite.

nodes: list or iterable

Nodes to project onto (the "bottom" nodes).

jaccard: Bool (default=True)

Returns Graph: NetworkX graph

A graph that is the projection onto the given nodes.

## See also:

```
is_bipartite, is_bipartite_node_set, sets, weighted_projected_graph,
collaboration_weighted_projected_graph,
projected_graph
```

#### **Notes**

No attempt is made to verify that the input graph B is bipartite. The graph and node properties are (shallow) copied to the projected graph.

#### References

[R187]

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> B = nx.path_graph(5)
>>> G = bipartite.overlap_weighted_projected_graph(B, [0, 2, 4])
>>> print(G.nodes())
[0, 2, 4]
>>> print(G.edges(data=True))
[(0, 2, {'weight': 0.5}), (2, 4, {'weight': 0.5})]
>>> G = bipartite.overlap_weighted_projected_graph(B, [0, 2, 4], jaccard=False)
>>> print(G.edges(data=True))
[(0, 2, {'weight': 1.0}), (2, 4, {'weight': 1.0})]
```

## generic weighted projected graph

```
generic_weighted_projected_graph (B, nodes, weight_function=None)
```

Weighted projection of B with a user-specified weight function.

The bipartite network B is projected on to the specified nodes with weights computed by a user-specified function. This function must accept as a parameter the neighborhood sets of two nodes and return an integer or a float.

The nodes retain their attributes and are connected in the resulting graph if they have an edge to a common node in the original graph.

## **Parameters B**: NetworkX graph

The input graph should be bipartite.

nodes: list or iterable

Nodes to project onto (the "bottom" nodes).

#### weight\_function: function

This function must accept as parameters the same input graph that this function, and two nodes; and return an integer or a float. The default function computes the number of shared neighbors.

## Returns Graph: NetworkX graph

A graph that is the projection onto the given nodes.

#### See also:

```
is_bipartite, is_bipartite_node_set, sets, weighted_projected_graph,
collaboration_weighted_projected_graph, overlap_weighted_projected_graph,
projected_graph
```

#### **Notes**

No attempt is made to verify that the input graph B is bipartite. The graph and node properties are (shallow) copied to the projected graph.

#### **Examples**

```
>>> from networkx.algorithms import bipartite
>>> # Define some custom weight functions
>>> def jaccard(G, u, v):
       unbrs = set(G[u])
       vnbrs = set(G[v])
. . .
       return float (len (unbrs & vnbrs)) / len (unbrs | vnbrs)
>>> def my_weight(G, u, v, weight='weight'):
       w = 0
        for nbr in set(G[u]) & set(G[v]):
. . .
            w += G.edge[u][nbr].get(weight, 1) + G.edge[v][nbr].get(weight, 1)
        return w
>>> # A complete bipartite graph with 4 nodes and 4 edges
>>> B = nx.complete_bipartite_graph(2,2)
>>> # Add some arbitrary weight to the edges
>>> for i, (u,v) in enumerate(B.edges()):
       B.edge[u][v]['weight'] = i + 1
. . .
. . .
>>> for edge in B.edges(data=True):
       print (edge)
. . .
. . .
(0, 2, {'weight': 1})
(0, 3, {'weight': 2})
(1, 2, {'weight': 3})
(1, 3, {'weight': 4})
>>> # Without specifying a function, the weight is equal to # shared partners
>>> G = bipartite.generic_weighted_projected_graph(B, [0, 1])
>>> print (G.edges (data=True))
[(0, 1, {'weight': 2})]
>>> # To specify a custom weight function use the weight_function parameter
>>> G = bipartite.generic_weighted_projected_graph(B, [0, 1], weight_function=jaccard)
>>> print (G.edges (data=True))
[(0, 1, {'weight': 1.0})]
>>> G = bipartite.generic_weighted_projected_graph(B, [0, 1], weight_function=my_weight)
>>> print (G.edges (data=True))
[(0, 1, {'weight': 10})]
```

# 4.3.4 Spectral

Spectral bipartivity measure.

spectral\_bipartivity(G[, nodes, weight]) Returns the spectral bipartivity.

## spectral bipartivity

```
spectral_bipartivity(G, nodes=None, weight='weight')
```

Returns the spectral bipartivity.

Parameters G: NetworkX graph

**nodes**: list or container optional(default is all nodes)

Nodes to return value of spectral bipartivity contribution.

weight : string or None optional (default = 'weight')

Edge data key to use for edge weights. If None, weights set to 1.

Returns sb: float or dict

A single number if the keyword nodes is not specified, or a dictionary keyed by node with the spectral bipartivity contribution of that node as the value.

## See also:

color

## **Notes**

This implementation uses Numpy (dense) matrices which are not efficient for storing large sparse graphs.

#### References

[R190]

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4)
>>> bipartite.spectral_bipartivity(G)
1.0
```

# 4.3.5 Clustering

clustering(G[, nodes, mode])	Compute a bipartite clustering coefficient for nodes.
$average\_clustering(G[, nodes, mode])$	Compute the average bipartite clustering coefficient.
$latapy\_clustering(G[, nodes, mode])$	Compute a bipartite clustering coefficient for nodes.
robins_alexander_clustering(G)	Compute the bipartite clustering of G.

# clustering

```
clustering(G, nodes=None, mode='dot')
```

Compute a bipartite clustering coefficient for nodes.

The bipartie clustering coefficient is a measure of local density of connections defined as [R180]:

$$c_u = \frac{\sum_{v \in N(N(v))} c_{uv}}{|N(N(u))|}$$

where N(N(u)) are the second order neighbors of u in G excluding u, and  $c_{uv}$  is the pairwise clustering coefficient between nodes u and v.

The mode selects the function for  $c_{uv}$  which can be:

dot:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{|N(u) \cup N(v)|}$$

min:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{\min(|N(u)|, |N(v)|)}$$

max:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{max(|N(u)|, |N(v)|)}$$

**Parameters G**: graph

A bipartite graph

nodes: list or iterable (optional)

Compute bipartite clustering for these nodes. The default is all nodes in G.

mode: string

The pariwise bipartite clustering method to be used in the computation. It must be "dot", "max", or "min".

Returns clustering: dictionary

A dictionary keyed by node with the clustering coefficient value.

## See also:

```
robins_alexander_clustering, square_clustering, average_clustering
```

#### References

[R180]

#### **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4) # path graphs are bipartite
>>> c = bipartite.clustering(G)
>>> c[0]
0.5
>>> c = bipartite.clustering(G, mode='min')
>>> c[0]
1.0
```

## average clustering

average\_clustering(G, nodes=None, mode='dot')

Compute the average bipartite clustering coefficient.

A clustering coefficient for the whole graph is the average,

$$C = \frac{1}{n} \sum_{v \in G} c_v,$$

where n is the number of nodes in G.

Similar measures for the two bipartite sets can be defined [R179]

$$C_X = \frac{1}{|X|} \sum_{v \in X} c_v,$$

where X is a bipartite set of G.

Parameters G: graph

a bipartite graph

nodes: list or iterable, optional

A container of nodes to use in computing the average. The nodes should be either the entire graph (the default) or one of the bipartite sets.

mode: string

The pariwise bipartite clustering method. It must be "dot", "max", or "min"

Returns clustering: float

The average bipartite clustering for the given set of nodes or the entire graph if no nodes are specified.

# See also:

clustering

#### **Notes**

The container of nodes passed to this function must contain all of the nodes in one of the bipartite sets ("top" or "bottom") in order to compute the correct average bipartite clustering coefficients.

#### References

[R179]

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G=nx.star_graph(3) # star graphs are bipartite
>>> bipartite.average_clustering(G)
0.75
>>> X,Y=bipartite.sets(G)
>>> bipartite.average_clustering(G,X)
```

```
0.0
>>> bipartite.average_clustering(G,Y)
1.0
```

# latapy\_clustering

latapy\_clustering(G, nodes=None, mode='dot')

Compute a bipartite clustering coefficient for nodes.

The bipartie clustering coefficient is a measure of local density of connections defined as [R181]:

$$c_u = \frac{\sum_{v \in N(N(v))} c_{uv}}{|N(N(u))|}$$

where N(N(u)) are the second order neighbors of u in G excluding u, and  $c_{uv}$  is the pairwise clustering coefficient between nodes u and v.

The mode selects the function for  $c_{uv}$  which can be:

dot:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{|N(u) \cup N(v)|}$$

min:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{min(|N(u)|, |N(v)|)}$$

max:

$$c_{uv} = \frac{|N(u) \cap N(v)|}{max(|N(u)|, |N(v)|)}$$

Parameters G: graph

A bipartite graph

nodes: list or iterable (optional)

Compute bipartite clustering for these nodes. The default is all nodes in G.

mode: string

The pariwise bipartite clustering method to be used in the computation. It must be "dot", "max", or "min".

Returns clustering: dictionary

A dictionary keyed by node with the clustering coefficient value.

See also:

```
robins_alexander_clustering, square_clustering, average_clustering
```

References

[R181]

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.path_graph(4) # path graphs are bipartite
>>> c = bipartite.clustering(G)
>>> c[0]
0.5
>>> c = bipartite.clustering(G, mode='min')
>>> c[0]
1.0
```

# robins\_alexander\_clustering

## $robins\_alexander\_clustering(G)$

Compute the bipartite clustering of G.

Robins and Alexander [R182] defined bipartite clustering coefficient as four times the number of four cycles  $C_4$  divided by the number of three paths  $L_3$  in a bipartite graph:

$$CC_4 = \frac{4 * C_4}{L_3}$$

**Parameters G**: graph

a bipartite graph

Returns clustering: float

The Robins and Alexander bipartite clustering for the input graph.

#### See also:

latapy\_clustering, square\_clustering

#### References

[R182]

## **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.davis_southern_women_graph()
>>> print(round(bipartite.robins_alexander_clustering(G), 3))
0.468
```

# 4.3.6 Redundancy

Node redundancy for bipartite graphs.

node\_redundancy(G[, nodes]) Compute bipartite node redundancy coefficient.

# node\_redundancy

# node\_redundancy (G, nodes=None)

Compute bipartite node redundancy coefficient.

The redundancy coefficient of a node v is the fraction of pairs of neighbors of v that are both linked to other nodes. In a one-mode projection these nodes would be linked together even if v were not there.

$$rc(v) = \frac{|\{\{u, w\} \subseteq N(v), \ \exists v' \neq v, \ (v', u) \in E \text{ and } (v', w) \in E\}|}{\frac{|N(v)|(|N(v)|-1)}{2}}$$

where N(v) are the neighbors of v in G.

Parameters G: graph

A bipartite graph

nodes: list or iterable (optional)

Compute redundancy for these nodes. The default is all nodes in G.

Returns redundancy: dictionary

A dictionary keyed by node with the node redundancy value.

#### References

[R189]

# **Examples**

```
>>> from networkx.algorithms import bipartite
>>> G = nx.cycle_graph(4)
>>> rc = bipartite.node_redundancy(G)
>>> rc[0]
1.0
```

Compute the average redundancy for the graph:

```
>>> sum(rc.values())/len(G)
1.0
```

Compute the average redundancy for a set of nodes:

```
>>> nodes = [0, 2]
>>> sum(rc[n] for n in nodes)/len(nodes)
1.0
```

# 4.3.7 Centrality

<pre>closeness_centrality(G, nodes[, normalized])</pre>	Compute the closeness centrality for nodes in a bipartite network.
$degree\_centrality(G, nodes)$	Compute the degree centrality for nodes in a bipartite network.
betweenness_centrality(G, nodes)	Compute betweenness centrality for nodes in a bipartite network.

#### closeness centrality

## closeness\_centrality(G, nodes, normalized=True)

Compute the closeness centrality for nodes in a bipartite network.

The closeness of a node is the distance to all other nodes in the graph or in the case that the graph is not connected to all other nodes in the connected component containing that node.

**Parameters G**: graph

A bipartite network

nodes: list or container

Container with all nodes in one bipartite node set.

normalized: bool, optional

If True (default) normalize by connected component size.

Returns closeness: dictionary

Dictionary keyed by node with bipartite closeness centrality as the value.

#### See also:

betweenness\_centrality, degree\_centrality, sets, is\_bipartite

#### **Notes**

The nodes input parameter must conatin all nodes in one bipartite node set, but the dictionary returned contains all nodes from both node sets.

Closeness centrality is normalized by the minimum distance possible. In the bipartite case the minimum distance for a node in one bipartite node set is 1 from all nodes in the other node set and 2 from all other nodes in its own set [R177]. Thus the closeness centrality for node v in the two bipartite sets U with v nodes and v with v nodes is

$$c_v = \frac{m+2(n-1)}{d}$$
, for  $v \in U$ ,  $c_v = \frac{n+2(m-1)}{d}$ , for  $v \in V$ ,

where d is the sum of the distances from v to all other nodes.

Higher values of closeness indicate higher centrality.

As in the unipartite case, setting normalized=True causes the values to normalized further to n-1 / size(G)-1 where n is the number of nodes in the connected part of graph containing the node. If the graph is not completely connected, this algorithm computes the closeness centrality for each connected part separately.

# References

[R177]

# degree\_centrality

## degree\_centrality(G, nodes)

Compute the degree centrality for nodes in a bipartite network.

The degree centrality for a node v is the fraction of nodes connected to it.

Parameters G: graph

A bipartite network **nodes**: list or container

Container with all nodes in one bipartite node set.

Returns centrality: dictionary

Dictionary keyed by node with bipartite degree centrality as the value.

#### See also:

betweenness\_centrality, closeness\_centrality, sets, is\_bipartite

#### **Notes**

The nodes input parameter must conatin all nodes in one bipartite node set, but the dictionary returned contains all nodes from both bipartite node sets.

For unipartite networks, the degree centrality values are normalized by dividing by the maximum possible degree (which is n-1 where n is the number of nodes in G).

In the bipartite case, the maximum possible degree of a node in a bipartite node set is the number of nodes in the opposite node set [R178]. The degree centrality for a node v in the bipartite sets U with n nodes and V with m nodes is

$$d_v = \frac{deg(v)}{m}, \text{ for } v \in U,$$
  
 $d_v = \frac{deg(v)}{n}, \text{ for } v \in V,$ 

where deg(v) is the degree of node v.

## References

[R178]

## betweenness centrality

## $betweenness\_centrality(G, nodes)$

Compute betweenness centrality for nodes in a bipartite network.

Betweenness centrality of a node v is the sum of the fraction of all-pairs shortest paths that pass through v.

Values of betweenness are normalized by the maximum possible value which for bipartite graphs is limited by the relative size of the two node sets [R176].

Let n be the number of nodes in the node set U and m be the number of nodes in the node set V, then nodes in U are normalized by dividing by

$$\frac{1}{2}[m^2(s+1)^2 + m(s+1)(2t-s-1) - t(2s-t+3)],$$

where

$$s = (n-1) \div m, t = (n-1) \mod m,$$

and nodes in V are normalized by dividing by

$$\frac{1}{2}[n^2(p+1)^2 + n(p+1)(2r-p-1) - r(2p-r+3)],$$

where,

$$p = (m-1) \div n, r = (m-1) \mod n.$$

Parameters G: graph

A bipartite graph

nodes: list or container

Container with all nodes in one bipartite node set.

Returns betweenness: dictionary

Dictionary keyed by node with bipartite betweenness centrality as the value.

#### See also:

degree\_centrality, closeness\_centrality, sets, is\_bipartite

## **Notes**

The nodes input parameter must contain all nodes in one bipartite node set, but the dictionary returned contains all nodes from both node sets.

## References

[R176]

## 4.3.8 Generators

Generators and functions for bipartite graphs.

<pre>complete_bipartite_graph(n1, n2[, create_using])</pre>	Return the complete bipartite graph $K_{n_1,n_2}$ .
configuration_model(aseq, bseq[,])	Return a random bipartite graph from two given degree sequences.
havel_hakimi_graph(aseq, bseq[, create_using])	Return a bipartite graph from two given degree sequences using a Ha
reverse_havel_hakimi_graph(aseq, bseq[,])	Return a bipartite graph from two given degree sequences using a Ha
alternating_havel_hakimi_graph(aseq, bseq[,])	Return a bipartite graph from two given degree sequences using an al
<pre>preferential_attachment_graph(aseq, p[,])</pre>	Create a bipartite graph with a preferential attachment model from a
random_graph(n, m, p[, seed, directed])	Return a bipartite random graph.
gnmk_random_graph(n, m, k[, seed, directed])	Return a random bipartite graph G_{n,m,k}.

#### complete bipartite graph

```
complete_bipartite_graph (n1, n2, create_using=None)
```

Return the complete bipartite graph  $K_{n_1,n_2}$ .

Composed of two partitions with  $n_1$  nodes in the first and  $n_2$  nodes in the second. Each node in the first is connected to each node in the second.

Parameters n1: integer

Number of nodes for node set A.

n2: integer

Number of nodes for node set B.

create\_using: NetworkX graph instance, optional

Return graph of this type.

#### **Notes**

Node labels are the integers 0 to  $n_1 + n_2 - 1$ .

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

# configuration\_model

configuration\_model (aseq, bseq, create\_using=None, seed=None)

Return a random bipartite graph from two given degree sequences.

Parameters aseq: list

Degree sequence for node set A.

**bseq**: list

Degree sequence for node set B.

create\_using: NetworkX graph instance, optional

Return graph of this type.

seed: integer, optional

Seed for random number generator.

Nodes from the set A are connected to nodes in the set B by choosing randomly from the possible free stubs, one in A and one in B.

## **Notes**

The sum of the two sequences must be equal: sum(aseq)=sum(bseq) If no graph type is specified use MultiGraph with parallel edges. If you want a graph with no parallel edges use create\_using=Graph() but then the resulting degree sequences might not be exact.

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

# havel\_hakimi\_graph

## havel\_hakimi\_graph (aseq, bseq, create\_using=None)

Return a bipartite graph from two given degree sequences using a Havel-Hakimi style construction.

Nodes from the set A are connected to nodes in the set B by connecting the highest degree nodes in set A to the highest degree nodes in set B until all stubs are connected.

## Parameters aseq: list

Degree sequence for node set A.

**bseq**: list

Degree sequence for node set B.

create\_using: NetworkX graph instance, optional

Return graph of this type.

#### **Notes**

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

The sum of the two sequences must be equal: sum(aseq)=sum(bseq) If no graph type is specified use MultiGraph with parallel edges. If you want a graph with no parallel edges use create\_using=Graph() but then the resulting degree sequences might not be exact.

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

# reverse\_havel\_hakimi\_graph

## reverse\_havel\_hakimi\_graph (aseq, bseq, create\_using=None)

Return a bipartite graph from two given degree sequences using a Havel-Hakimi style construction.

Nodes from set A are connected to nodes in the set B by connecting the highest degree nodes in set A to the lowest degree nodes in set B until all stubs are connected.

#### Parameters aseq: list

Degree sequence for node set A.

bseq: list

Degree sequence for node set B.

create\_using: NetworkX graph instance, optional

Return graph of this type.

#### **Notes**

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

The sum of the two sequences must be equal: sum(aseq)=sum(bseq) If no graph type is specified use MultiGraph with parallel edges. If you want a graph with no parallel edges use create\_using=Graph() but then the resulting degree sequences might not be exact.

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

# alternating\_havel\_hakimi\_graph

# alternating\_havel\_hakimi\_graph (aseq, bseq, create\_using=None)

Return a bipartite graph from two given degree sequences using an alternating Havel-Hakimi style construction.

Nodes from the set A are connected to nodes in the set B by connecting the highest degree nodes in set A to alternatively the highest and the lowest degree nodes in set B until all stubs are connected.

# Parameters aseq: list

Degree sequence for node set A.

**bseq**: list

Degree sequence for node set B.

create\_using: NetworkX graph instance, optional

Return graph of this type.

# Notes

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

The sum of the two sequences must be equal: sum(aseq)=sum(bseq) If no graph type is specified use MultiGraph with parallel edges. If you want a graph with no parallel edges use create\_using=Graph() but then the resulting degree sequences might not be exact.

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

#### preferential attachment graph

# $\verb|preferential_attachment_graph| (aseq, p, create\_using=None, seed=None)|$

Create a bipartite graph with a preferential attachment model from a given single degree sequence.

#### Parameters aseq: list

Degree sequence for node set A.

p: float

Probability that a new bottom node is added.

create\_using: NetworkX graph instance, optional

Return graph of this type.

**seed**: integer, optional

Seed for random number generator.

#### **Notes**

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

#### References

[R183]

# random\_graph

```
random_graph (n, m, p, seed=None, directed=False)
```

Return a bipartite random graph.

This is a bipartite version of the binomial (Erdős-Rényi) graph.

## Parameters n: int

The number of nodes in the first bipartite set.

m: int

The number of nodes in the second bipartite set.

p: float

Probability for edge creation.

seed: int, optional

Seed for random number generator (default=None).

directed: bool, optional (default=False)

If True return a directed graph

# See also:

```
gnp_random_graph, configuration_model
```

#### **Notes**

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

The bipartite random graph algorithm chooses each of the n\*m (undirected) or 2\*nm (directed) possible edges with probability p.

This algorithm is O(n+m) where m is the expected number of edges.

The nodes are assigned the attribute 'bipartite' with the value 0 or 1 to indicate which bipartite set the node belongs to.

#### References

[R184]

# gnmk\_random\_graph

```
gnmk_random_graph(n, m, k, seed=None, directed=False)
```

Return a random bipartite graph  $G_{n,m,k}$ .

Produces a bipartite graph chosen randomly out of the set of all graphs with n top nodes, m bottom nodes, and k edges.

#### Parameters n: int

The number of nodes in the first bipartite set.

m: int

The number of nodes in the second bipartite set.

k: int

The number of edges

seed: int, optional

Seed for random number generator (default=None).

directed : bool, optional (default=False)

If True return a directed graph

## See also:

gnm\_random\_graph

# Notes

This function is not imported in the main namespace. To use it you have to explicitly import the bipartite package.

If k > m \* n then a complete bipartite graph is returned.

This graph is a bipartite version of the  $G_{nm}$  random graph model.

#### **Examples**

from networkx.algorithms import bipartite  $G = bipartite.gnmk_random_graph(10,20,50)$ 

# 4.4 Blockmodeling

Functions for creating network blockmodels from node partitions.

Created by Drew Conway <a href="mailto:created-by-copyright">drew.conway@nyu.edu</a> Copyright (c) 2010. All rights reserved.

blockmodel(G, partitions[, multigraph]) Returns a reduced graph constructed using the generalized block modeling technique.

4.4. Blockmodeling

# 4.4.1 blockmodel

**blockmodel** (*G*, partitions, multigraph=False)

Returns a reduced graph constructed using the generalized block modeling technique.

The blockmodel technique collapses nodes into blocks based on a given partitioning of the node set. Each partition of nodes (block) is represented as a single node in the reduced graph.

Edges between nodes in the block graph are added according to the edges in the original graph. If the parameter multigraph is False (the default) a single edge is added with a weight equal to the sum of the edge weights between nodes in the original graph The default is a weight of 1 if weights are not specified. If the parameter multigraph is True then multiple edges are added each with the edge data from the original graph.

## **Parameters G**: graph

A networkx Graph or DiGraph

partitions: list of lists, or list of sets

The partition of the nodes. Must be non-overlapping.

multigraph: bool, optional

If True return a MultiGraph with the edge data of the original graph applied to each corresponding edge in the new graph. If False return a Graph with the sum of the edge weights, or a count of the edges if the original graph is unweighted.

Returns blockmodel: a Networkx graph object

#### References

[R191]

# **Examples**

```
>>> G=nx.path_graph(6)
>>> partition=[[0,1],[2,3],[4,5]]
>>> M=nx.blockmodel(G,partition)
```

# 4.5 Boundary

Routines to find the boundary of a set of nodes.

Edge boundaries are edges that have only one end in the set of nodes.

Node boundaries are nodes outside the set of nodes that have an edge to a node in the set.

edge_boundary(G, nbunch1[, nbunch2])	Return the edge boundary.
<pre>node_boundary(G, nbunch1[, nbunch2])</pre>	Return the node boundary.

# 4.5.1 edge\_boundary

```
edge_boundary (G, nbunch1, nbunch2=None) Return the edge boundary.
```

Edge boundaries are edges that have only one end in the given set of nodes.

Parameters G: graph

A networkx graph

nbunch1: list, container

Interior node set

nbunch2: list, container

Exterior node set. If None then it is set to all of the nodes in G not in nbunch1.

Returns elist: list

List of edges

## **Notes**

Nodes in nbunch1 and nbunch2 that are not in G are ignored.

nbunch1 and nbunch2 are usually meant to be disjoint, but in the interest of speed and generality, that is not required here.

# 4.5.2 node boundary

node\_boundary (G, nbunch1, nbunch2=None)

Return the node boundary.

The node boundary is all nodes in the edge boundary of a given set of nodes that are in the set.

**Parameters G**: graph

A networkx graph

nbunch1: list, container

Interior node set

nbunch2: list, container

Exterior node set. If None then it is set to all of the nodes in G not in nbunch1.

Returns nlist: list

List of nodes.

#### **Notes**

Nodes in nbunch1 and nbunch2 that are not in G are ignored.

nbunch1 and nbunch2 are usually meant to be disjoint, but in the interest of speed and generality, that is not required here.

4.5. Boundary 165

# 4.6 Centrality

# 4.6.1 Degree

$degree\_centrality(G)$	Compute the degree centrality for nodes.
$in\_degree\_centrality(G)$	Compute the in-degree centrality for nodes.
<pre>out_degree_centrality(G)</pre>	Compute the out-degree centrality for nodes.

# degree\_centrality

# $degree\_centrality(G)$

Compute the degree centrality for nodes.

The degree centrality for a node v is the fraction of nodes it is connected to.

**Parameters G**: graph

A networkx graph

Returns nodes: dictionary

Dictionary of nodes with degree centrality as the value.

#### See also:

betweenness\_centrality, load\_centrality, eigenvector\_centrality

#### **Notes**

The degree centrality values are normalized by dividing by the maximum possible degree in a simple graph n-1 where n is the number of nodes in G.

For multigraphs or graphs with self loops the maximum degree might be higher than n-1 and values of degree centrality greater than 1 are possible.

# in\_degree\_centrality

#### in\_degree\_centrality(G)

Compute the in-degree centrality for nodes.

The in-degree centrality for a node v is the fraction of nodes its incoming edges are connected to.

Parameters G: graph

A NetworkX graph

Returns nodes: dictionary

Dictionary of nodes with in-degree centrality as values.

Raises NetworkXError

If the graph is undirected.

## See also:

degree\_centrality, out\_degree\_centrality

4.6. Centrality 167

#### **Notes**

The degree centrality values are normalized by dividing by the maximum possible degree in a simple graph n-1 where n is the number of nodes in G.

For multigraphs or graphs with self loops the maximum degree might be higher than n-1 and values of degree centrality greater than 1 are possible.

# out\_degree\_centrality

## $out\_degree\_centrality(G)$

Compute the out-degree centrality for nodes.

The out-degree centrality for a node v is the fraction of nodes its outgoing edges are connected to.

Parameters G: graph

A NetworkX graph

Returns nodes: dictionary

Dictionary of nodes with out-degree centrality as values.

Raises NetworkXError

If the graph is undirected.

#### See also:

degree\_centrality, in\_degree\_centrality

#### **Notes**

The degree centrality values are normalized by dividing by the maximum possible degree in a simple graph n-1 where n is the number of nodes in G.

For multigraphs or graphs with self loops the maximum degree might be higher than n-1 and values of degree centrality greater than 1 are possible.

# 4.6.2 Closeness

 $closeness\_centrality(G[, u, distance, ...])$  Compute closeness centrality for nodes.

# closeness\_centrality

 $closeness\_centrality(G, u=None, distance=None, normalized=True)$ 

Compute closeness centrality for nodes.

Closeness centrality [R197] of a node u is the reciprocal of the sum of the shortest path distances from u to all n-1 other nodes. Since the sum of distances depends on the number of nodes in the graph, closeness is normalized by the sum of minimum possible distances n-1.

$$C(u) = \frac{n-1}{\sum_{v=1}^{n-1} d(v, u)},$$

where d(v, u) is the shortest-path distance between v and u, and n is the number of nodes in the graph.

Notice that higher values of closeness indicate higher centrality.

**Parameters G**: graph

A NetworkX graph

u: node, optional

Return only the value for node u

**distance**: edge attribute key, optional (default=None)

Use the specified edge attribute as the edge distance in shortest path calculations

normalized: bool, optional

If True (default) normalize by the number of nodes in the connected part of the graph.

Returns nodes: dictionary

Dictionary of nodes with closeness centrality as the value.

## See also:

```
betweenness_centrality, load_centrality, eigenvector_centrality, degree centrality
```

#### **Notes**

The closeness centrality is normalized to (n-1)/(|G|-1) where n is the number of nodes in the connected part of graph containing the node. If the graph is not completely connected, this algorithm computes the closeness centrality for each connected part separately.

If the 'distance' keyword is set to an edge attribute key then the shortest-path length will be computed using Dijkstra's algorithm with that edge attribute as the edge weight.

#### References

[R197]

## 4.6.3 Betweenness

betweenness_centrality( $G[, k, normalized,]$ )	Compute the shortest-path betweenness centrality for nodes.
edge_betweenness_centrality( $G[, k,]$ )	Compute betweenness centrality for edges.

## betweenness\_centrality

**betweenness\_centrality** (G, k=None, normalized=True, weight=None, endpoints=False, seed=None) Compute the shortest-path betweenness centrality for nodes.

Betweenness centrality of a node v is the sum of the fraction of all-pairs shortest paths that pass through v:

$$c_B(v) = \sum_{s,t \in V} \frac{\sigma(s,t|v)}{\sigma(s,t)}$$

where V is the set of nodes,  $\sigma(s,t)$  is the number of shortest (s,t)-paths, and  $\sigma(s,t|v)$  is the number of those paths passing through some node v other than s,t. If s=t,  $\sigma(s,t)=1$ , and if  $v\in s,t$ ,  $\sigma(s,t|v)=0$  [R194].

4.6. Centrality 169

Parameters G: graph

A NetworkX graph

k: int, optional (default=None)

If k is not None use k node samples to estimate betweenness. The value of  $k \le n$  where n is the number of nodes in the graph. Higher values give better approximation.

normalized: bool, optional

If True the betweenness values are normalized by 2/((n-1)(n-2)) for graphs, and 1/((n-1)(n-2)) for directed graphs where n is the number of nodes in G.

weight: None or string, optional

If None, all edge weights are considered equal. Otherwise holds the name of the edge attribute used as weight.

endpoints: bool, optional

If True include the endpoints in the shortest path counts.

Returns nodes: dictionary

Dictionary of nodes with betweenness centrality as the value.

#### See also:

edge\_betweenness\_centrality, load\_centrality

#### **Notes**

The algorithm is from Ulrik Brandes [R193]. See [R196] for the original first published version and [R194] for details on algorithms for variations and related metrics.

For approximate betweenness calculations set k=#samples to use k nodes ("pivots") to estimate the betweenness values. For an estimate of the number of pivots needed see [R195].

For weighted graphs the edge weights must be greater than zero. Zero edge weights can produce an infinite number of equal length paths between pairs of nodes.

#### References

[R193], [R194], [R195], [R196]

#### edge betweenness centrality

Betweenness centrality of an edge e is the sum of the fraction of all-pairs shortest paths that pass through e:

$$c_B(v) = \sum_{s,t \in V} \frac{\sigma(s,t|e)}{\sigma(s,t)}$$

where V is the set of nodes, 'sigma(s, t)' is the number of shortest (s, t)-paths, and  $\sigma(s, t|e)$  is the number of those paths passing through edge e [R212].

Parameters G: graph

A NetworkX graph

**k**: int, optional (default=None)

If k is not None use k node samples to estimate betweenness. The value of  $k \le n$  where n is the number of nodes in the graph. Higher values give better approximation.

normalized: bool, optional

If True the betweenness values are normalized by 2/(n(n-1)) for graphs, and 1/(n(n-1)) for directed graphs where n is the number of nodes in G.

weight: None or string, optional

If None, all edge weights are considered equal. Otherwise holds the name of the edge attribute used as weight.

Returns edges: dictionary

Dictionary of edges with betweenness centrality as the value.

#### See also:

betweenness\_centrality, edge\_load

#### **Notes**

The algorithm is from Ulrik Brandes [R211].

For weighted graphs the edge weights must be greater than zero. Zero edge weights can produce an infinite number of equal length paths between pairs of nodes.

# References

[R211], [R212]

# 4.6.4 Current Flow Closeness

 $current_flow_closeness_centrality(G[,...])$  Compute current-flow closeness centrality for nodes.

# current\_flow\_closeness\_centrality

```
current_flow_closeness_centrality (G, weight='weight', dtype=< type 'float'>, solver='lu') Compute current-flow closeness centrality for nodes.
```

Current-flow closeness centrality is variant of closeness centrality based on effective resistance between nodes in a network. This metric is also known as information centrality.

**Parameters G**: graph

A NetworkX graph

dtype: data type (float)

Default data type for internal matrices. Set to np.float32 for lower memory consumption.

solver: string (default='lu')

4.6. Centrality 171

Type of linear solver to use for computing the flow matrix. Options are "full" (uses most memory), "lu" (recommended), and "cg" (uses least memory).

Returns nodes: dictionary

Dictionary of nodes with current flow closeness centrality as the value.

#### See also:

```
closeness_centrality
```

#### **Notes**

The algorithm is from Brandes [R208].

See also [R209] for the original definition of information centrality.

#### References

[R208], [R209]

# 4.6.5 Current-Flow Betweenness

$\verb current_flow_betweenness_centrality  (G[,])$	Compute current-flow betweenness centrality for nodes.
${\tt edge\_current\_flow\_betweenness\_centrality}(G)$	Compute current-flow betweenness centrality for edges.
approximate_current_flow_betweenness_centrality(G)	Compute the approximate current-flow betweenness cent

#### current flow betweenness centrality

Compute current-flow betweenness centrality for nodes.

Current-flow betweenness centrality uses an electrical current model for information spreading in contrast to betweenness centrality which uses shortest paths.

Current-flow betweenness centrality is also known as random-walk betweenness centrality [R207].

# Parameters G: graph

A NetworkX graph

**normalized**: bool, optional (default=True)

If True the betweenness values are normalized by 2/[(n-1)(n-2)] where n is the number of nodes in G.

weight : string or None, optional (default='weight')

Key for edge data used as the edge weight. If None, then use 1 as each edge weight.

## dtype: data type (float)

Default data type for internal matrices. Set to np.float32 for lower memory consumption

```
solver: string (default='lu')
```

Type of linear solver to use for computing the flow matrix. Options are "full" (uses most memory), "lu" (recommended), and "cg" (uses least memory).

Returns nodes: dictionary

Dictionary of nodes with betweenness centrality as the value.

#### See also:

```
approximate_current_flow_betweenness_centrality, betweenness_centrality,
edge_betweenness_centrality
```

#### **Notes**

Current-flow betweenness can be computed in  $O(I(n-1) + mn \log n)$  time [R206], where I(n-1) is the time needed to compute the inverse Laplacian. For a full matrix this is  $O(n^3)$  but using sparse methods you can achieve  $O(nm\sqrt{k})$  where k is the Laplacian matrix condition number.

The space required is O(nw) where w is the width of the sparse Laplacian matrix. Worse case is w=n for  $O(n^2)$ .

If the edges have a 'weight' attribute they will be used as weights in this algorithm. Unspecified weights are set to 1

#### References

[R206], [R207]

# edge\_current\_flow\_betweenness\_centrality

```
\begin{tabular}{ll} \textbf{edge\_current\_flow\_betweenness\_centrality} (G, & normalized=True, & weight='weight', \\ & dtype=<type 'float'>, solver='full') \end{tabular}
```

Compute current-flow betweenness centrality for edges.

Current-flow betweenness centrality uses an electrical current model for information spreading in contrast to betweenness centrality which uses shortest paths.

Current-flow betweenness centrality is also known as random-walk betweenness centrality [R214].

#### **Parameters G**: graph

A NetworkX graph

normalized : bool, optional (default=True)

If True the betweenness values are normalized by 2/[(n-1)(n-2)] where n is the number of nodes in G.

**weight**: string or None, optional (default='weight')

Key for edge data used as the edge weight. If None, then use 1 as each edge weight.

# dtype: data type (float)

Default data type for internal matrices. Set to np.float32 for lower memory consumption.

```
solver: string (default='lu')
```

Type of linear solver to use for computing the flow matrix. Options are "full" (uses most memory), "lu" (recommended), and "cg" (uses least memory).

4.6. Centrality 173

#### Returns nodes: dictionary

Dictionary of edge tuples with betweenness centrality as the value.

#### See also:

betweenness\_centrality,edge\_betweenness\_centrality,current\_flow\_betweenness\_centrality

#### **Notes**

Current-flow betweenness can be computed in  $O(I(n-1) + mn \log n)$  time [R213], where I(n-1) is the time needed to compute the inverse Laplacian. For a full matrix this is  $O(n^3)$  but using sparse methods you can achieve  $O(nm\sqrt{k})$  where k is the Laplacian matrix condition number.

The space required is O(nw)where'w is the width of the sparse Laplacian matrix. Worse case is w=n for  $O(n^2)$ .

If the edges have a 'weight' attribute they will be used as weights in this algorithm. Unspecified weights are set to 1.

#### References

[R213], [R214]

## approximate current flow betweenness centrality

Compute the approximate current-flow betweenness centrality for nodes.

Approximates the current-flow betweenness centrality within absolute error of epsilon with high probability [R192].

# Parameters G: graph

A NetworkX graph

**normalized**: bool, optional (default=True)

If True the betweenness values are normalized by 2/[(n-1)(n-2)] where n is the number of nodes in G.

weight : string or None, optional (default='weight')

Key for edge data used as the edge weight. If None, then use 1 as each edge weight.

#### dtype: data type (float)

Default data type for internal matrices. Set to np.float32 for lower memory consumption

# solver: string (default='lu')

Type of linear solver to use for computing the flow matrix. Options are "full" (uses most memory), "lu" (recommended), and "cg" (uses least memory).

# epsilon: float

Absolute error tolerance.

kmax: int

Maximum number of sample node pairs to use for approximation.

Returns nodes: dictionary

Dictionary of nodes with betweenness centrality as the value.

#### See also:

```
current flow betweenness centrality
```

### **Notes**

The running time is  $O((1/\epsilon^2)m\sqrt{k}\log n)$  and the space required is O(m) for n nodes and m edges.

If the edges have a 'weight' attribute they will be used as weights in this algorithm. Unspecified weights are set to 1.

### References

[R192]

# 4.6.6 Eigenvector

eigenvector_centrality( $G[$ , max_iter, tol,])	Compute the eigenvector centrality for the graph G.
eigenvector_centrality_numpy( $G[$ , weight])	Compute the eigenvector centrality for the graph G.
katz_centrality(G[, alpha, beta, max_iter,])	Compute the Katz centrality for the nodes of the graph G.
$katz\_centrality\_numpy(G[, alpha, beta,])$	Compute the Katz centrality for the graph G.

# eigenvector centrality

eigenvector\_centrality(G, max\_iter=100, tol=1e-06, nstart=None, weight='weight')

Compute the eigenvector centrality for the graph G.

Eigenvector centrality computes the centrality for a node based on the centrality of its neighbors. The eigenvector centrality for node i is

$$\mathbf{A}\mathbf{x} = \lambda\mathbf{x}$$

where A is the adjacency matrix of the graph G with eigenvalue  $\lambda$ . By virtue of the Perron–Frobenius theorem, there is a unique and positive solution if  $\lambda$  is the largest eigenvalue associated with the eigenvector of the adjacency matrix A ([R216]).

Parameters G: graph

A networkx graph

max\_iter: interger, optional

Maximum number of iterations in power method.

tol: float, optional

Error tolerance used to check convergence in power method iteration.

nstart: dictionary, optional

4.6. Centrality 175

Starting value of eigenvector iteration for each node.

weight: None or string, optional

If None, all edge weights are considered equal. Otherwise holds the name of the edge attribute used as weight.

Returns nodes: dictionary

Dictionary of nodes with eigenvector centrality as the value.

### See also:

```
eigenvector_centrality_numpy, pagerank, hits
```

#### **Notes**

The measure was introduced by [R215].

The eigenvector calculation is done by the power iteration method and has no guarantee of convergence. The iteration will stop after max\_iter iterations or an error tolerance of number\_of\_nodes (G) \*tol has been reached.

For directed graphs this is "left" eigenvector centrality which corresponds to the in-edges in the graph. For out-edges eigenvector centrality first reverse the graph with G.reverse().

### **Examples**

```
>>> G = nx.path_graph(4)
>>> centrality = nx.eigenvector_centrality(G)
>>> print(['%s %0.2f'%(node,centrality[node]) for node in centrality])
['0 0.37', '1 0.60', '2 0.60', '3 0.37']
```

# eigenvector\_centrality\_numpy

```
eigenvector_centrality_numpy(G, weight='weight')
```

Compute the eigenvector centrality for the graph G.

Eigenvector centrality computes the centrality for a node based on the centrality of its neighbors. The eigenvector centrality for node i is

$$\mathbf{A}\mathbf{x} = \lambda\mathbf{x}$$

where A is the adjacency matrix of the graph G with eigenvalue  $\lambda$ . By virtue of the Perron–Frobenius theorem, there is a unique and positive solution if  $\lambda$  is the largest eigenvalue associated with the eigenvector of the adjacency matrix A ([R218]).

Parameters G: graph

A networkx graph

weight: None or string, optional

The name of the edge attribute used as weight. If None, all edge weights are considered equal.

Returns nodes: dictionary

Dictionary of nodes with eigenvector centrality as the value.

### See also:

```
eigenvector_centrality, pagerank, hits
```

#### **Notes**

The measure was introduced by [R217].

This algorithm uses the SciPy sparse eigenvalue solver (ARPACK) to find the largest eigenvalue/eigenvector pair.

For directed graphs this is "left" eigenvector centrality which corresponds to the in-edges in the graph. For out-edges eigenvector centrality first reverse the graph with G.reverse().

### **Examples**

```
>>> G = nx.path_graph(4)
>>> centrality = nx.eigenvector_centrality_numpy(G)
>>> print(['%s %0.2f'%(node,centrality[node]) for node in centrality])
['0 0.37', '1 0.60', '2 0.60', '3 0.37']
```

# katz\_centrality

Compute the Katz centrality for the nodes of the graph G.

Katz centrality computes the centrality for a node based on the centrality of its neighbors. It is a generalization of the eigenvector centrality. The Katz centrality for node i is

$$x_i = \alpha \sum_j A_{ij} x_j + \beta,$$

where A is the adjacency matrix of the graph G with eigenvalues  $\lambda$ .

The parameter  $\beta$  controls the initial centrality and

$$\alpha < \frac{1}{\lambda_{max}}.$$

Katz centrality computes the relative influence of a node within a network by measuring the number of the immediate neighbors (first degree nodes) and also all other nodes in the network that connect to the node under consideration through these immediate neighbors.

Extra weight can be provided to immediate neighbors through the parameter  $\beta$ . Connections made with distant neighbors are, however, penalized by an attenuation factor  $\alpha$  which should be strictly less than the inverse largest eigenvalue of the adjacency matrix in order for the Katz centrality to be computed correctly. More information is provided in [R220] .

**Parameters G**: graph

A NetworkX graph

alpha: float

Attenuation factor

**beta**: scalar or dictionary, optional (default=1.0)

4.6. Centrality 177

Weight attributed to the immediate neighborhood. If not a scalar, the dictionary must have an value for every node.

max\_iter : integer, optional (default=1000)

Maximum number of iterations in power method.

tol: float, optional (default=1.0e-6)

Error tolerance used to check convergence in power method iteration.

nstart: dictionary, optional

Starting value of Katz iteration for each node.

**normalized**: bool, optional (default=True)

If True normalize the resulting values.

weight: None or string, optional

If None, all edge weights are considered equal. Otherwise holds the name of the edge attribute used as weight.

**Returns nodes**: dictionary

Dictionary of nodes with Katz centrality as the value.

#### Raises NetworkXError

If the parameter beta is not a scalar but lacks a value for at least one node

#### See also:

katz\_centrality\_numpy, eigenvector\_centrality, eigenvector\_centrality\_numpy,
pagerank, hits

# **Notes**

Katz centrality was introduced by [R221].

This algorithm it uses the power method to find the eigenvector corresponding to the largest eigenvalue of the adjacency matrix of G. The constant alpha should be strictly less than the inverse of largest eigenvalue of the adjacency matrix for the algorithm to converge. The iteration will stop after max\_iter iterations or an error tolerance of number of nodes(G)\*tol has been reached.

When  $\alpha = 1/\lambda_{max}$  and  $\beta = 0$ , Katz centrality is the same as eigenvector centrality.

For directed graphs this finds "left" eigenvectors which corresponds to the in-edges in the graph. For out-edges Katz centrality first reverse the graph with G.reverse().

### References

[R220], [R221]

# **Examples**

```
>>> import math
>>> G = nx.path_graph(4)
>>> phi = (1+math.sqrt(5))/2.0 # largest eigenvalue of adj matrix
>>> centrality = nx.katz_centrality(G,1/phi-0.01)
>>> for n,c in sorted(centrality.items()):
... print("%d %0.2f"%(n,c))
0 0.37
1 0.60
2 0.60
3 0.37
```

# katz\_centrality\_numpy

**katz\_centrality\_numpy** (G, alpha=0.1, beta=1.0, normalized=True, weight='weight') Compute the Katz centrality for the graph G.

Katz centrality computes the centrality for a node based on the centrality of its neighbors. It is a generalization of the eigenvector centrality. The Katz centrality for node i is

$$x_i = \alpha \sum_j A_{ij} x_j + \beta,$$

where A is the adjacency matrix of the graph G with eigenvalues  $\lambda$ .

The parameter  $\beta$  controls the initial centrality and

$$\alpha < \frac{1}{\lambda_{max}}.$$

Katz centrality computes the relative influence of a node within a network by measuring the number of the immediate neighbors (first degree nodes) and also all other nodes in the network that connect to the node under consideration through these immediate neighbors.

Extra weight can be provided to immediate neighbors through the parameter  $\beta$ . Connections made with distant neighbors are, however, penalized by an attenuation factor  $\alpha$  which should be strictly less than the inverse largest eigenvalue of the adjacency matrix in order for the Katz centrality to be computed correctly. More information is provided in [R222].

**Parameters G**: graph

A NetworkX graph

alpha: float

Attenuation factor

**beta**: scalar or dictionary, optional (default=1.0)

Weight attributed to the immediate neighborhood. If not a scalar the dictionary must have an value for every node.

normalized: bool

If True normalize the resulting values.

weight: None or string, optional

If None, all edge weights are considered equal. Otherwise holds the name of the edge attribute used as weight.

Returns nodes: dictionary

4.6. Centrality 179

Dictionary of nodes with Katz centrality as the value.

### Raises NetworkXError

If the parameter beta is not a scalar but lacks a value for at least one node

### See also:

```
katz_centrality, eigenvector_centrality_numpy, eigenvector_centrality,
pagerank, hits
```

### **Notes**

Katz centrality was introduced by [R223].

This algorithm uses a direct linear solver to solve the above equation. The constant alpha should be strictly less than the inverse of largest eigenvalue of the adjacency matrix for there to be a solution. When  $\alpha=1/\lambda_{max}$  and  $\beta=0$ , Katz centrality is the same as eigenvector centrality.

For directed graphs this finds "left" eigenvectors which corresponds to the in-edges in the graph. For out-edges Katz centrality first reverse the graph with G.reverse().

### References

```
[R222], [R223]
```

#### **Examples**

```
>>> import math
>>> G = nx.path_graph(4)
>>> phi = (1+math.sqrt(5))/2.0 # largest eigenvalue of adj matrix
>>> centrality = nx.katz_centrality_numpy(G,1/phi)
>>> for n,c in sorted(centrality.items()):
... print("%d %0.2f"%(n,c))
0 0.37
1 0.60
2 0.60
3 0.37
```

# 4.6.7 Communicability

$ ext{communicability}(G)$	Return communicability between all pairs of nodes in G.
extstyle  ext	Return communicability between all pairs of nodes in G.
communicability_centrality(G)	Return communicability centrality for each node in G.
$ ext{communicability\_centrality\_exp}(G)$	Return the communicability centrality for each node of G
communicability_betweenness_centrality( $G[,]$ )	Return communicability betweenness for all pairs of nodes in G.
$estrada\_index(G)$	Return the Estrada index of a the graph G.

### communicability

### communicability(G)

Return communicability between all pairs of nodes in G.

The communicability between pairs of nodes in G is the sum of closed walks of different lengths starting at node u and ending at node v.

# Parameters G: graph

Returns comm: dictionary of dictionaries

Dictionary of dictionaries keyed by nodes with communicability as the value.

### Raises NetworkXError

If the graph is not undirected and simple.

### See also:

**communicability\_centrality\_exp** Communicability centrality for each node of G using matrix exponential.

**communicability\_centrality** Communicability centrality for each node in G using spectral decomposition.

communicability Communicability between pairs of nodes in G.

### **Notes**

This algorithm uses a spectral decomposition of the adjacency matrix. Let G=(V,E) be a simple undirected graph. Using the connection between the powers of the adjacency matrix and the number of walks in the graph, the communicability between nodes u and v based on the graph spectrum is [R198]

$$C(u,v) = \sum_{j=1}^{n} \phi_j(u)\phi_j(v)e^{\lambda_j},$$

where  $\phi_j(u)$  is the uth element of the jth orthonormal eigenvector of the adjacency matrix associated with the eigenvalue  $\lambda_j$ .

### References

[R198]

# **Examples**

```
>>> G = nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> c = nx.communicability(G)
```

# communicability\_exp

### $communicability_exp(G)$

Return communicability between all pairs of nodes in G.

Communicability between pair of node (u,v) of node in G is the sum of closed walks of different lengths starting at node u and ending at node v.

# Parameters G: graph

Returns comm: dictionary of dictionaries

4.6. Centrality 181

Dictionary of dictionaries keyed by nodes with communicability as the value.

### Raises NetworkXError

If the graph is not undirected and simple.

### See also:

**communicability\_centrality\_exp** Communicability centrality for each node of G using matrix exponential.

**communicability\_centrality** Communicability centrality for each node in G using spectral decomposition.

**communicability\_exp** Communicability between all pairs of nodes in G using spectral decomposition.

### **Notes**

This algorithm uses matrix exponentiation of the adjacency matrix.

Let G=(V,E) be a simple undirected graph. Using the connection between the powers of the adjacency matrix and the number of walks in the graph, the communicability between nodes u and v is [R205],

$$C(u,v) = (e^A)_{uv},$$

where A is the adjacency matrix of G.

#### References

[R205]

# **Examples**

```
>>> G = nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> c = nx.communicability\_exp(G)
```

### communicability centrality

### $communicability\_centrality(G)$

Return communicability centrality for each node in G.

Communicability centrality, also called subgraph centrality, of a node n is the sum of closed walks of all lengths starting and ending at node n.

# Parameters G: graph

**Returns** nodes: dictionary

Dictionary of nodes with communicability centrality as the value.

# Raises NetworkXError

If the graph is not undirected and simple.

### See also:

**communicability** Communicability between all pairs of nodes in G.

communicability\_centrality Communicability centrality for each node of G.

### **Notes**

This version of the algorithm computes eigenvalues and eigenvectors of the adjacency matrix.

Communicability centrality of a node u in G can be found using a spectral decomposition of the adjacency matrix [R201] [R202],

$$SC(u) = \sum_{j=1}^{N} (v_j^u)^2 e^{\lambda_j},$$

where  $v_j$  is an eigenvector of the adjacency matrix A of G corresponding corresponding to the eigenvalue  $\lambda_j$ .

#### References

[R201], [R202]

# **Examples**

```
>>> G = nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> sc = nx.communicability centrality(G)
```

# communicability\_centrality\_exp

# $communicability\_centrality\_exp(G)$

Return the communicability centrality for each node of G

Communicability centrality, also called subgraph centrality, of a node n is the sum of closed walks of all lengths starting and ending at node n.

### Parameters G: graph

**Returns** nodes:dictionary

Dictionary of nodes with communicability centrality as the value.

### Raises NetworkXError

If the graph is not undirected and simple.

### See also:

communicability Communicability between all pairs of nodes in G.

communicability\_centrality Communicability centrality for each node of G.

#### **Notes**

This version of the algorithm exponentiates the adjacency matrix. The communicability centrality of a node u in G can be found using the matrix exponential of the adjacency matrix of G [R203] [R204],

$$SC(u) = (e^A)_{uu}.$$

4.6. Centrality 183

#### References

[R203], [R204]

### **Examples**

```
>>> G = nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> sc = nx.communicability_centrality_exp(G)
```

### communicability betweenness centrality

### communicability\_betweenness\_centrality(G, normalized=True)

Return communicability betweenness for all pairs of nodes in G.

Communicability betweenness measure makes use of the number of walks connecting every pair of nodes as the basis of a betweenness centrality measure.

# Parameters G: graph

**Returns** nodes:dictionary

Dictionary of nodes with communicability betweenness as the value.

#### Raises NetworkXError

If the graph is not undirected and simple.

### See also:

communicability Communicability between all pairs of nodes in G.

**communicability\_centrality** Communicability centrality for each node of G using matrix exponential.

**communicability\_centrality\_exp** Communicability centrality for each node in G using spectral decomposition.

# Notes

Let G=(V,E) be a simple undirected graph with n nodes and m edges, and A denote the adjacency matrix of G

Let G(r) = (V, E(r)) be the graph resulting from removing all edges connected to node r but not the node itself.

The adjacency matrix for G(r) is A + E(r), where E(r) has nonzeros only in row and column r.

The communicability betweenness of a node r is [R200]

$$\omega_r = \frac{1}{C} \sum_{p} \sum_{q} \frac{G_{prq}}{G_{pq}}, p \neq q, q \neq r,$$

where  $G_{prq} = (e_{pq}^A - (e^{A+E(r)})_{pq})$  is the number of walks involving node r,  $G_{pq} = (e^A)_{pq}$  is the number of closed walks starting at node p and ending at node q, and  $C = (n-1)^2 - (n-1)$  is a normalization factor equal to the number of terms in the sum.

The resulting  $\omega_r$  takes values between zero and one. The lower bound cannot be attained for a connected graph, and the upper bound is attained in the star graph.

### References

[R200]

# **Examples**

```
>>> G = nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> cbc = nx.communicability_betweenness_centrality(G)
```

### estrada index

### estrada index(G)

Return the Estrada index of a the graph G.

Parameters G: graph

Returns estrada index: float

Raises NetworkXError

If the graph is not undirected and simple.

### See also:

estrada\_index\_exp

### **Notes**

Let G = (V, E) be a simple undirected graph with n nodes and let  $\lambda_1 \leq \lambda_2 \leq \cdots \leq \lambda_n$  be a non-increasing ordering of the eigenvalues of its adjacency matrix A. The Estrada index is

$$EE(G) = \sum_{j=1}^{n} e^{\lambda_j}.$$

### References

[R219]

# **Examples**

```
>>> G=nx.Graph([(0,1),(1,2),(1,5),(5,4),(2,4),(2,3),(4,3),(3,6)])
>>> ei=nx.estrada_index(G)
```

# 4.6.8 Load

<pre>load_centrality(G[, v, cutoff, normalized,])</pre>	Compute load centrality for nodes.
$edge\_load(G[, nodes, cutoff])$	Compute edge load.

4.6. Centrality 185

## load centrality

load\_centrality(G, v=None, cutoff=None, normalized=True, weight=None)

Compute load centrality for nodes.

The load centrality of a node is the fraction of all shortest paths that pass through that node.

Parameters G: graph

A networkx graph

normalized: bool, optional

If True the betweenness values are normalized by b=b/(n-1)(n-2) where n is the number of nodes in G.

weight: None or string, optional

If None, edge weights are ignored. Otherwise holds the name of the edge attribute used as weight.

cutoff: bool, optional

If specified, only consider paths of length <= cutoff.

Returns nodes: dictionary

Dictionary of nodes with centrality as the value.

#### See also:

betweenness\_centrality

### **Notes**

Load centrality is slightly different than betweenness. It was originally introduced by [R225]. For this load algorithm see [R224].

### References

[R224], [R225]

### edge load

edge\_load (G, nodes=None, cutoff=False)

Compute edge load.

WARNING:

This module is for demonstration and testing purposes.

# 4.6.9 Dispersion

dispersion(G[, u, v, normalized, alpha, b, c]) Calculate dispersion between u and v in G.

## dispersion

dispersion(G, u=None, v=None, normalized=True, alpha=1.0, b=0.0, c=0.0)

Calculate dispersion between u and v in G.

A link between two actors (u and v) has a high dispersion when their mutual ties (s and t) are not well connected with each other.

**Parameters G**: graph

A NetworkX graph.

u: node, optional

The source for the dispersion score (e.g. ego node of the network).

v: node, optional

The target of the dispersion score if specified.

normalized: bool

If True (default) normalize by the embeddeness of the nodes (u and v).

Returns nodes: dictionary

If u (v) is specified, returns a dictionary of nodes with dispersion score for all "target" ("source") nodes. If neither u nor v is specified, returns a dictionary of dictionaries for all nodes 'u' in the graph with a dispersion score for each node 'v'.

### **Notes**

This implementation follows Lars Backstrom and Jon Kleinberg [R210]. Typical usage would be to run dispersion on the ego network  $G_u$  if u were specified. Running dispersion () with neither u nor v specified can take some time to complete.

### References

[R210]

# 4.7 Chordal

Algorithms for chordal graphs.

A graph is chordal if every cycle of length at least 4 has a chord (an edge joining two nodes not adjacent in the cycle). http://en.wikipedia.org/wiki/Chordal\_graph

is_chordal(G)	Checks whether G is a chordal graph.
chordal_graph_cliques(G)	Returns the set of maximal cliques of a chordal graph.
${ t chordal\_graph\_treewidth}(G)$	Returns the treewidth of the chordal graph G.
<pre>find_induced_nodes(G, s, t[, treewidth_bound])</pre>	Returns the set of induced nodes in the path from s to t.

4.7. Chordal 187

# 4.7.1 is chordal

# $is\_chordal(G)$

Checks whether G is a chordal graph.

A graph is chordal if every cycle of length at least 4 has a chord (an edge joining two nodes not adjacent in the cycle).

Parameters G: graph

A NetworkX graph.

Returns chordal: bool

True if G is a chordal graph and False otherwise.

#### Raises NetworkXError

The algorithm does not support DiGraph, MultiGraph and MultiDiGraph. If the input graph is an instance of one of these classes, a NetworkXError is raised.

#### **Notes**

The routine tries to go through every node following maximum cardinality search. It returns False when it finds that the separator for any node is not a clique. Based on the algorithms in [R228].

#### References

[R228]

### **Examples**

```
>>> import networkx as nx
>>> e=[(1,2),(1,3),(2,3),(2,4),(3,4),(3,5),(3,6),(4,5),(4,6),(5,6)]
>>> G=nx.Graph(e)
>>> nx.is_chordal(G)
True
```

# 4.7.2 chordal graph cliques

# $chordal\_graph\_cliques(G)$

Returns the set of maximal cliques of a chordal graph.

The algorithm breaks the graph in connected components and performs a maximum cardinality search in each component to get the cliques.

Parameters G: graph

A NetworkX graph

**Returns** cliques: A set containing the maximal cliques in G.

Raises NetworkXError

The algorithm does not support DiGraph, MultiGraph and MultiDiGraph. If the input graph is an instance of one of these classes, a NetworkXError is raised. The algorithm can only be applied to chordal graphs. If the input graph is found to be non-chordal, a NetworkXError is raised.

### **Examples**

```
>>> import networkx as nx
>>> e= [(1,2),(1,3),(2,3),(2,4),(3,4),(3,5),(3,6),(4,5),(4,6),(5,6),(7,8)]
>>> G = nx.Graph(e)
>>> G.add_node(9)
>>> setlist = nx.chordal_graph_cliques(G)
```

# 4.7.3 chordal graph treewidth

# ${\tt chordal\_graph\_treewidth}\,(G)$

Returns the treewidth of the chordal graph G.

Parameters G: graph

A NetworkX graph

Returns treewidth: int

The size of the largest clique in the graph minus one.

### Raises NetworkXError

The algorithm does not support DiGraph, MultiGraph and MultiDiGraph. If the input graph is an instance of one of these classes, a NetworkXError is raised. The algorithm can only be applied to chordal graphs. If the input graph is found to be non-chordal, a NetworkXError is raised.

### References

[R226]

# **Examples**

```
>>> import networkx as nx
>>> e = [(1,2),(1,3),(2,3),(2,4),(3,4),(3,5),(3,6),(4,5),(4,6),(5,6),(7,8)]
>>> G = nx.Graph(e)
>>> G.add_node(9)
>>> nx.chordal_graph_treewidth(G)
3
```

# 4.7.4 find induced nodes

```
find_induced_nodes (G, s, t, treewidth_bound=9223372036854775807)
```

Returns the set of induced nodes in the path from s to t.

Parameters G: graph

A chordal NetworkX graph

4.7. Chordal 189

#### s: node

Source node to look for induced nodes

### t: node

Destination node to look for induced nodes

### treewith bound: float

Maximum treewidth acceptable for the graph H. The search for induced nodes will end as soon as the treewidth\_bound is exceeded.

### Returns I: Set of nodes

The set of induced nodes in the path from s to t in G

### Raises NetworkXError

The algorithm does not support DiGraph, MultiGraph and MultiDiGraph. If the input graph is an instance of one of these classes, a NetworkXError is raised. The algorithm can only be applied to chordal graphs. If the input graph is found to be non-chordal, a NetworkXError is raised.

### **Notes**

G must be a chordal graph and (s,t) an edge that is not in G.

If a treewidth\_bound is provided, the search for induced nodes will end as soon as the treewidth\_bound is exceeded.

The algorithm is inspired by Algorithm 4 in [R227]. A formal definition of induced node can also be found on that reference.

### References

[R227]

# **Examples**

```
>>> import networkx as nx
>>> G=nx.Graph()
>>> G = nx.generators.classic.path_graph(10)
>>> I = nx.find_induced_nodes(G,1,9,2)
>>> list(I)
[1, 2, 3, 4, 5, 6, 7, 8, 9]
```

# 4.8 Clique

Find and manipulate cliques of graphs.

Note that finding the largest clique of a graph has been shown to be an NP-complete problem; the algorithms here could take a long time to run.

http://en.wikipedia.org/wiki/Clique\_problem

enumerate_all_cliques(G)	Returns all cliques in an undirected graph.
$find\_cliques(G)$	Search for all maximal cliques in a graph.
$make\_max\_clique\_graph(G[, create\_using, name])$	Create the maximal clique graph of a graph.
$make\_clique\_bipartite(G[, fpos,])$	Create a bipartite clique graph from a graph G.
$graph\_clique\_number(G[, cliques])$	Return the clique number (size of the largest clique) for G.
<pre>graph_number_of_cliques(G[, cliques])</pre>	Returns the number of maximal cliques in G.
<pre>node_clique_number(G[, nodes, cliques])</pre>	Returns the size of the largest maximal clique containing each given node
<pre>number_of_cliques(G[, nodes, cliques])</pre>	Returns the number of maximal cliques for each node.
cliques_containing_node(G[, nodes, cliques])	Returns a list of cliques containing the given node.

# 4.8.1 enumerate\_all\_cliques

### $enumerate\_all\_cliques(G)$

Returns all cliques in an undirected graph.

This method returns cliques of size (cardinality) k = 1, 2, 3, ..., maxDegree - 1.

Where maxDegree is the maximal degree of any node in the graph.

### Parameters G: undirected graph

**Returns** generator of lists: generator of list for each clique.

### **Notes**

To obtain a list of all cliques, use list (enumerate\_all\_cliques(G)).

Based on the algorithm published by Zhang et al. (2005) [R229] and adapted to output all cliques discovered.

This algorithm is not applicable on directed graphs.

This algorithm ignores self-loops and parallel edges as clique is not conventionally defined with such edges.

There are often many cliques in graphs. This algorithm however, hopefully, does not run out of memory since it only keeps candidate sublists in memory and continuously removes exhausted sublists.

### References

[R229]

# 4.8.2 find\_cliques

### $find_cliques(G)$

Search for all maximal cliques in a graph.

Maximal cliques are the largest complete subgraph containing a given node. The largest maximal clique is sometimes called the maximum clique.

Returns generator of lists: genetor of member list for each maximal clique

#### See also:

find\_cliques\_recursive, A

4.8. Clique 191

#### **Notes**

To obtain a list of cliques, use list(find\_cliques(G)).

Based on the algorithm published by Bron & Kerbosch (1973) [R230] as adapted by Tomita, Tanaka and Takahashi (2006) [R231] and discussed in Cazals and Karande (2008) [R232]. The method essentially unrolls the recursion used in the references to avoid issues of recursion stack depth.

This algorithm is not suitable for directed graphs.

This algorithm ignores self-loops and parallel edges as clique is not conventionally defined with such edges.

There are often many cliques in graphs. This algorithm can run out of memory for large graphs.

#### References

[R230], [R231], [R232]

# 4.8.3 make max clique graph

make\_max\_clique\_graph (G, create\_using=None, name=None)

Create the maximal clique graph of a graph.

Finds the maximal cliques and treats these as nodes. The nodes are connected if they have common members in the original graph. Theory has done a lot with clique graphs, but I haven't seen much on maximal clique graphs.

### **Notes**

This should be the same as make\_clique\_bipartite followed by project\_up, but it saves all the intermediate steps.

# 4.8.4 make clique bipartite

make\_clique\_bipartite(G, fpos=None, create\_using=None, name=None)

Create a bipartite clique graph from a graph G.

Nodes of G are retained as the "bottom nodes" of B and cliques of G become "top nodes" of B. Edges are present if a bottom node belongs to the clique represented by the top node.

Returns a Graph with additional attribute dict B.node\_type which is keyed by nodes to "Bottom" or "Top" appropriately.

if fpos is not None, a second additional attribute dict B.pos is created to hold the position tuple of each node for viewing the bipartite graph.

# 4.8.5 graph clique number

# ${\tt graph\_clique\_number}\,(\textit{G},\textit{cliques=None})$

Return the clique number (size of the largest clique) for G.

An optional list of cliques can be input if already computed.

# 4.8.6 graph\_number\_of\_cliques

# graph\_number\_of\_cliques(G, cliques=None)

Returns the number of maximal cliques in G.

An optional list of cliques can be input if already computed.

# 4.8.7 node clique number

# node\_clique\_number(G, nodes=None, cliques=None)

Returns the size of the largest maximal clique containing each given node.

Returns a single or list depending on input nodes. Optional list of cliques can be input if already computed.

# 4.8.8 number of cliques

# number\_of\_cliques (G, nodes=None, cliques=None)

Returns the number of maximal cliques for each node.

Returns a single or list depending on input nodes. Optional list of cliques can be input if already computed.

# 4.8.9 cliques containing node

# $\verb|cliques_containing_node| (G, nodes=None, cliques=None)|$

Returns a list of cliques containing the given node.

Returns a single list or list of lists depending on input nodes. Optional list of cliques can be input if already computed.

# 4.9 Clustering

Algorithms to characterize the number of triangles in a graph.

triangles(G[, nodes])	Compute the number of triangles.
transitivity(G)	Compute graph transitivity, the fraction of all possible triangles present in G.
<pre>clustering(G[, nodes, weight])</pre>	Compute the clustering coefficient for nodes.
average_clustering(G[, nodes, weight,])	Compute the average clustering coefficient for the graph G.
square_clustering(G[, nodes])	Compute the squares clustering coefficient for nodes.

# 4.9.1 triangles

# triangles (G, nodes=None)

Compute the number of triangles.

Finds the number of triangles that include a node as one vertex.

Parameters G: graph

A networkx graph

**nodes**: container of nodes, optional (default= all nodes in G)

Compute triangles for nodes in this container.

4.9. Clustering

## Returns out: dictionary

Number of triangles keyed by node label.

### **Notes**

When computing triangles for the entire graph each triangle is counted three times, once at each node. Self loops are ignored.

### **Examples**

```
>>> G=nx.complete_graph(5)
>>> print(nx.triangles(G,0))
6
>>> print(nx.triangles(G))
{0: 6, 1: 6, 2: 6, 3: 6, 4: 6}
>>> print(list(nx.triangles(G,(0,1)).values()))
[6, 6]
```

# 4.9.2 transitivity

### transitivity(G)

Compute graph transitivity, the fraction of all possible triangles present in G.

Possible triangles are identified by the number of "triads" (two edges with a shared vertex).

The transitivity is

$$T=3\frac{\#triangles}{\#triads}.$$

Parameters G: graph

Returns out: float

Transitivity

# **Examples**

```
>>> G = nx.complete_graph(5)
>>> print(nx.transitivity(G))
1.0
```

# 4.9.3 clustering

clustering(G, nodes=None, weight=None)

Compute the clustering coefficient for nodes.

For unweighted graphs, the clustering of a node u is the fraction of possible triangles through that node that exist,

$$c_u = \frac{2T(u)}{deg(u)(deg(u) - 1)},$$

where T(u) is the number of triangles through node u and deg(u) is the degree of u.

For weighted graphs, the clustering is defined as the geometric average of the subgraph edge weights [R235],

$$c_u = \frac{1}{deg(u)(deg(u) - 1)} \sum_{uv} (\hat{w}_{uv} \hat{w}_{uw} \hat{w}_{vw})^{1/3}.$$

The edge weights  $\hat{w}_{uv}$  are normalized by the maximum weight in the network  $\hat{w}_{uv} = w_{uv}/\max(w)$ .

The value of  $c_u$  is assigned to 0 if deg(u) < 2.

Parameters G: graph

**nodes**: container of nodes, optional (default=all nodes in G)

Compute clustering for nodes in this container.

weight: string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

Returns out: float, or dictionary

Clustering coefficient at specified nodes

### **Notes**

Self loops are ignored.

### References

[R235]

### **Examples**

```
>>> G=nx.complete_graph(5)
>>> print(nx.clustering(G,0))
1.0
>>> print(nx.clustering(G))
{0: 1.0, 1: 1.0, 2: 1.0, 3: 1.0, 4: 1.0}
```

# 4.9.4 average clustering

 $\textbf{average\_clustering} \ (G, \textit{nodes=None}, \textit{weight=None}, \textit{count\_zeros=True})$ 

Compute the average clustering coefficient for the graph G.

The clustering coefficient for the graph is the average,

$$C = \frac{1}{n} \sum_{v \in G} c_v,$$

where n is the number of nodes in G.

Parameters G: graph

nodes: container of nodes, optional (default=all nodes in G)

4.9. Clustering 195

Compute average clustering for nodes in this container.

weight : string or None, optional (default=None)

The edge attribute that holds the numerical value used as a weight. If None, then each edge has weight 1.

count\_zeros: bool

If False include only the nodes with nonzero clustering in the average.

Returns avg: float

Average clustering

#### **Notes**

This is a space saving routine; it might be faster to use the clustering function to get a list and then take the average.

Self loops are ignored.

#### References

[R233], [R234]

# **Examples**

```
>>> G=nx.complete_graph(5)
>>> print(nx.average_clustering(G))
1.0
```

# 4.9.5 square\_clustering

square\_clustering(G, nodes=None)

Compute the squares clustering coefficient for nodes.

For each node return the fraction of possible squares that exist at the node [R236]

$$C_4(v) = \frac{\sum_{u=1}^{k_v} \sum_{w=u+1}^{k_v} q_v(u, w)}{\sum_{u=1}^{k_v} \sum_{w=u+1}^{k_v} [a_v(u, w) + q_v(u, w)]},$$

where  $q_v(u, w)$  are the number of common neighbors of u and w other than v (ie squares), and  $a_v(u, w) = (k_u - (1 + q_v(u, w) + \theta_{uv}))(k_w - (1 + q_v(u, w) + \theta_{uw}))$ , where  $\theta_{uw} = 1$  if u and w are connected and 0 otherwise.

Parameters G: graph

**nodes**: container of nodes, optional (default=all nodes in G)

Compute clustering for nodes in this container.

Returns c4: dictionary

A dictionary keyed by node with the square clustering coefficient value.

### **Notes**

While  $C_3(v)$  (triangle clustering) gives the probability that two neighbors of node v are connected with each other,  $C_4(v)$  is the probability that two neighbors of node v share a common neighbor different from v. This algorithm can be applied to both bipartite and unipartite networks.

### References

[R236]

# **Examples**

```
>>> G=nx.complete_graph(5)
>>> print(nx.square_clustering(G,0))
1.0
>>> print(nx.square_clustering(G))
{0: 1.0, 1: 1.0, 2: 1.0, 3: 1.0, 4: 1.0}
```

# 4.10 Coloring

greedy\_color(G[, strategy, interchange]) Color a graph using various strategies of greedy graph coloring.

# 4.10.1 greedy\_color

**greedy\_color** (*G*, *strategy=*<*function strategy\_largest\_first at 0x2ada9b0*>, *interchange=False*)

Color a graph using various strategies of greedy graph coloring. The strategies are described in [R237].

Attempts to color a graph using as few colors as possible, where no neighbours of a node can have same color as the node itself.

# Parameters G: NetworkX graph

strategy : function(G, colors)

A function that provides the coloring strategy, by returning nodes in the ordering they should be colored. G is the graph, and colors is a dict of the currently assigned colors, keyed by nodes.

You can pass your own ordering function, or use one of the built in:

- · strategy\_largest\_first
- strategy\_random\_sequential
- strategy\_smallest\_last
- strategy\_independent\_set
- strategy\_connected\_sequential\_bfs
- strategy\_connected\_sequential\_dfs
- strategy\_connected\_sequential (alias of strategy\_connected\_sequential\_bfs)
- strategy\_saturation\_largest\_first (also known as DSATUR)

4.10. Coloring 197

### interchange: bool

Will use the color interchange algorithm described by [R238] if set to true.

Note that saturation largest first and independent set do not work with interchange. Furthermore, if you use interchange with your own strategy function, you cannot rely on the values in the colors argument.

**Returns** A dictionary with keys representing nodes and values representing corresponding coloring.

#### References

[R237], [R238]

### **Examples**

```
>>> G = nx.cycle_graph(4)
>>> d = nx.coloring.greedy_color(G, strategy=nx.coloring.strategy_largest_first)
>>> d in [{0: 0, 1: 1, 2: 0, 3: 1}, {0: 1, 1: 0, 2: 1, 3: 0}]
True
```

# 4.11 Communities

# 4.11.1 K-Clique

k\_clique\_communities(G, k[, cliques]) Find k-clique communities in graph using the percolation method.

### k clique communities

# $k\_clique\_communities(G, k, cliques=None)$

Find k-clique communities in graph using the percolation method.

A k-clique community is the union of all cliques of size k that can be reached through adjacent (sharing k-1 nodes) k-cliques.

```
Parameters G: NetworkX graph
```

k: int

Size of smallest clique

# cliques: list or generator

Precomputed cliques (use networkx.find\_cliques(G))

**Returns** Yields sets of nodes, one for each k-clique community.

#### References

[R239]

### **Examples**

```
>>> G = nx.complete_graph(5)
>>> K5 = nx.convert_node_labels_to_integers(G, first_label=2)
>>> G.add_edges_from(K5.edges())
>>> c = list(nx.k_clique_communities(G, 4))
>>> list(c[0])
[0, 1, 2, 3, 4, 5, 6]
>>> list(nx.k_clique_communities(G, 6))
[]
```

# 4.12 Components

# 4.12.1 Connectivity

Connected components.

$is\_connected(G)$	Return True if the graph is connected, false otherwise.
$number\_connected\_components(G)$	Return the number of connected components.
$ ext{connected\_components}(G)$	Generate connected components.
$connected\_component\_subgraphs(G[, copy])$	Generate connected components as subgraphs.
node_connected_component(G, n)	Return the nodes in the component of graph containing node n.

# is\_connected

# $is\_connected(G)$

Return True if the graph is connected, false otherwise.

Parameters G: NetworkX Graph

An undirected graph.

Returns connected: bool

True if the graph is connected, false otherwise.

### See also:

```
connected_components
```

# **Notes**

For undirected graphs only.

### **Examples**

```
>>> G = nx.path_graph(4)
>>> print(nx.is_connected(G))
True
```

4.12. Components

# number\_connected\_components

# $number\_connected\_components(G)$

Return the number of connected components.

Parameters G: NetworkX graph

An undirected graph.

**Returns** n: integer

Number of connected components

### See also:

```
connected_components
```

### **Notes**

For undirected graphs only.

# connected\_components

### $connected\_components(G)$

Generate connected components.

Parameters G: NetworkX graph

An undirected graph

**Returns** comp: generator of lists

A list of nodes for each component of G.

# See also:

```
strongly_connected_components
```

# **Notes**

For undirected graphs only.

# **Examples**

Generate a sorted list of connected components, largest first.

```
>>> G = nx.path_graph(4)
>>> G.add_path([10, 11, 12])
>>> sorted(nx.connected_components(G), key = len, reverse=True)
[[0, 1, 2, 3], [10, 11, 12]]
```

# connected component subgraphs

# connected\_component\_subgraphs (G, copy=True)

Generate connected components as subgraphs.

Parameters G: NetworkX graph

An undirected graph.

### copy: bool (default=True)

If True make a copy of the graph attributes

Returns comp: generator

A generator of graphs, one for each connected component of G.

### See also:

```
connected_components
```

#### **Notes**

For undirected graphs only. Graph, node, and edge attributes are copied to the subgraphs by default.

# **Examples**

```
>>> G = nx.path_graph(4)
>>> G.add_edge(5,6)
>>> graphs = list(nx.connected_component_subgraphs(G))
```

# node\_connected\_component

# $node\_connected\_component(G, n)$

Return the nodes in the component of graph containing node n.

**Parameters G**: NetworkX Graph

An undirected graph.

**n**: node label

A node in G

Returns comp: lists

A list of nodes in component of G containing node n.

### See also:

```
connected_components
```

### **Notes**

For undirected graphs only.

# 4.12.2 Strong connectivity

Strongly connected components.

$is\_strongly\_connected(G)$	Test directed graph for strong connectivity.
${\tt number\_strongly\_connected\_components}(G)$	Return number of strongly connected components in graph.
	Continued on next page

4.12. Components 201

Table 4.38 – continued from previous page

${ t strongly\_connected\_components}(G)$	Generate nodes in strongly connected components of graph.
$strongly\_connected\_component\_subgraphs(G[, copy])$	Generate strongly connected components as subgraphs.
$strongly\_connected\_components\_recursive(G)$	Generate nodes in strongly connected components of graph.
kosaraju_strongly_connected_components( $G[,]$ )	Generate nodes in strongly connected components of graph.
$ ext{condensation}(G[, ext{scc}])$	Returns the condensation of G.

# is\_strongly\_connected

# $is\_strongly\_connected(G)$

Test directed graph for strong connectivity.

Parameters G: NetworkX Graph

A directed graph.

Returns connected: bool

True if the graph is strongly connected, False otherwise.

### See also:

strongly\_connected\_components

### **Notes**

For directed graphs only.

# number\_strongly\_connected\_components

# $number\_strongly\_connected\_components(G)$

Return number of strongly connected components in graph.

**Parameters G**: NetworkX graph

A directed graph.

Returns n: integer

Number of strongly connected components

# See also:

connected\_components

### Notes

For directed graphs only.

# strongly\_connected\_components

### $strongly\_connected\_components(G)$

Generate nodes in strongly connected components of graph.

Parameters G: NetworkX Graph

An directed graph.

Returns comp: generator of lists

A list of nodes for each strongly connected component of G.

Raises NetworkXNotImplemented: If G is undirected.

### See also:

connected\_components, weakly\_connected\_components

### **Notes**

Uses Tarjan's algorithm with Nuutila's modifications. Nonrecursive version of algorithm.

#### References

[R245], [R246]

## strongly\_connected\_component\_subgraphs

# strongly\_connected\_component\_subgraphs (G, copy=True)

Generate strongly connected components as subgraphs.

Parameters G: NetworkX Graph

A graph.

copy: boolean, optional

if copy is True, Graph, node, and edge attributes are copied to the subgraphs.

**Returns** comp: generator of lists

A list of graphs, one for each strongly connected component of G.

### See also:

connected component subgraphs

### strongly connected components recursive

### $strongly\_connected\_components\_recursive(G)$

Generate nodes in strongly connected components of graph.

Recursive version of algorithm.

Parameters G: NetworkX Graph

An directed graph.

**Returns** comp: generator of lists

A list of nodes for each component of G. The list is ordered from largest connected component to smallest.

Raises NetworkXNotImplemented : If G is undirected

### See also:

connected\_components

4.12. Components 203

### **Notes**

Uses Tarjan's algorithm with Nuutila's modifications.

### References

[R247], [R248]

# kosaraju\_strongly\_connected\_components

# kosaraju\_strongly\_connected\_components(G, source=None)

Generate nodes in strongly connected components of graph.

Parameters G: NetworkX Graph

An directed graph.

**Returns** comp: generator of lists

A list of nodes for each component of G.

Raises NetworkXNotImplemented: If G is undirected.

### See also:

connected components

### **Notes**

Uses Kosaraju's algorithm.

#### condensation

### condensation (G, scc=None)

Returns the condensation of G.

The condensation of G is the graph with each of the strongly connected components contracted into a single node.

# Parameters G: NetworkX DiGraph

A directed graph.

### scc: list or generator (optional, default=None)

Strongly connected components. If provided, the elements in scc must partition the nodes in G. If not provided, it will be calculated as  $scc=nx.strongly\_connected\_components(G)$ .

# Returns C: NetworkX DiGraph

The condensation graph C of G. The node labels are integers corresponding to the index of the component in the list of strongly connected components of G. C has a graph attribute named 'mapping' with a dictionary mapping the original nodes to the nodes in C to which they belong. Each node in C also has a node attribute 'members' with the list of original nodes in G that form the SCC that the node in C represents.

Raises NetworkXNotImplemented: If G is not directed

### **Notes**

After contracting all strongly connected components to a single node, the resulting graph is a directed acyclic graph.

# 4.12.3 Weak connectivity

Weakly connected components.

is_weakly_connected(G)	Test directed graph for weak connectivity.
${\tt number\_weakly\_connected\_components}(G)$	Return the number of connected components in G.
${\tt weakly\_connected\_components}(G)$	Generate weakly connected components of G.
weakly_connected_component_subgraphs $(G[, copy])$	Generate weakly connected components as subgraphs.

# is\_weakly\_connected

### $is\_weakly\_connected(G)$

Test directed graph for weak connectivity.

A directed graph is weakly connected if, and only if, the graph is connected when the direction of the edge between nodes is ignored.

Parameters G: NetworkX Graph

A directed graph.

Returns connected: bool

True if the graph is weakly connected, False otherwise.

### See also:

 $\verb|is_strongly_connected|, \verb|is_semiconnected|, \verb|is_connected||$ 

### **Notes**

For directed graphs only.

# number\_weakly\_connected\_components

# ${\tt number\_weakly\_connected\_components}\,(G)$

Return the number of connected components in G. For directed graphs only.

# weakly\_connected\_components

# $weakly\_connected\_components(G)$

Generate weakly connected components of G.

# weakly\_connected\_component\_subgraphs

# weakly\_connected\_component\_subgraphs (G, copy=True)

Generate weakly connected components as subgraphs.

4.12. Components 205

Parameters G: NetworkX Graph

A directed graph.

copy: bool

If copy is True, graph, node, and edge attributes are copied to the subgraphs.

# 4.12.4 Attracting components

### Attracting components.

$is\_attracting\_component(G)$	Returns True if $G$ consists of a single attracting component.
$number\_attracting\_components(G)$	Returns the number of attracting components in $G$ .
$\overline{ ext{attracting\_components}(G)}$	Generates a list of attracting components in $G$ .
attracting_component_subgraphs(G[, copy])	Generates a list of attracting component subgraphs from $G$ .

# is attracting component

### $is\_attracting\_component(G)$

Returns True if G consists of a single attracting component.

Parameters G: DiGraph, MultiDiGraph

The graph to be analyzed.

Returns attracting: bool

True if G has a single attracting component. Otherwise, False.

### See also:

attracting\_components, number\_attracting\_components, attracting\_component\_subgraphs

# number\_attracting\_components

### $number_attracting_components(G)$

Returns the number of attracting components in G.

Parameters G: DiGraph, MultiDiGraph

The graph to be analyzed.

Returns n: int

The number of attracting components in G.

### See also:

attracting\_components, is\_attracting\_component, attracting\_component\_subgraphs

# attracting\_components

### $attracting\_components(G)$

Generates a list of attracting components in G.

An attracting component in a directed graph G is a strongly connected component with the property that a random walker on the graph will never leave the component, once it enters the component.

The nodes in attracting components can also be thought of as recurrent nodes. If a random walker enters the attractor containing the node, then the node will be visited infinitely often.

Parameters G: DiGraph, MultiDiGraph

The graph to be analyzed.

Returns attractors: generator of list

The list of attracting components, sorted from largest attracting component to smallest attracting component.

### See also:

```
number_attracting_components,
attracting_component_subgraphs
```

is\_attracting\_component,

# attracting component subgraphs

### attracting\_component\_subgraphs (G, copy=True)

Generates a list of attracting component subgraphs from G.

Parameters G: DiGraph, MultiDiGraph

The graph to be analyzed.

Returns subgraphs: list

A list of node-induced subgraphs of the attracting components of G.

copy: bool

If copy is True, graph, node, and edge attributes are copied to the subgraphs.

### See also:

attracting\_components, number\_attracting\_components, is\_attracting\_component

# 4.12.5 Biconnected components

Biconnected components and articulation points.

is_biconnected(G)	Return True if the graph is biconnected, False otherwise.
$ ext{biconnected\_components}(G)$	Return a generator of sets of nodes, one set for each biconnected
${\tt biconnected\_component\_edges}(G)$	Return a generator of lists of edges, one list for each biconnected compor
$\verb biconnected_component_subgraphs(G[,copy]) $	Return a generator of graphs, one graph for each biconnected component
$\operatorname{articulation\_points}(G)$	Return a generator of articulation points, or cut vertices, of a graph.

### is biconnected

### $is\_biconnected(G)$

Return True if the graph is biconnected, False otherwise.

A graph is biconnected if, and only if, it cannot be disconnected by removing only one node (and all edges incident on that node). If removing a node increases the number of disconnected components in the graph, that node is called an articulation point, or cut vertex. A biconnected graph has no articulation points.

**Parameters G**: NetworkX Graph

An undirected graph.

4.12. Components 207

#### Returns biconnected: bool

True if the graph is biconnected, False otherwise.

# Raises NetworkXNotImplemented:

If the input graph is not undirected.

#### See also:

```
biconnected_components, articulation_points, biconnected_component_edges, biconnected_component_subgraphs
```

#### **Notes**

The algorithm to find articulation points and biconnected components is implemented using a non-recursive depth-first-search (DFS) that keeps track of the highest level that back edges reach in the DFS tree. A node n is an articulation point if, and only if, there exists a subtree rooted at n such that there is no back edge from any successor of n that links to a predecessor of n in the DFS tree. By keeping track of all the edges traversed by the DFS we can obtain the biconnected components because all edges of a bicomponent will be traversed consecutively between articulation points.

#### References

[R244]

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> print(nx.is_biconnected(G))
False
>>> G.add_edge(0,3)
>>> print(nx.is_biconnected(G))
True
```

### biconnected\_components

### $biconnected\_components(G)$

Return a generator of sets of nodes, one set for each biconnected component of the graph

Biconnected components are maximal subgraphs such that the removal of a node (and all edges incident on that node) will not disconnect the subgraph. Note that nodes may be part of more than one biconnected component. Those nodes are articulation points, or cut vertices. The removal of articulation points will increase the number of connected components of the graph.

Notice that by convention a dyad is considered a biconnected component.

Parameters G: NetworkX Graph

An undirected graph.

Returns nodes: generator

Generator of sets of nodes, one set for each biconnected component.

### Raises NetworkXNotImplemented:

If the input graph is not undirected.

#### See also:

```
is_biconnected, articulation_points, biconnected_component_edges, biconnected component subgraphs
```

#### **Notes**

The algorithm to find articulation points and biconnected components is implemented using a non-recursive depth-first-search (DFS) that keeps track of the highest level that back edges reach in the DFS tree. A node n is an articulation point if, and only if, there exists a subtree rooted at n such that there is no back edge from any successor of n that links to a predecessor of n in the DFS tree. By keeping track of all the edges traversed by the DFS we can obtain the biconnected components because all edges of a bicomponent will be traversed consecutively between articulation points.

### References

[R243]

### **Examples**

```
>>> G = nx.barbell_graph(4,2)
>>> print(nx.is_biconnected(G))
False
>>> components = nx.biconnected_components(G)
>>> G.add_edge(2,8)
>>> print(nx.is_biconnected(G))
True
>>> components = nx.biconnected_components(G)
```

### biconnected\_component\_edges

### $biconnected\_component\_edges(G)$

Return a generator of lists of edges, one list for each biconnected component of the input graph.

Biconnected components are maximal subgraphs such that the removal of a node (and all edges incident on that node) will not disconnect the subgraph. Note that nodes may be part of more than one biconnected component. Those nodes are articulation points, or cut vertices. However, each edge belongs to one, and only one, biconnected component.

Notice that by convention a dyad is considered a biconnected component.

Parameters G: NetworkX Graph

An undirected graph.

Returns edges: generator of lists

Generator of lists of edges, one list for each bicomponent.

### Raises NetworkXNotImplemented:

If the input graph is not undirected.

4.12. Components 209

### See also:

```
is_biconnected, biconnected_components, articulation_points, biconnected component subgraphs
```

#### **Notes**

The algorithm to find articulation points and biconnected components is implemented using a non-recursive depth-first-search (DFS) that keeps track of the highest level that back edges reach in the DFS tree. A node n is an articulation point if, and only if, there exists a subtree rooted at n such that there is no back edge from any successor of n that links to a predecessor of n in the DFS tree. By keeping track of all the edges traversed by the DFS we can obtain the biconnected components because all edges of a bicomponent will be traversed consecutively between articulation points.

#### References

[R241]

### **Examples**

```
>>> G = nx.barbell_graph(4,2)
>>> print(nx.is_biconnected(G))
False
>>> components = nx.biconnected_component_edges(G)
>>> G.add_edge(2,8)
>>> print(nx.is_biconnected(G))
True
>>> components = nx.biconnected_component_edges(G)
```

### biconnected component subgraphs

### biconnected\_component\_subgraphs (*G*, *copy=True*)

Return a generator of graphs, one graph for each biconnected component of the input graph.

Biconnected components are maximal subgraphs such that the removal of a node (and all edges incident on that node) will not disconnect the subgraph. Note that nodes may be part of more than one biconnected component. Those nodes are articulation points, or cut vertices. The removal of articulation points will increase the number of connected components of the graph.

Notice that by convention a dyad is considered a biconnected component.

**Parameters G**: NetworkX Graph

An undirected graph.

Returns graphs: generator

Generator of graphs, one graph for each biconnected component.

### Raises NetworkXNotImplemented:

If the input graph is not undirected.

#### See also:

```
is_biconnected, articulation_points, biconnected_component_edges, biconnected components
```

#### **Notes**

The algorithm to find articulation points and biconnected components is implemented using a non-recursive depth-first-search (DFS) that keeps track of the highest level that back edges reach in the DFS tree. A node n is an articulation point if, and only if, there exists a subtree rooted at n such that there is no back edge from any successor of n that links to a predecessor of n in the DFS tree. By keeping track of all the edges traversed by the DFS we can obtain the biconnected components because all edges of a bicomponent will be traversed consecutively between articulation points.

Graph, node, and edge attributes are copied to the subgraphs.

#### References

[R242]

#### **Examples**

```
>>> G = nx.barbell_graph(4,2)
>>> print(nx.is_biconnected(G))
False
>>> subgraphs = list(nx.biconnected_component_subgraphs(G))
```

## articulation\_points

## $articulation\_points(G)$

Return a generator of articulation points, or cut vertices, of a graph.

An articulation point or cut vertex is any node whose removal (along with all its incident edges) increases the number of connected components of a graph. An undirected connected graph without articulation points is biconnected. Articulation points belong to more than one biconnected component of a graph.

Notice that by convention a dyad is considered a biconnected component.

```
Parameters G: NetworkX Graph
```

An undirected graph.

**Returns articulation points**: generator

generator of nodes

# Raises NetworkXNotImplemented:

If the input graph is not undirected.

#### See also:

```
is_biconnected, biconnected_components, biconnected_component_edges, biconnected component subgraphs
```

4.12. Components 211

#### **Notes**

The algorithm to find articulation points and biconnected components is implemented using a non-recursive depth-first-search (DFS) that keeps track of the highest level that back edges reach in the DFS tree. A node n is an articulation point if, and only if, there exists a subtree rooted at n such that there is no back edge from any successor of n that links to a predecessor of n in the DFS tree. By keeping track of all the edges traversed by the DFS we can obtain the biconnected components because all edges of a bicomponent will be traversed consecutively between articulation points.

#### References

[R240]

#### **Examples**

```
>>> G = nx.barbell_graph(4,2)
>>> print(nx.is_biconnected(G))
False
>>> list(nx.articulation_points(G))
[6, 5, 4, 3]
>>> G.add_edge(2,8)
>>> print(nx.is_biconnected(G))
True
>>> list(nx.articulation_points(G))
[]
```

### 4.12.6 Semiconnectedness

Semiconnectedness.

is\_semiconnected(G) Return True if the graph is semiconnected, False otherwise.

### is semiconnected

### $is\_semiconnected(G)$

Return True if the graph is semiconnected, False otherwise.

A graph is semiconnected if, and only if, for any pair of nodes, either one is reachable from the other, or they are mutually reachable.

Parameters G: NetworkX graph

A directed graph.

Returns semiconnected: bool

True if the graph is semiconnected, False otherwise.

# Raises NetworkXNotImplemented:

If the input graph is not directed.

### **NetworkXPointlessConcept:**

If the graph is empty.

# See also:

```
is_strongly_connected, is_weakly_connected
```

# **Examples**

```
>>> G=nx.path_graph(4,create_using=nx.DiGraph())
>>> print(nx.is_semiconnected(G))
True
>>> G=nx.DiGraph([(1, 2), (3, 2)])
>>> print(nx.is_semiconnected(G))
False
```

# 4.13 Connectivity

Connectivity and cut algorithms

# 4.13.1 Flow-based Connectivity

Flow based connectivity algorithms

average_node_connectivity(G[, flow_func])	Returns the average connectivity of a graph G.
all_pairs_node_connectivity( $G[$ , $nbunch$ ,])	Compute node connectivity between all pairs of nodes of G.
$\verb  edge_connectivity(G[, s, t, flow_func])  \\$	Returns the edge connectivity of the graph or digraph G.
$local\_edge\_connectivity(G, u, v[,])$	Returns local edge connectivity for nodes s and t in G.
$local_node_connectivity(G, s, t[,])$	Computes local node connectivity for nodes s and t.
<pre>node_connectivity(G[, s, t, flow_func])</pre>	Returns node connectivity for a graph or digraph G.

# average\_node\_connectivity

average\_node\_connectivity(G, flow\_func=None)

Returns the average connectivity of a graph G.

The average connectivity  $\bar{\kappa}$  of a graph G is the average of local node connectivity over all pairs of nodes of G [R249] .

$$\bar{\kappa}(G) = \frac{\sum_{u,v} \kappa_G(u,v)}{\binom{n}{2}}$$

Parameters G: NetworkX graph

Undirected graph

flow\_func : function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See local\_node\_connectivity() for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

#### Returns K: float

Average node connectivity

### See also:

#### References

[R249]

# all pairs node connectivity

# all\_pairs\_node\_connectivity(G, nbunch=None, flow\_func=None)

Compute node connectivity between all pairs of nodes of G.

### **Parameters G**: NetworkX graph

Undirected graph

#### nbunch: container

Container of nodes. If provided node connectivity will be computed only over pairs of nodes in nbunch.

# flow\_func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

# Returns all\_pairs: dict

A dictionary with node connectivity between all pairs of nodes in G, or in nbunch if provided.

# See also:

```
local_node_connectivity(), edge_connectivity(), local_edge_connectivity(),
maximum_flow(),edmonds_karp(),preflow_push(),shortest_augmenting_path()
```

# edge\_connectivity

### edge\_connectivity(G, s=None, t=None, flow\_func=None)

Returns the edge connectivity of the graph or digraph G.

The edge connectivity is equal to the minimum number of edges that must be removed to disconnect G or render it trivial. If source and target nodes are provided, this function returns the local edge connectivity: the minimum number of edges that must be removed to break all paths from source to target in G.

#### **Parameters G**: NetworkX graph

Undirected or directed graph

s: node

Source node. Optional. Default value: None.

t: node

Target node. Optional. Default value: None.

flow\_func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

Returns K: integer

Edge connectivity for G, or local edge connectivity if source and target were provided

#### See also:

```
local_edge_connectivity(), local_node_connectivity(), node_connectivity(),
maximum_flow(),edmonds_karp(),preflow_push(),shortest_augmenting_path()
```

#### **Notes**

This is a flow based implementation of global edge connectivity. For undirected graphs the algorithm works by finding a 'small' dominating set of nodes of G (see algorithm 7 in [R250]) and computing local maximum flow (see local\_edge\_connectivity()) between an arbitrary node in the dominating set and the rest of nodes in it. This is an implementation of algorithm 6 in [R250]. For directed graphs, the algorithm does n calls to the maximum flow function. This is an implementation of algorithm 8 in [R250].

#### References

[R250]

# **Examples**

```
>>> # Platonic icosahedral graph is 5-edge-connected
>>> G = nx.icosahedral_graph()
>>> nx.edge_connectivity(G)
5
```

You can use alternative flow algorithms for the underlying maximum flow computation. In dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp(), which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> nx.edge_connectivity(G, flow_func=shortest_augmenting_path)
5
```

If you specify a pair of nodes (source and target) as parameters, this function returns the value of local edge connectivity.

```
>>> nx.edge_connectivity(G, 3, 7)
5
```

If you need to perform several local computations among different pairs of nodes on the same graph, it is recommended that you reuse the data structures used in the maximum flow computations. See <code>local\_edge\_connectivity()</code> for details.

# local\_edge\_connectivity

 $local\_edge\_connectivity$  (G, u, v,  $flow\_func=None$ , auxiliary=None, residual=None, cutoff=None) Returns local edge connectivity for nodes s and t in G.

Local edge connectivity for two nodes s and t is the minimum number of edges that must be removed to disconnect them.

This is a flow based implementation of edge connectivity. We compute the maximum flow on an auxiliary digraph build from the original network (see below for details). This is equal to the local edge connectivity because the value of a maximum s-t-flow is equal to the capacity of a minimum s-t-cut (Ford and Fulkerson theorem) [R251].

## **Parameters G**: NetworkX graph

Undirected or directed graph

s: node

Source node

t: node

Target node

flow func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

# auxiliary: NetworkX DiGraph

Auxiliary digraph for computing flow based edge connectivity. If provided it will be reused instead of recreated. Default value: None.

### residual: NetworkX DiGraph

Residual network to compute maximum flow. If provided it will be reused instead of recreated. Default value: None.

# cutoff: integer, float

If specified, the maximum flow algorithm will terminate when the flow value reaches or exceeds the cutoff. This is only for the algorithms that support the cutoff parameter: edmonds\_karp() and shortest\_augmenting\_path(). Other algorithms will ignore this parameter. Default value: None.

# Returns K: integer

local edge connectivity for nodes s and t.

#### See also:

```
edge_connectivity(), local_node_connectivity(), node_connectivity(),
maximum_flow(),edmonds_karp(),preflow_push(),shortest_augmenting_path()
```

#### **Notes**

This is a flow based implementation of edge connectivity. We compute the maximum flow using, by default, the edmonds\_karp() algorithm on an auxiliary digraph build from the original input graph:

If the input graph is undirected, we replace each edge (u, v) with two reciprocal arcs (u, v) and (v, u) and then we set the attribute 'capacity' for each arc to 1. If the input graph is directed we simply add the 'capacity' attribute. This is an implementation of algorithm 1 in [R251].

The maximum flow in the auxiliary network is equal to the local edge connectivity because the value of a maximum s-t-flow is equal to the capacity of a minimum s-t-cut (Ford and Fulkerson theorem).

#### References

[R251]

### **Examples**

This function is not imported in the base NetworkX namespace, so you have to explicitly import it from the connectivity package:

```
>>> from networkx.algorithms.connectivity import local_edge_connectivity
```

We use in this example the platonic icosahedral graph, which has edge connectivity 5.

```
>>> G = nx.icosahedral_graph()
>>> local_edge_connectivity(G, 0, 6)
5
```

If you need to compute local connectivity on several pairs of nodes in the same graph, it is recommended that you reuse the data structures that NetworkX uses in the computation: the auxiliary digraph for edge connectivity, and the residual network for the underlying maximum flow computation.

Example of how to compute local edge connectivity among all pairs of nodes of the platonic icosahedral graph reusing the data structures.

```
>>> import itertools
>>> # You also have to explicitly import the function for
>>> # building the auxiliary digraph from the connectivity package
>>> from networkx.algorithms.connectivity import (
       build_auxiliary_edge_connectivity)
>>> H = build_auxiliary_edge_connectivity(G)
>>> # And the function for building the residual network from the
>>> # flow package
>>> from networkx.algorithms.flow import build_residual_network
>>> # Note that the auxiliary digraph has an edge attribute named capacity
>>> R = build_residual_network(H, 'capacity')
>>> result = dict.fromkeys(G, dict())
>>> # Reuse the auxiliary digraph and the residual network by passing them
>>> # as parameters
>>> for u, v in itertools.combinations(G, 2):
       k = local_edge_connectivity(G, u, v, auxiliary=H, residual=R)
       result[u][v] = k
>>> all(result[u][v] == 5 for u, v in itertools.combinations(G, 2))
True
```

You can also use alternative flow algorithms for computing edge connectivity. For instance, in dense networks the algorithm <code>shortest\_augmenting\_path()</code> will usually perform better than the default <code>edmonds\_karp()</code> which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> local_edge_connectivity(G, 0, 6, flow_func=shortest_augmenting_path)
5
```

# local node connectivity

**local\_node\_connectivity** (*G*, *s*, *t*, *flow\_func=None*, *auxiliary=None*, *residual=None*, *cutoff=None*) Computes local node connectivity for nodes s and t.

Local node connectivity for two non adjacent nodes s and t is the minimum number of nodes that must be removed (along with their incident edges) to disconnect them.

This is a flow based implementation of node connectivity. We compute the maximum flow on an auxiliary digraph build from the original input graph (see below for details).

# **Parameters G**: NetworkX graph

Undirected graph

s: node

Source node

t: node

Target node

### flow\_func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

#### auxiliary: NetworkX DiGraph

Auxiliary digraph to compute flow based node connectivity. It has to have a graph attribute called mapping with a dictionary mapping node names in G and in the auxiliary digraph. If provided it will be reused instead of recreated. Default value: None.

### residual: NetworkX DiGraph

Residual network to compute maximum flow. If provided it will be reused instead of recreated. Default value: None.

# cutoff: integer, float

If specified, the maximum flow algorithm will terminate when the flow value reaches or exceeds the cutoff. This is only for the algorithms that support the cutoff parameter: edmonds\_karp() and shortest\_augmenting\_path(). Other algorithms will ignore this parameter. Default value: None.

#### Returns K: integer

local node connectivity for nodes s and t

#### See also:

#### **Notes**

This is a flow based implementation of node connectivity. We compute the maximum flow using, by default, the edmonds\_karp() algorithm (see: maximum\_flow()) on an auxiliary digraph build from the original input graph:

For an undirected graph G having n nodes and m edges we derive a directed graph H with 2n nodes and 2m+n arcs by replacing each original node v with two nodes  $v_A$ ,  $v_B$  linked by an (internal) arc in H. Then for each edge (u,v) in G we add two arcs  $(u_B,v_A)$  and  $(v_B,u_A)$  in H. Finally we set the attribute capacity = 1 for each arc in H [R253].

For a directed graph G having n nodes and m arcs we derive a directed graph H with 2n nodes and m + n arcs by replacing each original node v with two nodes  $v_A$ ,  $v_B$  linked by an (internal) arc  $(v_A, v_B)$  in H. Then for each arc (u, v) in G we add one arc  $(u_B, v_A)$  in H. Finally we set the attribute capacity = 1 for each arc in H.

This is equal to the local node connectivity because the value of a maximum s-t-flow is equal to the capacity of a minimum s-t-cut.

#### References

[R253]

# **Examples**

This function is not imported in the base NetworkX namespace, so you have to explicitly import it from the connectivity package:

```
>>> from networkx.algorithms.connectivity import local_node_connectivity
```

We use in this example the platonic icosahedral graph, which has node connectivity 5.

```
>>> G = nx.icosahedral_graph()
>>> local_node_connectivity(G, 0, 6)
5
```

If you need to compute local connectivity on several pairs of nodes in the same graph, it is recommended that you reuse the data structures that NetworkX uses in the computation: the auxiliary digraph for node connectivity, and the residual network for the underlying maximum flow computation.

Example of how to compute local node connectivity among all pairs of nodes of the platonic icosahedral graph reusing the data structures.

```
>>> import itertools
>>> # You also have to explicitly import the function for
>>> # building the auxiliary digraph from the connectivity package
>>> from networkx.algorithms.connectivity import (
... build_auxiliary_node_connectivity)
>>> H = build_auxiliary_node_connectivity(G)
>>> # And the function for building the residual network from the
>>> # flow package
>>> from networkx.algorithms.flow import build_residual_network
```

You can also use alternative flow algorithms for computing node connectivity. For instance, in dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp() which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> local_node_connectivity(G, 0, 6, flow_func=shortest_augmenting_path)
5
```

# node\_connectivity

```
node_connectivity (G, s=None, t=None, flow_func=None) Returns node connectivity for a graph or digraph G.
```

Node connectivity is equal to the minimum number of nodes that must be removed to disconnect G or render it trivial. If source and target nodes are provided, this function returns the local node connectivity: the minimum number of nodes that must be removed to break all paths from source to target in G.

## Parameters G: NetworkX graph

Undirected graph

s: node

Source node. Optional. Default value: None.

t: node

Target node. Optional. Default value: None.

flow func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

### Returns K: integer

Node connectivity of G, or local node connectivity if source and target are provided.

# See also:

```
local_node_connectivity(), edge_connectivity(), maximum_flow(),
edmonds_karp(),preflow_push(),shortest_augmenting_path()
```

#### **Notes**

This is a flow based implementation of node connectivity. The algorithm works by solving  $O((n-\delta-1+\delta(\delta-1)/2))$  maximum flow problems on an auxiliary digraph. Where  $\delta$  is the minimum degree of G. For details about the auxiliary digraph and the computation of local node connectivity see <code>local\_node\_connectivity()</code>. This implementation is based on algorithm 11 in [R254].

#### References

[R254]

#### **Examples**

```
>>> # Platonic icosahedral graph is 5-node-connected
>>> G = nx.icosahedral_graph()
>>> nx.node_connectivity(G)
5
```

You can use alternative flow algorithms for the underlying maximum flow computation. In dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp(), which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> nx.node_connectivity(G, flow_func=shortest_augmenting_path)
5
```

If you specify a pair of nodes (source and target) as parameters, this function returns the value of local node connectivity.

```
>>> nx.node_connectivity(G, 3, 7)
```

If you need to perform several local computations among different pairs of nodes on the same graph, it is recommended that you reuse the data structures used in the maximum flow computations. See local node connectivity() for details.

# 4.13.2 Flow-based Minimum Cuts

#### Flow based cut algorithms

$minimum\_edge\_cut(G[, s, t, flow\_func])$	Returns a set of edges of minimum cardinality that disconnects G.
$minimum\_node\_cut(G[, s, t, flow\_func])$	Returns a set of nodes of minimum cardinality that disconnects G.
$minimum_st_edge_cut(G, s, t[, flow_func,])$	Returns the edges of the cut-set of a minimum (s, t)-cut.
minimum_st_node_cut(G, s, t[, flow_func,])	Returns a set of nodes of minimum cardinality that disconnect source from targ

### minimum\_edge\_cut

```
minimum_edge_cut (G, s=None, t=None, flow_func=None)

Returns a set of edges of minimum cardinality that disconnects G.
```

If source and target nodes are provided, this function returns the set of edges of minimum cardinality that, if

removed, would break all paths among source and target in G. If not, it returns a set of edges of minimum cardinality that disconnects G.

### **Parameters G**: NetworkX graph

s: node

Source node. Optional. Default value: None.

t · node

Target node. Optional. Default value: None.

flow\_func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

#### Returns cutset: set

Set of edges that, if removed, would disconnect G. If source and target nodes are provided, the set contians the edges that if removed, would destroy all paths between source and target.

#### See also:

# **Notes**

This is a flow based implementation of minimum edge cut. For undirected graphs the algorithm works by finding a 'small' dominating set of nodes of G (see algorithm 7 in [R255]) and computing the maximum flow between an arbitrary node in the dominating set and the rest of nodes in it. This is an implementation of algorithm 6 in [R255]. For directed graphs, the algorithm does n calls to the max flow function. It is an implementation of algorithm 8 in [R255].

#### References

[R255]

#### **Examples**

```
>>> # Platonic icosahedral graph has edge connectivity 5
>>> G = nx.icosahedral_graph()
>>> len(nx.minimum_edge_cut(G))
5
```

You can use alternative flow algorithms for the underlying maximum flow computation. In dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp(), which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> len(nx.minimum_edge_cut(G, flow_func=shortest_augmenting_path))
5
```

If you specify a pair of nodes (source and target) as parameters, this function returns the value of local edge connectivity.

```
>>> nx.edge_connectivity(G, 3, 7)
5
```

If you need to perform several local computations among different pairs of nodes on the same graph, it is recommended that you reuse the data structures used in the maximum flow computations. See local\_edge\_connectivity() for details.

# minimum node cut

```
minimum_node_cut (G, s=None, t=None, flow_func=None)
```

Returns a set of nodes of minimum cardinality that disconnects G.

If source and target nodes are provided, this function returns the set of nodes of minimum cardinality that, if removed, would destroy all paths among source and target in G. If not, it returns a set of nodes of minimum cardinality that disconnects G.

# Parameters G: NetworkX graph

s: node

Source node. Optional. Default value: None.

t: node

Target node. Optional. Default value: None.

### flow\_func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

### Returns cutset: set

Set of nodes that, if removed, would disconnect G. If source and target nodes are provided, the set contians the nodes that if removed, would destroy all paths between source and target.

#### See also:

```
minimum_st_node_cut(), minimum_cut(), minimum_edge_cut(), stoer_wagner(),
node_connectivity(), edge_connectivity(), maximum_flow(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

#### **Notes**

This is a flow based implementation of minimum node cut. The algorithm is based in solving a number of maximum flow computations to determine the capacity of the minimum cut on an auxiliary directed network that

corresponds to the minimum node cut of G. It handles both directed and undirected graphs. This implementation is based on algorithm 11 in [R256].

#### References

[R256]

#### **Examples**

```
>>> # Platonic icosahedral graph has node connectivity 5
>>> G = nx.icosahedral_graph()
>>> node_cut = nx.minimum_node_cut(G)
>>> len(node_cut)
5
```

You can use alternative flow algorithms for the underlying maximum flow computation. In dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp(), which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> node_cut == nx.minimum_node_cut(G, flow_func=shortest_augmenting_path)
True
```

If you specify a pair of nodes (source and target) as parameters, this function returns a local st node cut.

```
>>> len(nx.minimum_node_cut(G, 3, 7))
5
```

If you need to perform several local st cuts among different pairs of nodes on the same graph, it is recommended that you reuse the data structures used in the maximum flow computations. See minimum\_st\_node\_cut() for details.

#### minimum st edge cut

```
minimum_st_edge_cut(G, s, t, flow_func=None, auxiliary=None, residual=None)
Returns the edges of the cut-set of a minimum (s, t)-cut.
```

This function returns the set of edges of minimum cardinality that, if removed, would destroy all paths among source and target in G. Edge weights are not considered

# **Parameters G**: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

s: node

Source node for the flow.

t: node

224

Sink node for the flow.

auxiliary: NetworkX DiGraph

Auxiliary digraph to compute flow based node connectivity. It has to have a graph attribute called mapping with a dictionary mapping node names in G and in the auxiliary digraph. If provided it will be reused instead of recreated. Default value: None.

# flow\_func : function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See node\_connectivity() for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

### residual: NetworkX DiGraph

Residual network to compute maximum flow. If provided it will be reused instead of recreated. Default value: None.

#### Returns cutset: set

Set of edges that, if removed from the graph, will disconnect it.

#### See also:

```
minimum_cut(), minimum_node_cut(), minimum_edge_cut(), stoer_wagner(),
node_connectivity(), edge_connectivity(), maximum_flow(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

# **Examples**

This function is not imported in the base NetworkX namespace, so you have to explicitly import it from the connectivity package:

```
>>> from networkx.algorithms.connectivity import minimum_st_edge_cut
```

We use in this example the platonic icosahedral graph, which has edge connectivity 5.

```
>>> G = nx.icosahedral_graph()
>>> len(minimum_st_edge_cut(G, 0, 6))
5
```

If you need to compute local edge cuts on several pairs of nodes in the same graph, it is recommended that you reuse the data structures that NetworkX uses in the computation: the auxiliary digraph for edge connectivity, and the residual network for the underlying maximum flow computation.

Example of how to compute local edge cuts among all pairs of nodes of the platonic icosahedral graph reusing the data structures.

```
>>> import itertools
>>> # You also have to explicitly import the function for
>>> # building the auxiliary digraph from the connectivity package
>>> from networkx.algorithms.connectivity import (
... build_auxiliary_edge_connectivity)
>>> H = build_auxiliary_edge_connectivity(G)
>>> # And the function for building the residual network from the
>>> # flow package
>>> from networkx.algorithms.flow import build_residual_network
>>> # Note that the auxiliary digraph has an edge attribute named capacity
>>> R = build_residual_network(H, 'capacity')
>>> result = dict.fromkeys(G, dict())
```

```
>>> # Reuse the auxiliary digraph and the residual network by passing them
>>> # as parameters
>>> for u, v in itertools.combinations(G, 2):
... k = len(minimum_st_edge_cut(G, u, v, auxiliary=H, residual=R))
... result[u][v] = k
>>> all(result[u][v] == 5 for u, v in itertools.combinations(G, 2))
True
```

You can also use alternative flow algorithms for computing edge cuts. For instance, in dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp() which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> len(minimum_st_edge_cut(G, 0, 6, flow_func=shortest_augmenting_path))
5
```

# minimum\_st\_node\_cut

 $minimum\_st\_node\_cut(G, s, t, flow\_func=None, auxiliary=None, residual=None)$ 

Returns a set of nodes of minimum cardinality that disconnect source from target in G.

This function returns the set of nodes of minimum cardinality that, if removed, would destroy all paths among source and target in G.

# Parameters G: NetworkX graph

s: node

Source node.

t: node

Target node.

# flow func: function

A function for computing the maximum flow among a pair of nodes. The function has to accept at least three parameters: a Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see maximum\_flow() for details). If flow\_func is None, the default maximum flow function (edmonds\_karp()) is used. See below for details. The choice of the default function may change from version to version and should not be relied on. Default value: None.

# auxiliary: NetworkX DiGraph

Auxiliary digraph to compute flow based node connectivity. It has to have a graph attribute called mapping with a dictionary mapping node names in G and in the auxiliary digraph. If provided it will be reused instead of recreated. Default value: None.

#### residual: NetworkX DiGraph

Residual network to compute maximum flow. If provided it will be reused instead of recreated. Default value: None.

#### Returns cutset: set

Set of nodes that, if removed, would destroy all paths between source and target in G.

See also:

```
minimum_node_cut(), minimum_edge_cut(), stoer_wagner(), node_connectivity(),
edge_connectivity(), maximum_flow(), edmonds_karp(), preflow_push(),
shortest_augmenting_path()
```

#### **Notes**

This is a flow based implementation of minimum node cut. The algorithm is based in solving a number of maximum flow computations to determine the capacity of the minimum cut on an auxiliary directed network that corresponds to the minimum node cut of G. It handles both directed and undirected graphs. This implementation is based on algorithm 11 in [R258].

#### References

[R258]

#### **Examples**

This function is not imported in the base NetworkX namespace, so you have to explicitly import it from the connectivity package:

```
>>> from networkx.algorithms.connectivity import minimum_st_node_cut
```

We use in this example the platonic icosahedral graph, which has node connectivity 5.

```
>>> G = nx.icosahedral_graph()
>>> len(minimum_st_node_cut(G, 0, 6))
5
```

If you need to compute local st cuts between several pairs of nodes in the same graph, it is recommended that you reuse the data structures that NetworkX uses in the computation: the auxiliary digraph for node connectivity and node cuts, and the residual network for the underlying maximum flow computation.

Example of how to compute local st node cuts reusing the data structures:

```
>>> # You also have to explicitly import the function for
>>> # building the auxiliary digraph from the connectivity package
>>> from networkx.algorithms.connectivity import (
... build_auxiliary_node_connectivity)
>>> H = build_auxiliary_node_connectivity(G)
>>> # And the function for building the residual network from the
>>> # flow package
>>> from networkx.algorithms.flow import build_residual_network
>>> # Note that the auxiliary digraph has an edge attribute named capacity
>>> R = build_residual_network(H, 'capacity')
>>> # Reuse the auxiliary digraph and the residual network by passing them
>>> # as parameters
>>> len(minimum_st_node_cut(G, 0, 6, auxiliary=H, residual=R))
```

You can also use alternative flow algorithms for computing minimum st node cuts. For instance, in dense networks the algorithm shortest\_augmenting\_path() will usually perform better than the default edmonds\_karp() which is faster for sparse networks with highly skewed degree distributions. Alternative flow functions have to be explicitly imported from the flow package.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> len(minimum_st_node_cut(G, 0, 6, flow_func=shortest_augmenting_path))
5
```

# 4.13.3 Stoer-Wagner minimum cut

Stoer-Wagner minimum cut algorithm.

stoer\_wagner(G[, weight, heap]) Returns the weighted minimum edge cut using the Stoer-Wagner algorithm.

# stoer wagner

 $\verb|stoer_wagner| (G, weight='weight', heap=<\!class' inetwork x.utils.heaps.Binary Heap'>)|$ 

Returns the weighted minimum edge cut using the Stoer-Wagner algorithm.

Determine the minimum edge cut of a connected graph using the Stoer-Wagner algorithm. In weighted cases, all weights must be nonnegative.

The running time of the algorithm depends on the type of heaps used:

Type of heap	Running time
Binary heap	$O(n(m+n)\log n)$
Fibonacci heap	$O(nm + n^2 \log n)$
Pairing heap	$O(2^{2\sqrt{\log\log n}}nm + n^2\log n)$

### **Parameters G**: NetworkX graph

Edges of the graph are expected to have an attribute named by the weight parameter below. If this attribute is not present, the edge is considered to have unit weight.

#### weight: string

Name of the weight attribute of the edges. If the attribute is not present, unit weight is assumed. Default value: 'weight'.

# heap: class

Type of heap to be used in the algorithm. It should be a subclass of MinHeap or implement a compatible interface.

If a stock heap implementation is to be used, BinaryHeap is recommeded over PairingHeap for Python implementations without optimized attribute accesses (e.g., CPython) despite a slower asymptotic running time. For Python implementations with optimized attribute accesses (e.g., PyPy), PairingHeap provides better performance. Default value: BinaryHeap.

#### Returns cut\_value: integer or float

The sum of weights of edges in a minimum cut.

# partition: pair of node lists

A partitioning of the nodes that defines a minimum cut.

# Raises NetworkXNotImplemented

If the graph is directed or a multigraph.

#### NetworkXError

If the graph has less than two nodes, is not connected or has a negative-weighted edge.

# **Examples**

```
>>> G = nx.Graph()
>>> G.add_edge('x','a', weight=3)
>>> G.add_edge('x','b', weight=1)
>>> G.add_edge('a','c', weight=3)
>>> G.add_edge('b','c', weight=5)
>>> G.add_edge('b','d', weight=4)
>>> G.add_edge('d','e', weight=2)
>>> G.add_edge('c','y', weight=2)
>>> G.add_edge('e','y', weight=3)
>>> cut_value, partition = nx.stoer_wagner(G)
```

# 4.13.4 Utils for flow-based connectivity

Utilities for connectivity package

```
build_auxiliary_edge_connectivity(G) Auxiliary digraph for computing flow based edge connectivity

build_auxiliary_node_connectivity(G) Creates a directed graph D from an undirected graph G to compute flow based
```

# build\_auxiliary\_edge\_connectivity

### $build_auxiliary_edge_connectivity(G)$

Auxiliary digraph for computing flow based edge connectivity

If the input graph is undirected, we replace each edge (u, v) with two reciprocal arcs (u, v) and (v, u) and then we set the attribute 'capacity' for each arc to 1. If the input graph is directed we simply add the 'capacity' attribute. Part of algorithm 1 in [R260].

#### References

[R260]

# build auxiliary node connectivity

# build\_auxiliary\_node\_connectivity(G)

Creates a directed graph D from an undirected graph G to compute flow based node connectivity.

For an undirected graph G having n nodes and m edges we derive a directed graph D with 2n nodes and 2m+n arcs by replacing each original node v with two nodes vA, vB linked by an (internal) arc in D. Then for each edge (u, v) in G we add two arcs (uB, vA) and (vB, uA) in D. Finally we set the attribute capacity = 1 for each arc in D [R261].

For a directed graph having n nodes and m arcs we derive a directed graph D with 2n nodes and m+n arcs by replacing each original node v with two nodes vA, vB linked by an (internal) arc (vA, vB) in D. Then for each arc (u, v) in G we add one arc (uB, vA) in D. Finally we set the attribute capacity = 1 for each arc in D.

A dictionary with a mapping between nodes in the original graph and the auxiliary digraph is stored as a graph attribute: H.graph['mapping'].

#### References

[R261]

# **4.14 Cores**

Find the k-cores of a graph.

The k-core is found by recursively pruning nodes with degrees less than k.

See the following reference for details:

An O(m) Algorithm for Cores Decomposition of Networks Vladimir Batagelj and Matjaz Zaversnik, 2003. http://arxiv.org/abs/cs.DS/0310049

core_number(G)	Return the core number for each vertex.
	Return the core number for each vertex.
$k\_core(G[, k, core\_number])$	Return the k-core of G.
$k\_shell(G[, k, core\_number])$	Return the k-shell of G.
k_crust(G[, k, core_number])	Return the k-crust of G.
k_corona(G, k[, core_number])	Return the k-corona of G.

# 4.14.1 core\_number

# $core_number(G)$

Return the core number for each vertex.

A k-core is a maximal subgraph that contains nodes of degree k or more.

The core number of a node is the largest value k of a k-core containing that node.

Parameters G: NetworkX graph

A graph or directed graph

**Returns core\_number**: dictionary

A dictionary keyed by node to the core number.

# Raises NetworkXError

The k-core is not defined for graphs with self loops or parallel edges.

#### **Notes**

Not implemented for graphs with parallel edges or self loops.

For directed graphs the node degree is defined to be the in-degree + out-degree.

# References

[R262]

# 4.14.2 k core

 $k\_core(G, k=None, core\_number=None)$ 

Return the k-core of G.

A k-core is a maximal subgraph that contains nodes of degree k or more.

**Parameters G**: NetworkX graph

A graph or directed graph

k: int, optional

The order of the core. If not specified return the main core.

core\_number: dictionary, optional

Precomputed core numbers for the graph G.

**Returns G**: NetworkX graph

The k-core subgraph

Raises NetworkXError

The k-core is not defined for graphs with self loops or parallel edges.

#### See also:

core number

### **Notes**

The main core is the core with the largest degree.

Not implemented for graphs with parallel edges or self loops.

For directed graphs the node degree is defined to be the in-degree + out-degree.

Graph, node, and edge attributes are copied to the subgraph.

# References

[R263]

# 4.14.3 k shell

 $k\_shell(G, k=None, core\_number=None)$ 

Return the k-shell of G.

The k-shell is the subgraph of nodes in the k-core but not in the (k+1)-core.

Parameters G: NetworkX graph

A graph or directed graph.

k: int, optional

The order of the shell. If not specified return the main shell.

core\_number: dictionary, optional

Precomputed core numbers for the graph G.

4.14. Cores 231

**Returns** G: NetworkX graph

The k-shell subgraph

# Raises NetworkXError

The k-shell is not defined for graphs with self loops or parallel edges.

#### See also:

```
core number, k corona, -----
```

Shai Carmi, Shlomo Havlin, Scott Kirkpatrick, Yuval Shavitt, and Eran Shir, PNAS July 3, 2007 vol. 104 no. 27 11150-11154

http://www.pnas.org/content/104/27/11150.full

### Notes

This is similar to k\_corona but in that case only neighbors in the k-core are considered.

Not implemented for graphs with parallel edges or self loops.

For directed graphs the node degree is defined to be the in-degree + out-degree.

Graph, node, and edge attributes are copied to the subgraph.

# 4.14.4 k crust

**k\_crust** (*G*, *k=None*, *core\_number=None*)

Return the k-crust of G.

The k-crust is the graph G with the k-core removed.

**Parameters G**: NetworkX graph

A graph or directed graph.

k: int, optional

The order of the shell. If not specified return the main crust.

core\_number: dictionary, optional

Precomputed core numbers for the graph G.

Returns G: NetworkX graph

The k-crust subgraph

Raises NetworkXError

The k-crust is not defined for graphs with self loops or parallel edges.

# See also:

core\_number

#### **Notes**

This definition of k-crust is different than the definition in [R265]. The k-crust in [R265] is equivalent to the k+1 crust of this algorithm.

Not implemented for graphs with parallel edges or self loops.

For directed graphs the node degree is defined to be the in-degree + out-degree.

Graph, node, and edge attributes are copied to the subgraph.

#### References

[R265]

# 4.14.5 k corona

# k\_corona (G, k, core\_number=None)

Return the k-corona of G.

The k-corona is the subgraph of nodes in the k-core which have exactly k neighbours in the k-core.

**Parameters G**: NetworkX graph

A graph or directed graph

k: int

The order of the corona.

core\_number: dictionary, optional

Precomputed core numbers for the graph G.

Returns G: NetworkX graph

The k-corona subgraph

#### Raises NetworkXError

The k-cornoa is not defined for graphs with self loops or parallel edges.

# See also:

```
core_number
```

### Notes

Not implemented for graphs with parallel edges or self loops.

For directed graphs the node degree is defined to be the in-degree + out-degree.

Graph, node, and edge attributes are copied to the subgraph.

#### References

[R264]

4.14. Cores 233

# 4.15 Cycles

$cycle\_basis(G[,root])$	Returns a list of cycles which form a basis for cycles of G.
$simple\_cycles(G)$	Find simple cycles (elementary circuits) of a directed graph.
<pre>find_cycle(G[, source, orientation])</pre>	Returns the edges of a cycle found via a directed, depth-first traversal.

# 4.15.1 cycle basis

### cycle\_basis(G, root=None)

Returns a list of cycles which form a basis for cycles of G.

A basis for cycles of a network is a minimal collection of cycles such that any cycle in the network can be written as a sum of cycles in the basis. Here summation of cycles is defined as "exclusive or" of the edges. Cycle bases are useful, e.g. when deriving equations for electric circuits using Kirchhoff's Laws.

Parameters G: NetworkX Graph

root: node, optional

Specify starting node for basis.

**Returns** A list of cycle lists. Each cycle list is a list of nodes

which forms a cycle (loop) in G.

### See also:

```
simple_cycles
```

#### **Notes**

This is adapted from algorithm CACM 491 [R266].

#### References

[R266]

#### **Examples**

```
>>> G=nx.Graph()
>>> G.add_cycle([0,1,2,3])
>>> G.add_cycle([0,3,4,5])
>>> print(nx.cycle_basis(G,0))
[[3, 4, 5, 0], [1, 2, 3, 0]]
```

# 4.15.2 simple\_cycles

# $simple\_cycles(G)$

Find simple cycles (elementary circuits) of a directed graph.

An simple cycle, or elementary circuit, is a closed path where no node appears twice, except that the first and last node are the same. Two elementary circuits are distinct if they are not cyclic permutations of each other.

This is a nonrecursive, iterator/generator version of Johnson's algorithm [R267]. There may be better algorithms for some cases [R268] [R269].

Parameters G: NetworkX DiGraph

A directed graph

Returns cycle\_generator: generator

A generator that produces elementary cycles of the graph. Each cycle is a list of nodes with the first and last nodes being the same.

#### See also:

```
cycle_basis
```

#### **Notes**

The implementation follows pp. 79-80 in [R267].

The time complexity is O((n+e)(c+1)) for n nodes, e edges and c elementary circuits.

To filter the cycles so that they don't include certain nodes or edges, copy your graph and eliminate those nodes or edges before calling:

```
>>> copyG = G.copy()
>>> copyG.remove_nodes_from([1])
>>> copyG.remove_edges_from([(0,1)])
>>> list(nx.simple_cycles(copyG))
[[2], [2, 0], [0]]
```

#### References

```
[R267], [R268], [R269]
```

# **Examples**

```
>>> G = nx.DiGraph([(0, 0), (0, 1), (0, 2), (1, 2), (2, 0), (2, 1), (2, 2)])
>>> list(nx.simple_cycles(G))
[[2], [2, 1], [2, 0], [2, 0, 1], [0]]
```

# 4.15.3 find\_cycle

```
find_cycle (G, source=None, orientation='original')
```

Returns the edges of a cycle found via a directed, depth-first traversal.

Parameters G: graph

A directed/undirected graph/multigraph.

```
source: node, list of nodes
```

The node from which the traversal begins. If None, then a source is chosen arbitrarily and repeatedly until all edges from each node in the graph are searched.

```
orientation : 'original' | 'reverse' | 'ignore'
```

4.15. Cycles 235

For directed graphs and directed multigraphs, edge traversals need not respect the original orientation of the edges. When set to 'reverse', then every edge will be traversed in the reverse direction. When set to 'ignore', then each directed edge is treated as a single undirected edge that can be traversed in either direction. For undirected graphs and undirected multigraphs, this parameter is meaningless and is not consulted by the algorithm.

# Returns edges: directed edges

A list of directed edges indicating the path taken for the loop. If no cycle is found, then edges will be an empty list. For graphs, an edge is of the form (u, v) where u and v are the tail and head of the edge as determined by the traversal. For multigraphs, an edge is of the form (u, v, key), where key is the key of the edge. When the graph is directed, then u and v are always in the order of the actual directed edge. If orientation is 'ignore', then an edge takes the form (u, v, key, direction) where direction indicates if the edge was followed in the forward (tail to head) or reverse (head to tail) direction. When the direction is forward, the value of direction is 'forward'. When the direction is reverse, the value of direction is 'reverse'.

# **Examples**

In this example, we construct a DAG and find, in the first call, that there are no directed cycles, and so an exception is raised. In the second call, we ignore edge orientations and find that there is an undirected cycle. Note that the second call finds a directed cycle while effectively traversing an undirected graph, and so, we found an "undirected cycle". This means that this DAG structure does not form a directed tree (which is also known as a polytree).

```
>>> import networkx as nx
>>> G = nx.DiGraph([(0,1), (0,2), (1,2)])
>>> try:
... find_cycle(G, orientation='original')
... except:
... pass
...
>>> list(find_cycle(G, orientation='ignore'))
[(0, 1, 'forward'), (1, 2, 'forward'), (0, 2, 'reverse')]
```

# 4.16 Directed Acyclic Graphs

ancestors(G, source)	Return all nodes having a path to <i>source</i> in G.
descendants(G, source)	Return all nodes reachable from <i>source</i> in G.
topological_sort(G[, nbunch, reverse])	Return a list of nodes in topological sort order.
topological_sort_recursive(G[, nbunch, reverse])	Return a list of nodes in topological sort order.
is_directed_acyclic_graph(G)	Return True if the graph G is a directed acyclic graph (DAG) or False
is_aperiodic(G)	Return True if G is aperiodic.

# 4.16.1 ancestors

ancestors(G, source)

Return all nodes having a path to *source* in G.

Parameters G: NetworkX DiGraph

source : node in G
Returns ancestors : set()

The ancestors of source in G

# 4.16.2 descendants

### descendants (G, source)

Return all nodes reachable from source in G.

Parameters G: NetworkX DiGraph

source : node in G
Returns des : set()

The descendants of source in G

# 4.16.3 topological sort

# topological\_sort (G, nbunch=None, reverse=False)

Return a list of nodes in topological sort order.

A topological sort is a nonunique permutation of the nodes such that an edge from u to v implies that u appears before v in the topological sort order.

# Parameters G: NetworkX digraph

A directed graph

**nbunch**: container of nodes (optional)

Explore graph in specified order given in nbunch

reverse: bool, optional

Return postorder instead of preorder if True. Reverse mode is a bit more efficient.

# Raises NetworkXError

Topological sort is defined for directed graphs only. If the graph G is undirected, a NetworkXError is raised.

#### NetworkXUnfeasible

If G is not a directed acyclic graph (DAG) no topological sort exists and a NetworkX-Unfeasible exception is raised.

# See also:

```
is_directed_acyclic_graph
```

#### **Notes**

This algorithm is based on a description and proof in The Algorithm Design Manual [R271].

### References

[R271]

# 4.16.4 topological sort recursive

### topological\_sort\_recursive(G, nbunch=None, reverse=False)

Return a list of nodes in topological sort order.

A topological sort is a nonunique permutation of the nodes such that an edge from u to v implies that u appears before v in the topological sort order.

Parameters G: NetworkX digraph

**nbunch**: container of nodes (optional)

Explore graph in specified order given in nbunch

reverse: bool, optional

Return postorder instead of preorder if True. Reverse mode is a bit more efficient.

#### Raises NetworkXError

Topological sort is defined for directed graphs only. If the graph G is undirected, a NetworkXError is raised.

#### NetworkXUnfeasible

If G is not a directed acyclic graph (DAG) no topological sort exists and a NetworkX-Unfeasible exception is raised.

#### See also:

```
topological_sort, is_directed_acyclic_graph
```

#### **Notes**

This is a recursive version of topological sort.

# 4.16.5 is directed acyclic graph

# $is\_directed\_acyclic\_graph(G)$

Return True if the graph G is a directed acyclic graph (DAG) or False if not.

**Parameters G**: NetworkX graph

A graph

Returns is\_dag: bool

True if G is a DAG, false otherwise

# 4.16.6 is aperiodic

#### $is\_aperiodic(G)$

Return True if G is aperiodic.

A directed graph is aperiodic if there is no integer k > 1 that divides the length of every cycle in the graph.

**Parameters G**: NetworkX DiGraph

Graph

Returns aperiodic: boolean

True if the graph is aperiodic False otherwise

### Raises NetworkXError

If G is not directed

#### **Notes**

This uses the method outlined in [R270], which runs in O(m) time given m edges in G. Note that a graph is not aperiodic if it is acyclic as every integer trivial divides length 0 cycles.

#### References

[R270]

# 4.17 Distance Measures

Graph diameter, radius, eccentricity and other properties.

center(G[,e])	Return the center of the graph G.
diameter(G[,e])	Return the diameter of the graph G.
eccentricity( $G[, v, sp]$ )	Return the eccentricity of nodes in G.
periphery(G[,e])	Return the periphery of the graph G.
radius(G[,e])	Return the radius of the graph G.

# 4.17.1 center

center(G, e=None)

Return the center of the graph G.

The center is the set of nodes with eccentricity equal to radius.

Parameters G: NetworkX graph

A graph

 $\mathbf{e}$ : eccentricity dictionary, optional

A precomputed dictionary of eccentricities.

Returns c: list

List of nodes in center

# 4.17.2 diameter

diameter(G, e=None)

Return the diameter of the graph G.

The diameter is the maximum eccentricity.

Parameters G: NetworkX graph

A graph

e: eccentricity dictionary, optional

A precomputed dictionary of eccentricities.

Returns d: integer

Diameter of graph

See also:

eccentricity

# 4.17.3 eccentricity

eccentricity(G, v=None, sp=None)

Return the eccentricity of nodes in G.

The eccentricity of a node v is the maximum distance from v to all other nodes in G.

**Parameters G**: NetworkX graph

A graph

v: node, optional

Return value of specified node

sp: dict of dicts, optional

All pairs shortest path lengths as a dictionary of dictionaries

**Returns ecc**: dictionary

A dictionary of eccentricity values keyed by node.

# 4.17.4 periphery

periphery(G, e=None)

Return the periphery of the graph G.

The periphery is the set of nodes with eccentricity equal to the diameter.

**Parameters G**: NetworkX graph

A graph

e: eccentricity dictionary, optional

A precomputed dictionary of eccentricities.

Returns p: list

List of nodes in periphery

# 4.17.5 radius

radius(G, e=None)

Return the radius of the graph G.

The radius is the minimum eccentricity.

**Parameters G**: NetworkX graph

A graph

e: eccentricity dictionary, optional

A precomputed dictionary of eccentricities.

Returns r: integer

Radius of graph

# 4.18 Distance-Regular Graphs

is_distance_regular(G)	Returns True if the graph is distance regular, False otherwise.
$intersection\_array(G)$	Returns the intersection array of a distance-regular graph.
global_parameters(b, c)	Return global parameters for a given intersection array.

# 4.18.1 is\_distance\_regular

# $is\_distance\_regular(G)$

Returns True if the graph is distance regular, False otherwise.

A connected graph G is distance-regular if for any nodes x,y and any integers i,j=0,1,...,d (where d is the graph diameter), the number of vertices at distance i from x and distance j from y depends only on i,j and the graph distance between x and y, independently of the choice of x and y.

# Parameters G: Networkx graph (undirected)

Returns bool

True if the graph is Distance Regular, False otherwise

# See also:

```
intersection_array, global_parameters
```

# **Notes**

For undirected and simple graphs only

# References

[R274], [R275]

# **Examples**

```
>>> G=nx.hypercube_graph(6)
>>> nx.is_distance_regular(G)
True
```

# 4.18.2 intersection array

### $intersection\_array(G)$

Returns the intersection array of a distance-regular graph.

Given a distance-regular graph G with integers  $b_i$ ,  $c_i$ , i = 0,...,d such that for any 2 vertices x,y in G at a distance i=d(x,y), there are exactly  $c_i$  neighbors of y at a distance of i-1 from x and  $b_i$  neighbors of y at a distance of i+1 from x.

A distance regular graph's intersection array is given by, [b\_0,b\_1,....b\_{d-1};c\_1,c\_2,....c\_d]

# **Parameters G: Networkx graph (undirected)**

Returns b,c: tuple of lists

#### See also:

```
global_parameters
```

#### References

[R273]

### **Examples**

```
>>> G=nx.icosahedral_graph()
>>> nx.intersection_array(G)
([5, 2, 1], [1, 2, 5])
```

# 4.18.3 global\_parameters

#### global\_parameters(b, c)

Return global parameters for a given intersection array.

Given a distance-regular graph G with integers  $b_i$ ,  $c_i$ , i = 0,...,d such that for any 2 vertices x,y in G at a distance i=d(x,y), there are exactly  $c_i$  neighbors of y at a distance of i-1 from x and  $b_i$  neighbors of y at a distance of i+1 from x.

Thus, a distance regular graph has the global parameters,  $[[c_0,a_0,b_0],[c_1,a_1,b_1],....,[c_d,a_d,b_d]]$  for the intersection array  $[b_0,b_1,....b_{d-1};c_1,c_2,....c_d]$  where  $a_i+b_i+c_i=k$ , k=degree of every vertex.

# Parameters b,c: tuple of lists

**Returns p** : list of three-tuples

#### See also:

```
intersection array
```

#### References

[R272]

# **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> b,c=nx.intersection_array(G)
>>> list(nx.global_parameters(b,c))
[(0, 0, 3), (1, 0, 2), (1, 1, 1), (1, 1, 1), (2, 0, 1), (3, 0, 0)]
```

# 4.19 Dominance

Dominance algorithms.

<pre>immediate_dominators(G, start)</pre>	Returns the immediate dominators of all nodes of a directed graph.
<pre>dominance_frontiers(G, start)</pre>	Returns the dominance frontiers of all nodes of a directed graph.

# 4.19.1 immediate\_dominators

### immediate\_dominators(G, start)

Returns the immediate dominators of all nodes of a directed graph.

Parameters G: a DiGraph or MultiDiGraph

The graph where dominance is to be computed.

start: node

The start node of dominance computation.

Returns idom: dict keyed by nodes

A dict containing the immediate dominators of each node reachable from start.

### Raises NetworkXNotImplemented

If G is undirected.

## NetworkXError

If start is not in G.

# **Notes**

Except for start, the immediate dominators are the parents of their corresponding nodes in the dominator tree.

# References

[R277]

# **Examples**

```
>>> G = nx.DiGraph([(1, 2), (1, 3), (2, 5), (3, 4), (4, 5)])
>>> sorted(nx.immediate_dominators(G, 1).items())
[(1, 1), (2, 1), (3, 1), (4, 3), (5, 1)]
```

4.19. Dominance 243

# 4.19.2 dominance frontiers

### dominance\_frontiers(G, start)

Returns the dominance frontiers of all nodes of a directed graph.

Parameters G: a DiGraph or MultiDiGraph

The graph where dominance is to be computed.

start: node

The start node of dominance computation.

**Returns df**: dict keyed by nodes

A dict containing the dominance frontiers of each node reachable from start as lists.

# Raises NetworkXNotImplemented

If G is undirected.

# NetworkXError

If start is not in G.

#### References

[R276]

### **Examples**

```
>>> G = nx.DiGraph([(1, 2), (1, 3), (2, 5), (3, 4), (4, 5)])
>>> sorted((u, sorted(df)) for u, df in nx.dominance_frontiers(G, 1).items())
[(1, []), (2, [5]), (3, [5]), (4, [5]), (5, [])]
```

# 4.20 Dominating Sets

$dominating\_set(G[, start\_with])$	Finds a dominating set for the graph G.
<pre>is_dominating_set(G, nbunch)</pre>	Checks if nodes in nbunch are a dominating set for G.

# 4.20.1 dominating\_set

```
dominating_set (G, start_with=None)
```

Finds a dominating set for the graph G.

A dominating set for a graph G = (V, E) is a node subset D of V such that every node not in D is adjacent to at least one member of D [R278].

Parameters G: NetworkX graph

start\_with : Node (default=None)

Node to use as a starting point for the algorithm.

Returns D: set

A dominating set for G.

### See also:

```
is_dominating_set
```

### **Notes**

This function is an implementation of algorithm 7 in [R279] which finds some dominating set, not necessarily the smallest one.

#### References

[R278], [R279]

# 4.20.2 is dominating set

# $\verb|is_dominating_set|(G, nbunch)|$

Checks if nodes in nbunch are a dominating set for G.

A dominating set for a graph G = (V, E) is a node subset D of V such that every node not in D is adjacent to at least one member of D [R280].

Parameters G: NetworkX graph

nbunch: Node container

### See also:

dominating\_set

### References

[R280]

# 4.21 Eulerian

Eulerian circuits and graphs.

is_eulerian(G)	Return True if G is an Eulerian graph, False otherwise.
$eulerian\_circuit(G[, source])$	Return the edges of an Eulerian circuit in G.

# 4.21.1 is\_eulerian

# $is\_eulerian(G)$

Return True if G is an Eulerian graph, False otherwise.

An Eulerian graph is a graph with an Eulerian circuit.

Parameters G: graph

A NetworkX Graph

4.21. Eulerian 245

#### **Notes**

This implementation requires the graph to be connected (or strongly connected for directed graphs).

# **Examples**

```
>>> nx.is_eulerian(nx.DiGraph({0:[3], 1:[2], 2:[3], 3:[0, 1]}))
True
>>> nx.is_eulerian(nx.complete_graph(5))
True
>>> nx.is_eulerian(nx.petersen_graph())
False
```

# 4.21.2 eulerian circuit

```
eulerian circuit(G, source=None)
```

Return the edges of an Eulerian circuit in G.

An Eulerian circuit is a path that crosses every edge in G exactly once and finishes at the starting node.

Parameters G: NetworkX Graph or DiGraph

A directed or undirected graph

source: node, optional

Starting node for circuit.

Returns edges: generator

A generator that produces edges in the Eulerian circuit.

#### Raises NetworkXError

If the graph is not Eulerian.

# See also:

```
is_eulerian
```

# **Notes**

Linear time algorithm, adapted from [R281]. General information about Euler tours [R282].

## References

```
[R281], [R282]
```

# **Examples**

```
>>> G=nx.complete_graph(3)
>>> list(nx.eulerian_circuit(G))
[(0, 2), (2, 1), (1, 0)]
>>> list(nx.eulerian_circuit(G, source=1))
[(1, 2), (2, 0), (0, 1)]
```

```
>>> [u for u,v in nx.eulerian_circuit(G)] # nodes in circuit
[0, 2, 1]
```

## **4.22 Flows**

## 4.22.1 Maximum Flow

<pre>maximum_flow(G, s, t[, capacity, flow_func])</pre>	Find a maximum single-commodity flow.
$maximum\_flow\_value(G, s, t[, capacity,])$	Find the value of maximum single-commodity flow.
<pre>minimum_cut(G, s, t[, capacity, flow_func])</pre>	Compute the value and the node partition of a minimum (s, t)-cut.
<pre>minimum_cut_value(G, s, t[, capacity, flow_func])</pre>	Compute the value of a minimum (s, t)-cut.

## maximum flow

**maximum\_flow** (*G*, *s*, *t*, *capacity='capacity'*, *flow\_func=None*, \*\*kwargs) Find a maximum single-commodity flow.

## Parameters G: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

#### $\mathbf{s}$ : node

Source node for the flow.

#### t: node

Sink node for the flow.

## capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

#### flow\_func : function

A function for computing the maximum flow among a pair of nodes in a capacitated graph. The function has to accept at least three parameters: a Graph or Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see Notes). If flow\_func is None, the default maximum flow function (preflow\_push()) is used. See below for alternative algorithms. The choice of the default function may change from version to version and should not be relied on. Default value: None.

kwargs: Any other keyword parameter is passed to the function that

computes the maximum flow.

## Returns flow\_value : integer, float

Value of the maximum flow, i.e., net outflow from the source.

flow\_dict: dict

A dictionary containing the value of the flow that went through each edge.

#### Raises NetworkXError

The algorithm does not support MultiGraph and MultiDiGraph. If the input graph is an instance of one of these two classes, a NetworkXError is raised.

#### NetworkXUnbounded

If the graph has a path of infinite capacity, the value of a feasible flow on the graph is unbounded above and the function raises a NetworkXUnbounded.

#### See also:

```
maximum_flow_value(), minimum_cut(), minimum_cut_value(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

#### **Notes**

The function used in the flow\_func paramter has to return a residual network that follows NetworkX conventions:

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. Reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

Specific algorithms may store extra data in R.

The function should supports an optional boolean parameter value\_only. When True, it can optionally terminate the algorithm as soon as the maximum flow value and the minimum cut can be determined.

## **Examples**

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity=3.0)
>>> G.add_edge('x','b', capacity=1.0)
>>> G.add_edge('a','c', capacity=3.0)
>>> G.add_edge('b','c', capacity=5.0)
>>> G.add_edge('b','d', capacity=4.0)
>>> G.add_edge('d','e', capacity=2.0)
>>> G.add_edge('c','y', capacity=2.0)
>>> G.add_edge('e','y', capacity=3.0)
```

maximum\_flow returns both the value of the maximum flow and a dictionary with all flows.

```
>>> flow_value, flow_dict = nx.maximum_flow(G, 'x', 'y')
>>> flow_value
3.0
>>> print(flow_dict['x']['b'])
1.0
```

You can also use alternative algorithms for computing the maximum flow by using the flow\_func parameter.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> flow_value == nx.maximum_flow(G, 'x', 'y',
...
flow_func=shortest_augmenting_path)[0]
True
```

## maximum\_flow\_value

```
maximum_flow_value (G, s, t, capacity='capacity', flow_func=None, **kwargs) Find the value of maximum single-commodity flow.
```

## Parameters G: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

s: node

Source node for the flow.

t: node

Sink node for the flow.

capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

### flow\_func : function

A function for computing the maximum flow among a pair of nodes in a capacitated graph. The function has to accept at least three parameters: a Graph or Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see Notes). If flow\_func is None, the default maximum flow function (preflow\_push()) is used. See below for alternative algorithms. The choice of the default function may change from version to version and should not be relied on. Default value: None.

**kwargs**: Any other keyword parameter is passed to the function that

computes the maximum flow.

### Returns flow\_value: integer, float

Value of the maximum flow, i.e., net outflow from the source.

## Raises NetworkXError

The algorithm does not support MultiGraph and MultiDiGraph. If the input graph is an instance of one of these two classes, a NetworkXError is raised.

#### NetworkXUnbounded

If the graph has a path of infinite capacity, the value of a feasible flow on the graph is unbounded above and the function raises a NetworkXUnbounded.

### See also:

```
maximum_flow(), minimum_cut(), minimum_cut_value(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

#### **Notes**

The function used in the flow\_func paramter has to return a residual network that follows NetworkX conventions:

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. Reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

Specific algorithms may store extra data in R.

The function should supports an optional boolean parameter value\_only. When True, it can optionally terminate the algorithm as soon as the maximum flow value and the minimum cut can be determined.

## **Examples**

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity=3.0)
>>> G.add_edge('x','b', capacity=1.0)
>>> G.add_edge('a','c', capacity=3.0)
>>> G.add_edge('b','c', capacity=5.0)
>>> G.add_edge('b','d', capacity=4.0)
>>> G.add_edge('d','e', capacity=2.0)
>>> G.add_edge('c','y', capacity=2.0)
>>> G.add_edge('e','y', capacity=3.0)
```

maximum\_flow\_value computes only the value of the maximum flow:

```
>>> flow_value = nx.maximum_flow_value(G, 'x', 'y')
>>> flow_value
3.0
```

You can also use alternative algorithms for computing the maximum flow by using the flow func parameter.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> flow_value == nx.maximum_flow_value(G, 'x', 'y',
...
flow_func=shortest_augmenting_path)
True
```

## minimum cut

```
minimum_cut (G, s, t, capacity='capacity', flow_func=None, **kwargs)
```

Compute the value and the node partition of a minimum (s, t)-cut.

Use the max-flow min-cut theorem, i.e., the capacity of a minimum capacity cut is equal to the flow value of a maximum flow.

**Parameters G**: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

#### s: node

Source node for the flow.

#### t: node

Sink node for the flow.

### capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

#### flow func: function

A function for computing the maximum flow among a pair of nodes in a capacitated graph. The function has to accept at least three parameters: a Graph or Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see Notes). If flow\_func is None, the default maximum flow function (preflow\_push()) is used. See below for alternative algorithms. The choice of the default function may change from version to version and should not be relied on. Default value: None.

**kwargs**: Any other keyword parameter is passed to the function that

computes the maximum flow.

### Returns cut\_value: integer, float

Value of the minimum cut.

partition: pair of node sets

A partitioning of the nodes that defines a minimum cut.

### Raises NetworkXUnbounded

If the graph has a path of infinite capacity, all cuts have infinite capacity and the function raises a NetworkXError.

#### See also:

```
maximum_flow(), maximum_flow_value(), minimum_cut_value(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

#### **Notes**

The function used in the flow\_func paramter has to return a residual network that follows NetworkX conventions:

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. Reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

Specific algorithms may store extra data in R.

The function should supports an optional boolean parameter value\_only. When True, it can optionally terminate the algorithm as soon as the maximum flow value and the minimum cut can be determined.

### **Examples**

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity = 3.0)
>>> G.add_edge('x','b', capacity = 1.0)
>>> G.add_edge('a','c', capacity = 3.0)
>>> G.add_edge('b','c', capacity = 5.0)
>>> G.add_edge('b','d', capacity = 4.0)
>>> G.add_edge('d','e', capacity = 2.0)
>>> G.add_edge('c','y', capacity = 2.0)
>>> G.add_edge('e','y', capacity = 3.0)
```

minimum\_cut computes both the value of the minimum cut and the node partition:

```
>>> cut_value, partition = nx.minimum_cut(G, 'x', 'y')
>>> reachable, non_reachable = partition
```

'partition' here is a tuple with the two sets of nodes that define the minimum cut. You can compute the cut set of edges that induce the minimum cut as follows:

```
>>> cutset = set()
>>> for u, nbrs in ((n, G[n]) for n in reachable):
...    cutset.update((u, v) for v in nbrs if v in non_reachable)
>>> print(sorted(cutset))
[('c', 'y'), ('x', 'b')]
>>> cut_value == sum(G.edge[u][v]['capacity'] for (u, v) in cutset)
True
```

You can also use alternative algorithms for computing the minimum cut by using the flow\_func parameter.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> cut_value == nx.minimum_cut(G, 'x', 'y',
...
flow_func=shortest_augmenting_path)[0]
True
```

## minimum cut value

```
minimum_cut_value (G, s, t, capacity='capacity', flow_func=None, **kwargs)

Compute the value of a minimum (s, t)-cut.
```

Use the max-flow min-cut theorem, i.e., the capacity of a minimum capacity cut is equal to the flow value of a maximum flow.

#### **Parameters G**: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

```
s: node
```

Source node for the flow.

#### t: node

Sink node for the flow.

## capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

#### flow\_func : function

A function for computing the maximum flow among a pair of nodes in a capacitated graph. The function has to accept at least three parameters: a Graph or Digraph, a source node, and a target node. And return a residual network that follows NetworkX conventions (see Notes). If flow\_func is None, the default maximum flow function (preflow\_push()) is used. See below for alternative algorithms. The choice of the default function may change from version to version and should not be relied on. Default value: None.

**kwargs**: Any other keyword parameter is passed to the function that computes the maximum flow.

### Returns cut\_value: integer, float

Value of the minimum cut.

#### Raises NetworkXUnbounded

If the graph has a path of infinite capacity, all cuts have infinite capacity and the function raises a NetworkXError.

### See also:

```
maximum_flow(), maximum_flow_value(), minimum_cut(), edmonds_karp(),
preflow_push(), shortest_augmenting_path()
```

#### **Notes**

The function used in the flow\_func paramter has to return a residual network that follows NetworkX conventions:

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. Reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

Specific algorithms may store extra data in R.

The function should supports an optional boolean parameter value\_only. When True, it can optionally terminate the algorithm as soon as the maximum flow value and the minimum cut can be determined.

#### **Examples**

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity = 3.0)
>>> G.add_edge('x','b', capacity = 1.0)
>>> G.add_edge('a','c', capacity = 3.0)
>>> G.add_edge('b','c', capacity = 5.0)
>>> G.add_edge('b','d', capacity = 4.0)
>>> G.add_edge('d','e', capacity = 2.0)
>>> G.add_edge('c','y', capacity = 2.0)
>>> G.add_edge('e','y', capacity = 3.0)
```

minimum\_cut\_value computes only the value of the minimum cut:

```
>>> cut_value = nx.minimum_cut_value(G, 'x', 'y')
>>> cut_value
3.0
```

You can also use alternative algorithms for computing the minimum cut by using the flow\_func parameter.

```
>>> from networkx.algorithms.flow import shortest_augmenting_path
>>> cut_value == nx.minimum_cut_value(G, 'x', 'y',
...
flow_func=shortest_augmenting_path)
True
```

## 4.22.2 Edmonds-Karp

edmonds\_karp(G, s, t[, capacity, residual, ...]) Find a maximum single-commodity flow using the Edmonds-Karp algorithm.

### edmonds karp

**edmonds\_karp** (*G*, *s*, *t*, *capacity='capacity'*, *residual=None*, *value\_only=False*, *cutoff=None*) Find a maximum single-commodity flow using the Edmonds-Karp algorithm.

This function returns the residual network resulting after computing the maximum flow. See below for details about the conventions NetworkX uses for defining residual networks.

This algorithm has a running time of  $O(nm^2)$  for n nodes and m edges.

### **Parameters G**: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

s : node

Source node for the flow.

t: node

Sink node for the flow.

capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

#### residual: NetworkX graph

Residual network on which the algorithm is to be executed. If None, a new residual network is created. Default value: None.

### value\_only: bool

If True compute only the value of the maximum flow. This parameter will be ignored by this algorithm because it is not applicable.

#### cutoff: integer, float

If specified, the algorithm will terminate when the flow value reaches or exceeds the cutoff. In this case, it may be unable to immediately determine a minimum cut. Default value: None.

### Returns R: NetworkX DiGraph

Residual network after computing the maximum flow.

#### Raises NetworkXError

The algorithm does not support MultiGraph and MultiDiGraph. If the input graph is an instance of one of these two classes, a NetworkXError is raised.

#### NetworkXUnbounded

If the graph has a path of infinite capacity, the value of a feasible flow on the graph is unbounded above and the function raises a NetworkXUnbounded.

#### See also:

```
maximum_flow(), minimum_cut(), preflow_push(), shortest_augmenting_path()
```

#### **Notes**

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. If cutoff is not specified, reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

## **Examples**

```
>>> import networkx as nx
>>> from networkx.algorithms.flow import edmonds_karp
```

The functions that implement flow algorithms and output a residual network, such as this one, are not imported to the base NetworkX namespace, so you have to explicitly import them from the flow package.

```
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity=3.0)
>>> G.add_edge('x','b', capacity=1.0)
>>> G.add_edge('a','c', capacity=3.0)
>>> G.add_edge('b','c', capacity=5.0)
>>> G.add_edge('b','d', capacity=4.0)
>>> G.add_edge('d','e', capacity=2.0)
>>> G.add_edge('c','y', capacity=2.0)
>>> G.add_edge('c','y', capacity=3.0)
>>> R = edmonds_karp(G, 'x', 'y')
>>> flow_value = nx.maximum_flow_value(G, 'x', 'y')
>>> flow_value = R.graph['flow_value']
True
```

## 4.22.3 Shortest Augmenting Path

shortest\_augmenting\_path(G, s, t[, ...]) Find a maximum single-commodity flow using the shortest augmenting path algorithms.

## shortest\_augmenting\_path

```
\begin{tabular}{ll} \textbf{shortest\_augmenting\_path} (G, s, t, capacity='capacity', residual=None, value\_only=False, \\ two\_phase=False, cutoff=None) \end{tabular}
```

Find a maximum single-commodity flow using the shortest augmenting path algorithm.

This function returns the residual network resulting after computing the maximum flow. See below for details about the conventions NetworkX uses for defining residual networks.

This algorithm has a running time of  $O(n^2m)$  for n nodes and m edges.

## Parameters G: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

s: node

Source node for the flow.

t: node

Sink node for the flow.

capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

residual: NetworkX graph

Residual network on which the algorithm is to be executed. If None, a new residual network is created. Default value: None.

value\_only: bool

If True compute only the value of the maximum flow. This parameter will be ignored by this algorithm because it is not applicable.

#### two\_phase: bool

If True, a two-phase variant is used. The two-phase variant improves the running time on unit-capacity networks from O(nm) to  $O(\min(n^{2/3}, m^{1/2})m)$ . Default value: False.

#### cutoff: integer, float

If specified, the algorithm will terminate when the flow value reaches or exceeds the cutoff. In this case, it may be unable to immediately determine a minimum cut. Default value: None.

## Returns R: NetworkX DiGraph

Residual network after computing the maximum flow.

#### Raises NetworkXError

The algorithm does not support MultiGraph and MultiDiGraph. If the input graph is an instance of one of these two classes, a NetworkXError is raised.

#### NetworkXUnbounded

If the graph has a path of infinite capacity, the value of a feasible flow on the graph is unbounded above and the function raises a NetworkXUnbounded.

#### See also:

```
maximum_flow(), minimum_cut(), edmonds_karp(), preflow_push()
```

#### **Notes**

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. If cutoff is not specified, reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

### **Examples**

```
>>> import networkx as nx
>>> from networkx.algorithms.flow import shortest_augmenting_path
```

The functions that implement flow algorithms and output a residual network, such as this one, are not imported to the base NetworkX namespace, so you have to explicitly import them from the flow package.

```
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity=3.0)
>>> G.add_edge('x','b', capacity=1.0)
>>> G.add_edge('a','c', capacity=3.0)
>>> G.add_edge('b','c', capacity=5.0)
>>> G.add_edge('b','d', capacity=4.0)
```

```
>>> G.add_edge('d','e', capacity=2.0)
>>> G.add_edge('c','y', capacity=2.0)
>>> G.add_edge('e','y', capacity=3.0)
>>> R = shortest_augmenting_path(G, 'x', 'y')
>>> flow_value = nx.maximum_flow_value(G, 'x', 'y')
>>> flow_value
3.0
>>> flow_value == R.graph['flow_value']
True
```

### 4.22.4 Preflow-Push

preflow\_push(G, s, t[, capacity, residual, ...]) Find a maximum single-commodity flow using the highest-label preflow-push algorithms are similar to the sin

## preflow push

 $preflow_push(G, s, t, capacity='capacity', residual=None, global_relabel_freq=1, value_only=False)$  Find a maximum single-commodity flow using the highest-label preflow-push algorithm.

This function returns the residual network resulting after computing the maximum flow. See below for details about the conventions NetworkX uses for defining residual networks.

This algorithm has a running time of  $O(n^2\sqrt{m})$  for n nodes and m edges.

## Parameters G: NetworkX graph

Edges of the graph are expected to have an attribute called 'capacity'. If this attribute is not present, the edge is considered to have infinite capacity.

s: node

Source node for the flow.

t: node

Sink node for the flow.

capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

residual: NetworkX graph

Residual network on which the algorithm is to be executed. If None, a new residual network is created. Default value: None.

```
global_relabel_freq: integer, float
```

Relative frequency of applying the global relabeling heuristic to speed up the algorithm. If it is None, the heuristic is disabled. Default value: 1.

```
value_only: bool
```

If False, compute a maximum flow; otherwise, compute a maximum preflow which is enough for computing the maximum flow value. Default value: False.

#### **Returns R**: NetworkX DiGraph

Residual network after computing the maximum flow.

#### Raises NetworkXError

The algorithm does not support MultiGraph and MultiDiGraph. If the input graph is an instance of one of these two classes, a NetworkXError is raised.

#### NetworkXUnbounded

If the graph has a path of infinite capacity, the value of a feasible flow on the graph is unbounded above and the function raises a NetworkXUnbounded.

#### See also:

```
maximum_flow(), minimum_cut(), edmonds_karp(), shortest_augmenting_path()
```

#### **Notes**

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G. For each node u in R, R. node [u] ['excess'] represents the difference between flow into u and flow out of u.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. Reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

#### **Examples**

```
>>> import networkx as nx
>>> from networkx.algorithms.flow import preflow_push
```

The functions that implement flow algorithms and output a residual network, such as this one, are not imported to the base NetworkX namespace, so you have to explicitly import them from the flow package.

```
>>> G = nx.DiGraph()
>>> G.add_edge('x','a', capacity=3.0)
>>> G.add_edge('x','b', capacity=1.0)
>>> G.add_edge('a','c', capacity=3.0)
>>> G.add_edge('b','c', capacity=5.0)
>>> G.add_edge('b','d', capacity=4.0)
>>> G.add_edge('d','e', capacity=2.0)
>>> G.add_edge('c','y', capacity=2.0)
>>> G.add_edge('e','y', capacity=3.0)
>>> R = preflow_push(G, 'x', 'y')
>>> flow_value = nx.maximum_flow_value(G, 'x', 'y')
>>> flow_value == R.graph['flow_value']
>>> # preflow push also stores the maximum flow value
>>> # in the excess attribute of the sink node t
>>> flow_value == R.node['y']['excess']
True
>>> # For some problems, you might only want to compute a
```

```
>>> # maximum preflow.
>>> R = preflow_push(G, 'x', 'y', value_only=True)
>>> flow_value == R.graph['flow_value']
True
>>> flow_value == R.node['y']['excess']
True
```

## 4.22.5 Utils

build residual network(G, capacity) Build a residual network and initialize a zero flow.

## build residual network

#### build residual network(G, capacity)

Build a residual network and initialize a zero flow.

The residual network R from an input graph G has the same nodes as G. R is a DiGraph that contains a pair of edges (u, v) and (v, u) iff (u, v) is not a self-loop, and at least one of (u, v) and (v, u) exists in G.

For each edge (u, v) in R, R[u][v]['capacity'] is equal to the capacity of (u, v) in G if it exists in G or zero otherwise. If the capacity is infinite, R[u][v]['capacity'] will have a high arbitrary finite value that does not affect the solution of the problem. This value is stored in R.graph['inf']. For each edge (u, v) in R, R[u][v]['flow'] represents the flow function of (u, v) and satisfies R[u][v]['flow'] = -R[v][u]['flow'].

The flow value, defined as the total flow into t, the sink, is stored in R.graph['flow\_value']. If cutoff is not specified, reachability to t using only edges (u, v) such that R[u][v]['flow'] < R[u][v]['capacity'] induces a minimum s-t cut.

## 4.22.6 Network Simplex

<pre>network_simplex(G[, demand, capacity, weight])</pre>	Find a minimum cost flow satisfying all demands in digraph G.
min_cost_flow_cost(G[, demand, capacity, weight])	Find the cost of a minimum cost flow satisfying all demands in digraph
min_cost_flow(G[, demand, capacity, weight])	Return a minimum cost flow satisfying all demands in digraph G.
<pre>cost_of_flow(G, flowDict[, weight])</pre>	Compute the cost of the flow given by flowDict on graph G.
<pre>max_flow_min_cost(G, s, t[, capacity, weight])</pre>	Return a maximum (s, t)-flow of minimum cost.

## network\_simplex

 $\verb"network_simplex" (G, demand='demand', capacity='capacity', weight='weight')$ 

Find a minimum cost flow satisfying all demands in digraph G.

This is a primal network simplex algorithm that uses the leaving arc rule to prevent cycling.

G is a digraph with edge costs and capacities and in which nodes have demand, i.e., they want to send or receive some amount of flow. A negative demand means that the node wants to send flow, a positive demand means that the node want to receive flow. A flow on the digraph G satisfies all demand if the net flow into each node is equal to the demand of that node.

#### **Parameters G**: NetworkX graph

DiGraph on which a minimum cost flow satisfying all demands is to be found.

## demand: string

Nodes of the graph G are expected to have an attribute demand that indicates how much flow a node wants to send (negative demand) or receive (positive demand). Note that the sum of the demands should be 0 otherwise the problem in not feasible. If this attribute is not present, a node is considered to have 0 demand. Default value: 'demand'.

#### capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

## weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

## Returns flowCost: integer, float

Cost of a minimum cost flow satisfying all demands.

#### flowDict: dictionary

Dictionary of dictionaries keyed by nodes such that flowDict[u][v] is the flow edge (u, v).

#### Raises NetworkXError

This exception is raised if the input graph is not directed, not connected or is a multi-graph.

## NetworkXUnfeasible

## This exception is raised in the following situations:

- The sum of the demands is not zero. Then, there is no flow satisfying all demands.
- There is no flow satisfying all demand.

## NetworkXUnbounded

This exception is raised if the digraph G has a cycle of negative cost and infinite capacity. Then, the cost of a flow satisfying all demands is unbounded below.

#### See also:

```
cost_of_flow, max_flow_min_cost, min_cost_flow, min_cost_flow_cost
```

#### **Notes**

This algorithm is not guaranteed to work if edge weights are floating point numbers (overflows and roundoff errors can cause problems).

#### References

[R283], [R284]

#### **Examples**

A simple example of a min cost flow problem.

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_node('a', demand=-5)
>>> G.add_node('d', demand=5)
>>> G.add_edge('a', 'b', weight=3, capacity=4)
>>> G.add_edge('a', 'c', weight=6, capacity=10)
>>> G.add_edge('b', 'd', weight=1, capacity=9)
>>> G.add_edge('c', 'd', weight=2, capacity=5)
>>> flowCost, flowDict = nx.network_simplex(G)
>>> flowCost
24
>>> flowDict
{'a': {'c': 1, 'b': 4}, 'c': {'d': 1}, 'b': {'d': 4}, 'd': {}}}
```

The mincost flow algorithm can also be used to solve shortest path problems. To find the shortest path between two nodes u and v, give all edges an infinite capacity, give node u a demand of -1 and node v a demand a 1. Then run the network simplex. The value of a min cost flow will be the distance between u and v and edges carrying positive flow will indicate the path.

```
>>> G=nx.DiGraph()
>>> G.add_weighted_edges_from([('s', 'u' ,10), ('s' ,'x' ,5),
                                ('u', 'v',1), ('u','x',2),
                                ('v', 'y', 1), ('x', 'u', 3),
. . .
                                ('x', 'v', 5), ('x', 'y', 2),
. . .
                                ('y', 's',7), ('y','v',6)])
\rightarrow G.add_node('s', demand = -1)
>>> G.add_node('v', demand = 1)
>>> flowCost, flowDict = nx.network_simplex(G)
>>> flowCost == nx.shortest_path_length(G, 's', 'v', weight='weight')
>>> sorted([(u, v) for u in flowDict for v in flowDict[u] if flowDict[u][v] > 0])
[('s', 'x'), ('u', 'v'), ('x', 'u')]
>>> nx.shortest_path(G, 's', 'v', weight = 'weight')
['s', 'x', 'u', 'v']
```

It is possible to change the name of the attributes used for the algorithm.

```
>>> G = nx.DiGraph()
>>> G.add_node('p', spam=-4)
>>> G.add_node('q', spam=2)
>>> G.add_node('a', spam=-2)
>>> G.add_node('d', spam=-1)
>>> G.add_node('t', spam=2)
>>> G.add_node('w', spam=3)
>>> G.add_edge('p', 'q', cost=7, vacancies=5)
>>> G.add_edge('p', 'a', cost=1, vacancies=4)
>>> G.add_edge('q', 'd', cost=2, vacancies=3)
>>> G.add_edge('t', 'q', cost=1, vacancies=2)
>>> G.add_edge('a', 't', cost=2, vacancies=4)
>>> G.add_edge('d', 'w', cost=3, vacancies=4)
>>> G.add_edge('t', 'w', cost=4, vacancies=1)
>>> flowCost, flowDict = nx.network_simplex(G, demand='spam',
                                             capacity='vacancies',
. . .
                                             weight='cost')
. . .
>>> flowCost
```

```
37
>>> flowDict
{'a': {'t': 4}, 'd': {'w': 2}, 'q': {'d': 1}, 'p': {'q': 2, 'a': 2}, 't': {'q': 1, 'w': 1}, 'w':
```

#### min cost flow cost

min\_cost\_flow\_cost(G, demand='demand', capacity='capacity', weight='weight')

Find the cost of a minimum cost flow satisfying all demands in digraph G.

G is a digraph with edge costs and capacities and in which nodes have demand, i.e., they want to send or receive some amount of flow. A negative demand means that the node wants to send flow, a positive demand means that the node want to receive flow. A flow on the digraph G satisfies all demand if the net flow into each node is equal to the demand of that node.

## Parameters G: NetworkX graph

DiGraph on which a minimum cost flow satisfying all demands is to be found.

### demand: string

Nodes of the graph G are expected to have an attribute demand that indicates how much flow a node wants to send (negative demand) or receive (positive demand). Note that the sum of the demands should be 0 otherwise the problem in not feasible. If this attribute is not present, a node is considered to have 0 demand. Default value: 'demand'.

#### capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

### weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

### Returns flowCost: integer, float

Cost of a minimum cost flow satisfying all demands.

#### Raises NetworkXError

This exception is raised if the input graph is not directed or not connected.

### NetworkXUnfeasible

## This exception is raised in the following situations:

- The sum of the demands is not zero. Then, there is no flow satisfying all demands.
- There is no flow satisfying all demand.

#### NetworkXUnbounded

This exception is raised if the digraph G has a cycle of negative cost and infinite capacity. Then, the cost of a flow satisfying all demands is unbounded below.

## See also:

```
cost_of_flow, max_flow_min_cost, min_cost_flow, network_simplex
```

#### **Examples**

A simple example of a min cost flow problem.

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_node('a', demand = -5)
>>> G.add_node('d', demand = 5)
>>> G.add_edge('a', 'b', weight = 3, capacity = 4)
>>> G.add_edge('a', 'c', weight = 6, capacity = 10)
>>> G.add_edge('b', 'd', weight = 1, capacity = 9)
>>> G.add_edge('c', 'd', weight = 2, capacity = 5)
>>> flowCost = nx.min_cost_flow_cost(G)
>>> flowCost
```

## min\_cost\_flow

```
min_cost_flow (G, demand='demand', capacity='capacity', weight='weight')

Return a minimum cost flow satisfying all demands in digraph G.
```

G is a digraph with edge costs and capacities and in which nodes have demand, i.e., they want to send or receive some amount of flow. A negative demand means that the node wants to send flow, a positive demand means that the node want to receive flow. A flow on the digraph G satisfies all demand if the net flow into each node is equal to the demand of that node.

## **Parameters G**: NetworkX graph

DiGraph on which a minimum cost flow satisfying all demands is to be found.

#### demand: string

Nodes of the graph G are expected to have an attribute demand that indicates how much flow a node wants to send (negative demand) or receive (positive demand). Note that the sum of the demands should be 0 otherwise the problem in not feasible. If this attribute is not present, a node is considered to have 0 demand. Default value: 'demand'.

#### capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

## weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

## Returns flowDict: dictionary

Dictionary of dictionaries keyed by nodes such that flowDict[u][v] is the flow edge (u, v).

## Raises NetworkXError

This exception is raised if the input graph is not directed or not connected.

## NetworkXUnfeasible

This exception is raised in the following situations:

- The sum of the demands is not zero. Then, there is no flow satisfying all demands.
- There is no flow satisfying all demand.

### NetworkXUnbounded

This exception is raised if the digraph G has a cycle of negative cost and infinite capacity. Then, the cost of a flow satisfying all demands is unbounded below.

## See also:

```
cost_of_flow, max_flow_min_cost, min_cost_flow_cost, network_simplex
```

#### **Examples**

A simple example of a min cost flow problem.

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_node('a', demand = -5)
>>> G.add_node('d', demand = 5)
>>> G.add_edge('a', 'b', weight = 3, capacity = 4)
>>> G.add_edge('a', 'c', weight = 6, capacity = 10)
>>> G.add_edge('b', 'd', weight = 1, capacity = 9)
>>> G.add_edge('c', 'd', weight = 2, capacity = 5)
>>> flowDict = nx.min_cost_flow(G)
```

#### cost of flow

```
cost_of_flow(G, flowDict, weight='weight')
```

Compute the cost of the flow given by flowDict on graph G.

Note that this function does not check for the validity of the flow flowDict. This function will fail if the graph G and the flow don't have the same edge set.

## Parameters G: NetworkX graph

DiGraph on which a minimum cost flow satisfying all demands is to be found.

#### weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

## flowDict: dictionary

Dictionary of dictionaries keyed by nodes such that flowDict[u][v] is the flow edge (u, v).

Returns cost: Integer, float

The total cost of the flow. This is given by the sum over all edges of the product of the edge's flow and the edge's weight.

#### See also:

```
max_flow_min_cost, min_cost_flow, min_cost_flow_cost, network_simplex
```

## max\_flow\_min\_cost

```
{\tt max\_flow\_min\_cost} (G, s, t, capacity = 'capacity', weight = 'weight')
```

Return a maximum (s, t)-flow of minimum cost.

G is a digraph with edge costs and capacities. There is a source node s and a sink node t. This function finds a maximum flow from s to t whose total cost is minimized.

## Parameters G: NetworkX graph

DiGraph on which a minimum cost flow satisfying all demands is to be found.

#### s: node label

Source of the flow.

#### t: node label

Destination of the flow.

## capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

### weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

### Returns flowDict: dictionary

Dictionary of dictionaries keyed by nodes such that flowDict[u][v] is the flow edge (u, v).

## Raises NetworkXError

This exception is raised if the input graph is not directed or not connected.

#### NetworkXUnbounded

This exception is raised if there is an infinite capacity path from s to t in G. In this case there is no maximum flow. This exception is also raised if the digraph G has a cycle of negative cost and infinite capacity. Then, the cost of a flow is unbounded below.

## See also:

```
\verb|cost_of_flow|, \verb|min_cost_flow|, \verb|min_cost_flow_cost|, \verb|network_simplex||
```

### **Examples**

## 4.22.7 Capacity Scaling Minimum Cost Flow

capacity\_scaling(G[, demand, capacity, ...]) Find a minimum cost flow satisfying all demands in digraph G.

## capacity\_scaling

```
capacity_scaling(G, demand='demand', capacity='capacity', weight='weight', heap=<class 'net-
workx.utils.heaps.BinaryHeap'>)
```

Find a minimum cost flow satisfying all demands in digraph G.

This is a capacity scaling successive shortest augmenting path algorithm.

G is a digraph with edge costs and capacities and in which nodes have demand, i.e., they want to send or receive some amount of flow. A negative demand means that the node wants to send flow, a positive demand means that the node want to receive flow. A flow on the digraph G satisfies all demand if the net flow into each node is equal to the demand of that node.

### **Parameters G**: NetworkX graph

DiGraph or MultiDiGraph on which a minimum cost flow satisfying all demands is to be found.

## demand: string

Nodes of the graph G are expected to have an attribute demand that indicates how much flow a node wants to send (negative demand) or receive (positive demand). Note that the sum of the demands should be 0 otherwise the problem in not feasible. If this attribute is not present, a node is considered to have 0 demand. Default value: 'demand'.

### capacity: string

Edges of the graph G are expected to have an attribute capacity that indicates how much flow the edge can support. If this attribute is not present, the edge is considered to have infinite capacity. Default value: 'capacity'.

### weight: string

Edges of the graph G are expected to have an attribute weight that indicates the cost incurred by sending one unit of flow on that edge. If not present, the weight is considered to be 0. Default value: 'weight'.

heap: class

Type of heap to be used in the algorithm. It should be a subclass of MinHeap or implement a compatible interface.

If a stock heap implementation is to be used, BinaryHeap is recommeded over PairingHeap for Python implementations without optimized attribute accesses (e.g., CPython) despite a slower asymptotic running time. For Python implementations with optimized attribute accesses (e.g., PyPy), PairingHeap provides better performance. Default value: BinaryHeap.

## Returns flowCost: integer

Cost of a minimum cost flow satisfying all demands.

### flowDict: dictionary

Dictionary of dictionaries keyed by nodes such that flowDict[u][v] is the flow edge (u, v) if G is a digraph.

Dictionary of dictionaries of dictionaries keyed by nodes such that flowDict[u][v][key] is the flow edge (u, v, key) if G is a multidigraph.

#### Raises NetworkXError

This exception is raised if the input graph is not directed, not connected.

#### NetworkXUnfeasible

### This exception is raised in the following situations:

- The sum of the demands is not zero. Then, there is no flow satisfying all demands.
- There is no flow satisfying all demand.

#### NetworkXUnbounded

This exception is raised if the digraph G has a cycle of negative cost and infinite capacity. Then, the cost of a flow satisfying all demands is unbounded below.

#### See also:

```
network_simplex()
```

#### **Notes**

This algorithm does not work if edge weights are floating-point numbers.

#### **Examples**

A simple example of a min cost flow problem.

```
>>> import networkx as nx
>>> G = nx.DiGraph()
>>> G.add_node('a', demand = -5)
>>> G.add_node('d', demand = 5)
>>> G.add_edge('a', 'b', weight = 3, capacity = 4)
>>> G.add_edge('a', 'c', weight = 6, capacity = 10)
>>> G.add_edge('b', 'd', weight = 1, capacity = 9)
>>> G.add_edge('c', 'd', weight = 2, capacity = 5)
>>> flowCost, flowDict = nx.capacity_scaling(G)
>>> flowCost
```

```
It is possible to change the name of the attributes used for the algorithm.
>>> G = nx.DiGraph()
>>> G.add_node('p', spam = -4)
>>> G.add_node('q', spam = 2)
>>> G.add_node('a', spam = -2)
\rightarrow \rightarrow G.add_node('d', spam = -1)
>>> G.add_node('t', spam = 2)
>>> G.add_node('w', spam = 3)
>>> G.add_edge('p', 'q', cost = 7, vacancies = 5)
>>> G.add_edge('p', 'a', cost = 1, vacancies = 4)
>>> G.add_edge('q', 'd', cost = 2, vacancies = 3)
>>> G.add_edge('t', 'q', cost = 1, vacancies = 2)
>>> G.add_edge('a', 't', cost = 2, vacancies = 4)
>>> G.add_edge('d', 'w', cost = 3, vacancies = 4)
>>> G.add_edge('t', 'w', cost = 4, vacancies = 1)
>>> flowCost, flowDict = nx.capacity_scaling(G, demand = 'spam',
                                                capacity = 'vacancies',
. . .
                                                weight = 'cost')
>>> flowCost
37
>>> flowDict
{'a': {'t': 4}, 'd': {'w': 2}, 'q': {'d': 1}, 'p': {'q': 2, 'a': 2}, 't': {'q': 1, 'w': 1}, 'w':
```

{'a': {'c': 1, 'b': 4}, 'c': {'d': 1}, 'b': {'d': 4}, 'd': {}}

# 4.23 Graphical degree sequence

Test sequences for graphiness.

>>> flowDict

<pre>is_graphical(sequence[, method])</pre>	Returns True if sequence is a valid degree sequence.
is_digraphical(in_sequence, out_sequence)	Returns True if some directed graph can realize the in- and out-degree so
is_multigraphical(sequence)	Returns True if some multigraph can realize the sequence.
is_pseudographical(sequence)	Returns True if some pseudograph can realize the sequence.
is_valid_degree_sequence_havel_hakimi()	Returns True if deg_sequence can be realized by a simple graph.
is_valid_degree_sequence_erdos_gallai()	Returns True if deg_sequence can be realized by a simple graph.

## 4.23.1 is graphical

is\_graphical (sequence, method='eg')

Returns True if sequence is a valid degree sequence.

A degree sequence is valid if some graph can realize it.

Parameters sequence: list or iterable container

A sequence of integer node degrees

method: "eg" | "hh"

The method used to validate the degree sequence. "eg" corresponds to the Erdős-Gallai algorithm, and "hh" to the Havel-Hakimi algorithm.

Returns valid: bool

True if the sequence is a valid degree sequence and False if not.

## References

Erdős-Gallai [EG1960], [choudum1986]

**Havel-Hakimi** [havel1955], [hakimi1962], [CL1996]

## **Examples**

```
>>> G = nx.path_graph(4)
>>> sequence = G.degree().values()
>>> nx.is_valid_degree_sequence(sequence)
True
```

## 4.23.2 is\_digraphical

### is\_digraphical (in\_sequence, out\_sequence)

Returns True if some directed graph can realize the in- and out-degree sequences.

Parameters in\_sequence: list or iterable container

A sequence of integer node in-degrees

out\_sequence : list or iterable container

A sequence of integer node out-degrees

Returns valid: bool

True if in and out-sequences are digraphic False if not.

#### **Notes**

This algorithm is from Kleitman and Wang [R285]. The worst case runtime is O(s \* log n) where s and n are the sum and length of the sequences respectively.

#### References

[R285]

## 4.23.3 is\_multigraphical

### is\_multigraphical(sequence)

Returns True if some multigraph can realize the sequence.

Parameters deg\_sequence : list

A list of integers

Returns valid: bool

True if deg\_sequence is a multigraphic degree sequence and False if not.

#### **Notes**

The worst-case run time is O(n) where n is the length of the sequence.

#### References

[R286]

## 4.23.4 is\_pseudographical

## is\_pseudographical(sequence)

Returns True if some pseudograph can realize the sequence.

Every nonnegative integer sequence with an even sum is pseudographical (see [R287]).

Parameters sequence: list or iterable container

A sequence of integer node degrees

Returns valid: bool

True if the sequence is a pseudographic degree sequence and False if not.

#### **Notes**

The worst-case run time is O(n) where n is the length of the sequence.

#### References

[R287]

## 4.23.5 is\_valid\_degree\_sequence\_havel\_hakimi

### is\_valid\_degree\_sequence\_havel\_hakimi (deg\_sequence)

Returns True if deg\_sequence can be realized by a simple graph.

The validation proceeds using the Havel-Hakimi theorem. Worst-case run time is: O(s) where s is the sum of the sequence.

Parameters deg\_sequence : list

A list of integers where each element specifies the degree of a node in a graph.

Returns valid: bool

True if deg\_sequence is graphical and False if not.

#### **Notes**

The ZZ condition says that for the sequence d if

$$|d| > = \frac{(\max(d) + \min(d) + 1)^2}{4 * \min(d)}$$

then d is graphical. This was shown in Theorem 6 in [R290].

#### References

[havel1955], [hakimi1962], [CL1996] [R290]

## 4.23.6 is valid degree sequence erdos gallai

is\_valid\_degree\_sequence\_erdos\_gallai(deg\_sequence)

Returns True if deg\_sequence can be realized by a simple graph.

The validation is done using the Erdős-Gallai theorem [EG1960].

Parameters deg\_sequence: list

A list of integers

Returns valid: bool

True if deg\_sequence is graphical and False if not.

#### **Notes**

This implementation uses an equivalent form of the Erdős-Gallai criterion. Worst-case run time is: O(n) where n is the length of the sequence.

Specifically, a sequence d is graphical if and only if the sum of the sequence is even and for all strong indices k in the sequence,

$$\sum_{i=1}^{k} d_i \le k(k-1) + \sum_{j=k+1}^{n} \min(d_i, k) = k(n-1) - (k \sum_{j=0}^{k-1} n_j - \sum_{j=0}^{k-1} j n_j)$$

A strong index k is any index where  $d_k \ge k$  and the value  $n_j$  is the number of occurrences of j in d. The maximal strong index is called the Durfee index.

This particular rearrangement comes from the proof of Theorem 3 in [R289].

The ZZ condition says that for the sequence d if

$$|d| > = \frac{(\max(d) + \min(d) + 1)^2}{4 * \min(d)}$$

then d is graphical. This was shown in Theorem 6 in [R289].

#### References

[EG1960], [choudum1986]

[R288], [R289]

# 4.24 Hierarchy

Flow Hierarchy.

 $flow_hierarchy(G[, weight])$  Returns the flow hierarchy of a directed network.

## 4.24.1 flow hierarchy

## flow\_hierarchy (G, weight=None)

Returns the flow hierarchy of a directed network.

Flow hierarchy is defined as the fraction of edges not participating in cycles in a directed graph [R291].

Parameters G: DiGraph or MultiDiGraph

A directed graph

weight : key,optional (default=None)

Attribute to use for node weights. If None the weight defaults to 1.

Returns h: float

Flow heirarchy value

#### **Notes**

The algorithm described in [R291] computes the flow hierarchy through exponentiation of the adjacency matrix. This function implements an alternative approach that finds strongly connected components. An edge is in a cycle if and only if it is in a strongly connected component, which can be found in O(m) time using Tarjan's algorithm.

#### References

[R291]

## 4.25 Isolates

Functions for identifying isolate (degree zero) nodes.

is_isolate(G, n)	Determine of node n is an isolate (degree zero).
isolates(G)	Return list of isolates in the graph.

## 4.25.1 is\_isolate

## $is\_isolate(G, n)$

Determine of node n is an isolate (degree zero).

**Parameters G**: graph

A networkx graph

n: node

A node in G

Returns isolate: bool

4.25. Isolates 273

True if n has no neighbors, False otherwise.

## **Examples**

```
>>> G=nx.Graph()
>>> G.add_edge(1,2)
>>> G.add_node(3)
>>> nx.is_isolate(G,2)
False
>>> nx.is_isolate(G,3)
True
```

## **4.25.2** isolates

## isolates(G)

Return list of isolates in the graph.

Isolates are nodes with no neighbors (degree zero).

Parameters G: graph

A networkx graph

Returns isolates: list

List of isolate nodes.

## **Examples**

```
>>> G = nx.Graph()
>>> G.add_edge(1,2)
>>> G.add_node(3)
>>> nx.isolates(G)
[3]
```

To remove all isolates in the graph use >>> G.remove\_nodes\_from(nx.isolates(G)) >>> G.nodes() [1, 2]

For digraphs isolates have zero in-degree and zero out\_degre >>>  $G = nx.DiGraph([(0,1),(1,2)]) >>> G.add_node(3) >>> nx.isolates(G) [3]$ 

# 4.26 Isomorphism

is_isomorphic(G1, G2[, node_match, edge_match])	Returns True if the graphs G1 and G2 are isomorphic and False otherwise
could_be_isomorphic(G1, G2)	Returns False if graphs are definitely not isomorphic.
$fast\_could\_be\_isomorphic(G1, G2)$	Returns False if graphs are definitely not isomorphic.
faster_could_be_isomorphic(G1, G2)	Returns False if graphs are definitely not isomorphic.

## 4.26.1 is isomorphic

```
\verb|is_isomorphic| (G1, G2, node\_match=None, edge\_match=None)|
```

Returns True if the graphs G1 and G2 are isomorphic and False otherwise.

#### Parameters G1, G2: graphs

The two graphs G1 and G2 must be the same type.

```
node match: callable
```

A function that returns True if node n1 in G1 and n2 in G2 should be considered equal during the isomorphism test. If node\_match is not specified then node attributes are not considered.

The function will be called like

```
node_match(G1.node[n1], G2.node[n2]).
```

That is, the function will receive the node attribute dictionaries for n1 and n2 as inputs.

```
edge_match: callable
```

A function that returns True if the edge attribute dictionary for the pair of nodes (u1, v1) in G1 and (u2, v2) in G2 should be considered equal during the isomorphism test. If edge\_match is not specified then edge attributes are not considered.

The function will be called like

```
edge_match(G1[u1][v1], G2[u2][v2]).
```

That is, the function will receive the edge attribute dictionaries of the edges under consideration.

#### See also:

```
numerical_node_match, numerical_edge_match, numerical_multiedge_match, categorical_node_match, categorical_edge_match, categorical_multiedge_match
```

#### **Notes**

Uses the vf2 algorithm [R292].

### References

[R292]

## **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
```

For digraphs G1 and G2, using 'weight' edge attribute (default: 1)

```
>>> G1 = nx.DiGraph()
>>> G2 = nx.DiGraph()
>>> G1.add_path([1,2,3,4],weight=1)
>>> G2.add_path([10,20,30,40],weight=2)
>>> em = iso.numerical_edge_match('weight', 1)
>>> nx.is_isomorphic(G1, G2) # no weights considered
True
>>> nx.is_isomorphic(G1, G2, edge_match=em) # match weights
False
```

For multidigraphs G1 and G2, using 'fill' node attribute (default: ")

```
>>> G1 = nx.MultiDiGraph()
>>> G2 = nx.MultiDiGraph()
>>> G1.add_nodes_from([1,2,3],fill='red')
>>> G2.add_nodes_from([10,20,30,40],fill='red')
>>> G1.add_path([1,2,3,4],weight=3, linewidth=2.5)
>>> G2.add_path([10,20,30,40],weight=3)
>>> nm = iso.categorical_node_match('fill', 'red')
>>> nx.is_isomorphic(G1, G2, node_match=nm)
True
```

For multidigraphs G1 and G2, using 'weight' edge attribute (default: 7)

```
>>> G1.add_edge(1,2, weight=7)
>>> G2.add_edge(10,20)
>>> em = iso.numerical_multiedge_match('weight', 7, rtol=1e-6)
>>> nx.is_isomorphic(G1, G2, edge_match=em)
True
```

For multigraphs G1 and G2, using 'weight' and 'linewidth' edge attributes with default values 7 and 2.5. Also using 'fill' node attribute with default value 'red'.

```
>>> em = iso.numerical_multiedge_match(['weight', 'linewidth'], [7, 2.5])
>>> nm = iso.categorical_node_match('fill', 'red')
>>> nx.is_isomorphic(G1, G2, edge_match=em, node_match=nm)
True
```

## 4.26.2 could be isomorphic

```
could_be_isomorphic(G1, G2)
```

Returns False if graphs are definitely not isomorphic. True does NOT guarantee isomorphism.

```
Parameters G1, G2: graphs
```

The two graphs G1 and G2 must be the same type.

#### **Notes**

Checks for matching degree, triangle, and number of cliques sequences.

## 4.26.3 fast\_could\_be\_isomorphic

```
fast\_could\_be\_isomorphic(G1, G2)
```

Returns False if graphs are definitely not isomorphic.

True does NOT guarantee isomorphism.

```
Parameters G1, G2: graphs
```

The two graphs G1 and G2 must be the same type.

#### **Notes**

Checks for matching degree and triangle sequences.

## 4.26.4 faster could be isomorphic

```
faster\_could\_be\_isomorphic(G1, G2)
```

Returns False if graphs are definitely not isomorphic.

True does NOT guarantee isomorphism.

```
Parameters G1, G2: graphs
```

The two graphs G1 and G2 must be the same type.

#### **Notes**

Checks for matching degree sequences.

## 4.26.5 Advanced Interface to VF2 Algorithm

## **VF2 Algorithm**

An implementation of VF2 algorithm for graph ismorphism testing.

The simplest interface to use this module is to call networkx.is\_isomorphic().

#### Introduction

The GraphMatcher and DiGraphMatcher are responsible for matching graphs or directed graphs in a predetermined manner. This usually means a check for an isomorphism, though other checks are also possible. For example, a subgraph of one graph can be checked for isomorphism to a second graph.

Matching is done via syntactic feasibility. It is also possible to check for semantic feasibility. Feasibility, then, is defined as the logical AND of the two functions.

To include a semantic check, the (Di)GraphMatcher class should be subclassed, and the semantic\_feasibility() function should be redefined. By default, the semantic feasibility function always returns True. The effect of this is that semantics are not considered in the matching of G1 and G2.

## **Examples**

Suppose G1 and G2 are isomorphic graphs. Verification is as follows:

```
>>> from networkx.algorithms import isomorphism
>>> G1 = nx.path_graph(4)
>>> G2 = nx.path_graph(4)
>>> GM = isomorphism.GraphMatcher(G1,G2)
>>> GM.is_isomorphic()
True
```

GM.mapping stores the isomorphism mapping from G1 to G2.

```
>>> GM.mapping {0: 0, 1: 1, 2: 2, 3: 3}
```

Suppose G1 and G2 are isomorphic directed graphs graphs. Verification is as follows:

4.26. Isomorphism 277

```
>>> G1 = nx.path_graph(4, create_using=nx.DiGraph())
>>> G2 = nx.path_graph(4, create_using=nx.DiGraph())
>>> DiGM = isomorphism.DiGraphMatcher(G1,G2)
>>> DiGM.is_isomorphic()
```

DiGM.mapping stores the isomorphism mapping from G1 to G2.

```
>>> DiGM.mapping {0: 0, 1: 1, 2: 2, 3: 3}
```

## Subgraph Isomorphism

Graph theory literature can be ambiguious about the meaning of the above statement, and we seek to clarify it now.

In the VF2 literature, a mapping M is said to be a graph-subgraph isomorphism iff M is an isomorphism between G2 and a subgraph of G1. Thus, to say that G1 and G2 are graph-subgraph isomorphic is to say that a subgraph of G1 is isomorphic to G2.

Other literature uses the phrase 'subgraph isomorphic' as in 'G1 does not have a subgraph isomorphic to G2'. Another use is as an in adverb for isomorphic. Thus, to say that G1 and G2 are subgraph isomorphic is to say that a subgraph of G1 is isomorphic to G2.

Finally, the term 'subgraph' can have multiple meanings. In this context, 'subgraph' always means a 'node-induced subgraph'. Edge-induced subgraph isomorphisms are not directly supported, but one should be able to perform the check by making use of nx.line\_graph(). For subgraphs which are not induced, the term 'monomorphism' is preferred over 'isomorphism'. Currently, it is not possible to check for monomorphisms.

Let G=(N,E) be a graph with a set of nodes N and set of edges E.

If G'=(N',E') is a subgraph, then: N' is a subset of N E' is a subset of E

- If G'=(N',E') is a node-induced subgraph, then: N' is a subset of N E' is the subset of edges in E relating nodes in N'
- If G'=(N',E') is an edge-induced subgrpah, then: N' is the subset of nodes in N related by edges in E' E' is a subset of E

## References

- [1] Luigi P. Cordella, Pasquale Foggia, Carlo Sansone, Mario Vento, "A (Sub)Graph Isomorphism Algorithm for Matching Large Graphs", IEEE Transactions on Pattern Analysis and Machine Intelligence, vol. 26, no. 10, pp. 1367-1372, Oct., 2004. http://ieeexplore.ieee.org/iel5/34/29305/01323804.pdf
- [2] L. P. Cordella, P. Foggia, C. Sansone, M. Vento, "An Improved Algorithm for Matching Large Graphs", 3rd IAPR-TC15 Workshop on Graph-based Representations in Pattern Recognition, Cuen, pp. 149-159, 2001. http://amalfi.dis.unina.it/graph/db/papers/vf-algorithm.pdf

#### See Also

syntactic\_feasibliity(), semantic\_feasibility()

#### **Notes**

Modified to handle undirected graphs. Modified to handle multiple edges.

In general, this problem is NP-Complete.

#### **Graph Matcher**

$GraphMatcher.\init\(G1, G2[, node\_match,])$	Initialize graph matcher.
GraphMatcher.initialize()	Reinitializes the state of the algorithm.
<pre>GraphMatcher.is_isomorphic()</pre>	Returns True if G1 and G2 are isomorphic graphs.
<pre>GraphMatcher.subgraph_is_isomorphic()</pre>	Returns True if a subgraph of G1 is isomorphic to G2.
<pre>GraphMatcher.isomorphisms_iter()</pre>	Generator over isomorphisms between G1 and G2.
GraphMatcher.subgraph_isomorphisms_iter()	Generator over isomorphisms between a subgraph of G1 and G2.
GraphMatcher.candidate_pairs_iter()	Iterator over candidate pairs of nodes in G1 and G2.
GraphMatcher.match()	Extends the isomorphism mapping.
GraphMatcher.semantic_feasibility(G1_node,)	Returns True if mapping G1_node to G2_node is semantically fea
GraphMatcher.syntactic_feasibility(G1_node,)	Returns True if adding (G1_node, G2_node) is syntactically feasib

## init

GraphMatcher.\_\_init\_\_(G1, G2, node\_match=None, edge\_match=None)
Initialize graph matcher.

## Parameters G1, G2: graph

The graphs to be tested.

### node match: callable

A function that returns True iff node n1 in G1 and n2 in G2 should be considered equal during the isomorphism test. The function will be called like:

```
node_match(G1.node[n1], G2.node[n2])
```

That is, the function will receive the node attribute dictionaries of the nodes under consideration. If None, then no attributes are considered when testing for an isomorphism.

### edge match: callable

A function that returns True iff the edge attribute dictionary for the pair of nodes (u1, v1) in G1 and (u2, v2) in G2 should be considered equal during the isomorphism test. The function will be called like:

```
edge_match(G1[u1][v1], G2[u2][v2])
```

That is, the function will receive the edge attribute dictionaries of the edges under consideration. If None, then no attributes are considered when testing for an isomorphism.

#### initialize

```
GraphMatcher.initialize()
```

Reinitializes the state of the algorithm.

This method should be redefined if using something other than GMState. If only subclassing GraphMatcher, a redefinition is not necessary.

4.26. Isomorphism 279

#### is isomorphic

```
GraphMatcher.is_isomorphic()
```

Returns True if G1 and G2 are isomorphic graphs.

#### subgraph\_is\_isomorphic

```
GraphMatcher.subgraph_is_isomorphic()
```

Returns True if a subgraph of G1 is isomorphic to G2.

## isomorphisms\_iter

```
GraphMatcher.isomorphisms_iter()
```

Generator over isomorphisms between G1 and G2.

## subgraph\_isomorphisms\_iter

```
GraphMatcher.subgraph_isomorphisms_iter()
```

Generator over isomorphisms between a subgraph of G1 and G2.

## candidate\_pairs\_iter

```
GraphMatcher.candidate_pairs_iter()
```

Iterator over candidate pairs of nodes in G1 and G2.

#### match

GraphMatcher.match()

Extends the isomorphism mapping.

This function is called recursively to determine if a complete isomorphism can be found between G1 and G2. It cleans up the class variables after each recursive call. If an isomorphism is found, we yield the mapping.

### semantic\_feasibility

```
GraphMatcher.semantic_feasibility(G1_node, G2_node)
```

Returns True if mapping G1\_node to G2\_node is semantically feasible.

#### syntactic feasibility

```
GraphMatcher.syntactic_feasibility(G1_node, G2_node)
```

Returns True if adding (G1\_node, G2\_node) is syntactically feasible.

This function returns True if it is adding the candidate pair to the current partial isomorphism mapping is allowable. The addition is allowable if the inclusion of the candidate pair does not make it impossible for an isomorphism to be found.

## **DiGraph Matcher**

DiGraphMatcher. $\_$ init $\_$ (G1,G2[,])	Initialize graph matcher.
DiGraphMatcher.initialize()	Reinitializes the state of the algorithm.
DiGraphMatcher.is_isomorphic()	Returns True if G1 and G2 are isomorphic graphs.
DiGraphMatcher.subgraph_is_isomorphic()	Returns True if a subgraph of G1 is isomorphic to G2.
DiGraphMatcher.isomorphisms_iter()	Generator over isomorphisms between G1 and G2.
DiGraphMatcher.subgraph_isomorphisms_iter()	Generator over isomorphisms between a subgraph of G1 and G2
	Continued on next

## Table 4.67 – continued from previous page

DiGraphMatcher.candidate_pairs_iter()	Iterator over candidate pairs of nodes in G1 and G2.
DiGraphMatcher.match()	Extends the isomorphism mapping.
$\label{limits} {\tt DiGraphMatcher.semantic\_feasibility} (G1\_node,)$	Returns True if mapping G1_node to G2_node is semantically fe
DiGraphMatcher.syntactic_feasibility()	Returns True if adding (G1_node, G2_node) is syntactically feas
	DiGraphMatcher.match() DiGraphMatcher.semantic_feasibility(G1_node,)

#### \_\_init\_\_

 $\label{linear_node_match} \mbox{DiGraphMatcher.} \mbox{$\_$initalize graph matcher.} \mbox{$(G1,G2,node\_match=None,edge\_match=None)$} \\ \mbox{Initialize graph matcher.} \mbox{$(G1,G2,node\_match=None,edge\_match=None)$} \\ \mbox{$=$$} \$ 

## Parameters G1, G2: graph

The graphs to be tested.

#### node match: callable

A function that returns True iff node n1 in G1 and n2 in G2 should be considered equal during the isomorphism test. The function will be called like:

```
node_match(G1.node[n1], G2.node[n2])
```

That is, the function will receive the node attribute dictionaries of the nodes under consideration. If None, then no attributes are considered when testing for an isomorphism.

## edge\_match: callable

A function that returns True iff the edge attribute dictionary for the pair of nodes (u1, v1) in G1 and (u2, v2) in G2 should be considered equal during the isomorphism test. The function will be called like:

```
edge_match(G1[u1][v1], G2[u2][v2])
```

That is, the function will receive the edge attribute dictionaries of the edges under consideration. If None, then no attributes are considered when testing for an isomorphism.

### initialize

DiGraphMatcher.initialize()

Reinitializes the state of the algorithm.

This method should be redefined if using something other than DiGMState. If only subclassing GraphMatcher, a redefinition is not necessary.

281

## is\_isomorphic

```
DiGraphMatcher.is_isomorphic()
```

Returns True if G1 and G2 are isomorphic graphs.

### subgraph\_is\_isomorphic

DiGraphMatcher.subgraph\_is\_isomorphic()

Returns True if a subgraph of G1 is isomorphic to G2.

## isomorphisms\_iter

```
DiGraphMatcher.isomorphisms_iter()
```

Generator over isomorphisms between G1 and G2.

4.26. Isomorphism

## subgraph\_isomorphisms\_iter

DiGraphMatcher.subgraph\_isomorphisms\_iter()

Generator over isomorphisms between a subgraph of G1 and G2.

### candidate\_pairs\_iter

DiGraphMatcher.candidate\_pairs\_iter()

Iterator over candidate pairs of nodes in G1 and G2.

#### match

DiGraphMatcher.match()

Extends the isomorphism mapping.

This function is called recursively to determine if a complete isomorphism can be found between G1 and G2. It cleans up the class variables after each recursive call. If an isomorphism is found, we yield the mapping.

### semantic\_feasibility

DiGraphMatcher.semantic\_feasibility(G1\_node, G2\_node)

Returns True if mapping G1\_node to G2\_node is semantically feasible.

#### syntactic feasibility

DiGraphMatcher.syntactic\_feasibility(G1\_node, G2\_node)

Returns True if adding (G1\_node, G2\_node) is syntactically feasible.

This function returns True if it is adding the candidate pair to the current partial isomorphism mapping is allowable. The addition is allowable if the inclusion of the candidate pair does not make it impossible for an isomorphism to be found.

## **Match helpers**

Returns a comparison function for a categorical node attribute.
Returns a comparison function for a categorical edge attribute.
Returns a comparison function for a categorical edge attribute.
Returns a comparison function for a numerical node attribute.
Returns a comparison function for a numerical edge attribute.
Returns a comparison function for a numerical edge attribute.
Returns a comparison function for a generic attribute.
Returns a comparison function for a generic attribute.
Returns a comparison function for a generic attribute.

## $categorical\_node\_match$

 ${\tt categorical\_node\_match}~(\textit{attr}, \textit{default})$ 

Returns a comparison function for a categorical node attribute.

The value(s) of the attr(s) must be hashable and comparable via the == operator since they are placed into a set([]) object. If the sets from G1 and G2 are the same, then the constructed function returns True.

Parameters attr: string | list

The categorical node attribute to compare, or a list of categorical node attributes to compare.

default : value | list

The default value for the categorical node attribute, or a list of default values for the categorical node attributes.

#### Returns match: function

The customized, categorical  $node_match$  function.

## **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.categorical_node_match('size', 1)
>>> nm = iso.categorical_node_match(['color', 'size'], ['red', 2])
```

## categorical\_edge\_match

```
categorical_edge_match (attr, default)
```

Returns a comparison function for a categorical edge attribute.

The value(s) of the attr(s) must be hashable and comparable via the == operator since they are placed into a set([]) object. If the sets from G1 and G2 are the same, then the constructed function returns True.

#### Parameters attr: string | list

The categorical edge attribute to compare, or a list of categorical edge attributes to compare.

## default: value | list

The default value for the categorical edge attribute, or a list of default values for the categorical edge attributes.

## Returns match: function

The customized, categorical  $edge_match$  function.

## **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.categorical_edge_match('size', 1)
>>> nm = iso.categorical_edge_match(['color', 'size'], ['red', 2])
```

## categorical\_multiedge\_match

```
categorical_multiedge_match (attr, default)
```

Returns a comparison function for a categorical edge attribute.

The value(s) of the attr(s) must be hashable and comparable via the == operator since they are placed into a set([]) object. If the sets from G1 and G2 are the same, then the constructed function returns True.

#### **Parameters attr**: string | list

The categorical edge attribute to compare, or a list of categorical edge attributes to compare.

#### default : value | list

The default value for the categorical edge attribute, or a list of default values for the categorical edge attributes.

#### Returns match: function

The customized, categorical  $edge_match$  function.

## **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.categorical_multiedge_match('size', 1)
>>> nm = iso.categorical_multiedge_match(['color', 'size'], ['red', 2])
```

## numerical\_node\_match

```
numerical_node_match (attr, default, rtol=1e-05, atol=1e-08)
```

Returns a comparison function for a numerical node attribute.

The value(s) of the attr(s) must be numerical and sortable. If the sorted list of values from G1 and G2 are the same within some tolerance, then the constructed function returns True.

## Parameters attr: string | list

The numerical node attribute to compare, or a list of numerical node attributes to compare.

default: value | list

The default value for the numerical node attribute, or a list of default values for the numerical node attributes.

rtol: float

The relative error tolerance.

atol: float

The absolute error tolerance.

## Returns match: function

The customized, numerical  $node_match$  function.

# **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.numerical_node_match('weight', 1.0)
>>> nm = iso.numerical_node_match(['weight', 'linewidth'], [.25, .5])
```

## numerical\_edge\_match

```
numerical_edge_match (attr, default, rtol=1e-05, atol=1e-08)
```

Returns a comparison function for a numerical edge attribute.

The value(s) of the attr(s) must be numerical and sortable. If the sorted list of values from G1 and G2 are the same within some tolerance, then the constructed function returns True.

**Parameters attr**: string | list

The numerical edge attribute to compare, or a list of numerical edge attributes to compare.

default : value | list

The default value for the numerical edge attribute, or a list of default values for the numerical edge attributes.

rtol: float

The relative error tolerance.

atol: float

The absolute error tolerance.

Returns match: function

The customized, numerical  $edge_match$  function.

#### **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.numerical_edge_match('weight', 1.0)
>>> nm = iso.numerical_edge_match(['weight', 'linewidth'], [.25, .5])
```

#### numerical\_multiedge\_match

```
numerical_multiedge_match (attr, default, rtol=1e-05, atol=1e-08)
```

Returns a comparison function for a numerical edge attribute.

The value(s) of the attr(s) must be numerical and sortable. If the sorted list of values from G1 and G2 are the same within some tolerance, then the constructed function returns True.

**Parameters attr**: string | list

The numerical edge attribute to compare, or a list of numerical edge attributes to compare.

default: value | list

The default value for the numerical edge attribute, or a list of default values for the numerical edge attributes.

rtol: float

The relative error tolerance.

atol: float

The absolute error tolerance.

Returns match: function

The customized, numerical  $edge_match$  function.

## **Examples**

```
>>> import networkx.algorithms.isomorphism as iso
>>> nm = iso.numerical_multiedge_match('weight', 1.0)
>>> nm = iso.numerical_multiedge_match(['weight', 'linewidth'], [.25, .5])
```

#### generic\_node\_match

```
generic_node_match (attr, default, op)
```

Returns a comparison function for a generic attribute.

The value(s) of the attr(s) are compared using the specified operators. If all the attributes are equal, then the constructed function returns True.

#### Parameters attr: string | list

The node attribute to compare, or a list of node attributes to compare.

default: value | list

The default value for the node attribute, or a list of default values for the node attributes.

op: callable | list

The operator to use when comparing attribute values, or a list of operators to use when comparing values for each attribute.

## Returns match: function

The customized, generic  $node_match$  function.

## **Examples**

```
>>> from operator import eq
>>> from networkx.algorithms.isomorphism.matchhelpers import close
>>> from networkx.algorithms.isomorphism import generic_node_match
>>> nm = generic_node_match('weight', 1.0, close)
>>> nm = generic_node_match('color', 'red', eq)
>>> nm = generic_node_match(['weight', 'color'], [1.0, 'red'], [close, eq])
```

## generic\_edge\_match

```
generic_edge_match (attr, default, op)
```

Returns a comparison function for a generic attribute.

The value(s) of the attr(s) are compared using the specified operators. If all the attributes are equal, then the constructed function returns True.

# Parameters attr: string | list

The edge attribute to compare, or a list of edge attributes to compare.

default: value | list

The default value for the edge attribute, or a list of default values for the edge attributes.

op: callable | list

The operator to use when comparing attribute values, or a list of operators to use when comparing values for each attribute.

# Returns match: function

The customized, generic  $edge_match$  function.

## **Examples**

```
>>> from operator import eq
>>> from networkx.algorithms.isomorphism.matchhelpers import close
>>> from networkx.algorithms.isomorphism import generic_edge_match
>>> nm = generic_edge_match('weight', 1.0, close)
>>> nm = generic_edge_match('color', 'red', eq)
>>> nm = generic_edge_match(['weight', 'color'], [1.0, 'red'], [close, eq])
```

# $generic\_multiedge\_match$

```
generic_multiedge_match (attr, default, op)
```

Returns a comparison function for a generic attribute.

The value(s) of the attr(s) are compared using the specified operators. If all the attributes are equal, then the constructed function returns True. Potentially, the constructed edge\_match function can be slow since it must verify that no isomorphism exists between the multiedges before it returns False.

#### Parameters attr: string | list

The edge attribute to compare, or a list of node attributes to compare.

#### default: value | list

The default value for the edge attribute, or a list of default values for the dgeattributes.

```
op: callable | list
```

The operator to use when comparing attribute values, or a list of operators to use when comparing values for each attribute.

## Returns match: function

The customized, generic  $edge_match$  function.

# **Examples**

# 4.27 Link Analysis

# 4.27.1 PageRank

PageRank analysis of graph structure.

pagerank(G[, alpha, personalization,])	Return the PageRank of the nodes in the graph.
pagerank_numpy(G[, alpha, personalization,])	Return the PageRank of the nodes in the graph.
	Continued on next page

4.27. Link Analysis 287

# Table 4.69 – continued from previous page

<pre>pagerank_scipy(G[, alpha, personalization,])</pre>	Return the PageRank of the nodes in the graph.
<pre>google_matrix(G[, alpha, personalization,])</pre>	Return the Google matrix of the graph.

# pagerank

Return the PageRank of the nodes in the graph.

PageRank computes a ranking of the nodes in the graph G based on the structure of the incoming links. It was originally designed as an algorithm to rank web pages.

# Parameters G: graph

A NetworkX graph. Undirected graphs will be converted to a directed graph with two directed edges for each undirected edge.

alpha: float, optional

Damping parameter for PageRank, default=0.85.

## personalization: dict, optional

The "personalization vector" consisting of a dictionary with a key for every graph node and nonzero personalization value for each node. By default, a uniform distribution is used.

max\_iter: integer, optional

Maximum number of iterations in power method eigenvalue solver.

tol: float, optional

Error tolerance used to check convergence in power method solver.

nstart: dictionary, optional

Starting value of PageRank iteration for each node.

weight: key, optional

Edge data key to use as weight. If None weights are set to 1.

# dangling: dict, optional

The outedges to be assigned to any "dangling" nodes, i.e., nodes without any outedges. The dict key is the node the outedge points to and the dict value is the weight of that outedge. By default, dangling nodes are given outedges according to the personalization vector (uniform if not specified). This must be selected to result in an irreducible transition matrix (see notes under google\_matrix). It may be common to have the dangling dict to be the same as the personalization dict.

## Returns pagerank: dictionary

Dictionary of nodes with PageRank as value

#### See also:

pagerank\_numpy, pagerank\_scipy, google\_matrix

#### **Notes**

The eigenvector calculation is done by the power iteration method and has no guarantee of convergence. The iteration will stop after max\_iter iterations or an error tolerance of number\_of\_nodes(G)\*tol has been reached.

The PageRank algorithm was designed for directed graphs but this algorithm does not check if the input graph is directed and will execute on undirected graphs by converting each edge in the directed graph to two edges.

#### References

[R299], [R300]

#### **Examples**

```
>>> G = nx.DiGraph(nx.path_graph(4))
>>> pr = nx.pagerank(G, alpha=0.9)
```

## pagerank\_numpy

**pagerank\_numpy** (*G*, *alpha=0.85*, *personalization=None*, *weight='weight'*, *dangling=None*) Return the PageRank of the nodes in the graph.

PageRank computes a ranking of the nodes in the graph G based on the structure of the incoming links. It was originally designed as an algorithm to rank web pages.

## **Parameters G**: graph

A NetworkX graph. Undirected graphs will be converted to a directed graph with two directed edges for each undirected edge.

## alpha: float, optional

Damping parameter for PageRank, default=0.85.

## personalization: dict, optional

The "personalization vector" consisting of a dictionary with a key for every graph node and nonzero personalization value for each node. By default, a uniform distribution is used.

#### weight: key, optional

Edge data key to use as weight. If None weights are set to 1.

# dangling: dict, optional

The outedges to be assigned to any "dangling" nodes, i.e., nodes without any outedges. The dict key is the node the outedge points to and the dict value is the weight of that outedge. By default, dangling nodes are given outedges according to the personalization vector (uniform if not specified) This must be selected to result in an irreducible transition matrix (see notes under google\_matrix). It may be common to have the dangling dict to be the same as the personalization dict.

## Returns pagerank: dictionary

Dictionary of nodes with PageRank as value.

4.27. Link Analysis 289

#### See also:

```
pagerank, pagerank_scipy, google_matrix
```

#### **Notes**

The eigenvector calculation uses NumPy's interface to the LAPACK eigenvalue solvers. This will be the fastest and most accurate for small graphs.

This implementation works with Multi(Di)Graphs. For multigraphs the weight between two nodes is set to be the sum of all edge weights between those nodes.

#### References

[R301], [R302]

#### **Examples**

```
>>> G = nx.DiGraph(nx.path_graph(4))
>>> pr = nx.pagerank_numpy(G, alpha=0.9)
```

# pagerank\_scipy

 $pagerank\_scipy$  (G, alpha=0.85, personalization=None,  $max\_iter=100$ , tol=1e-06, weight='weight', dangling=None)

Return the PageRank of the nodes in the graph.

PageRank computes a ranking of the nodes in the graph G based on the structure of the incoming links. It was originally designed as an algorithm to rank web pages.

# Parameters G: graph

A NetworkX graph. Undirected graphs will be converted to a directed graph with two directed edges for each undirected edge.

alpha: float, optional

Damping parameter for PageRank, default=0.85.

# personalization: dict, optional

The "personalization vector" consisting of a dictionary with a key for every graph node and nonzero personalization value for each node. By default, a uniform distribution is used.

max\_iter: integer, optional

Maximum number of iterations in power method eigenvalue solver.

tol: float, optional

Error tolerance used to check convergence in power method solver.

weight: key, optional

Edge data key to use as weight. If None weights are set to 1.

dangling: dict, optional

The outedges to be assigned to any "dangling" nodes, i.e., nodes without any outedges. The dict key is the node the outedge points to and the dict value is the weight of that outedge. By default, dangling nodes are given outedges according to the personalization vector (uniform if not specified) This must be selected to result in an irreducible transition matrix (see notes under google\_matrix). It may be common to have the dangling dict to be the same as the personalization dict.

# Returns pagerank: dictionary

Dictionary of nodes with PageRank as value

#### See also:

```
pagerank_numpy, google_matrix
```

#### **Notes**

The eigenvector calculation uses power iteration with a SciPy sparse matrix representation.

This implementation works with Multi(Di)Graphs. For multigraphs the weight between two nodes is set to be the sum of all edge weights between those nodes.

#### References

```
[R303], [R304]
```

### **Examples**

```
>>> G = nx.DiGraph(nx.path_graph(4))
>>> pr = nx.pagerank_scipy(G, alpha=0.9)
```

#### google matrix

```
google_matrix(G, alpha=0.85, personalization=None, nodelist=None, weight='weight', dan-gling=None)
Return the Google matrix of the graph.
```

# Parameters G: graph

A NetworkX graph. Undirected graphs will be converted to a directed graph with two directed edges for each undirected edge.

alpha: float

The damping factor.

#### personalization: dict, optional

The "personalization vector" consisting of a dictionary with a key for every graph node and nonzero personalization value for each node. By default, a uniform distribution is used.

nodelist: list, optional

The rows and columns are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

weight: key, optional

Edge data key to use as weight. If None weights are set to 1.

# dangling: dict, optional

The outedges to be assigned to any "dangling" nodes, i.e., nodes without any outedges. The dict key is the node the outedge points to and the dict value is the weight of that outedge. By default, dangling nodes are given outedges according to the personalization vector (uniform if not specified) This must be selected to result in an irreducible transition matrix (see notes below). It may be common to have the dangling dict to be the same as the personalization dict.

## **Returns** A: NumPy matrix

Google matrix of the graph

#### See also:

pagerank, pagerank\_numpy, pagerank\_scipy

#### **Notes**

The matrix returned represents the transition matrix that describes the Markov chain used in PageRank. For PageRank to converge to a unique solution (i.e., a unique stationary distribution in a Markov chain), the transition matrix must be irreducible. In other words, it must be that there exists a path between every pair of nodes in the graph, or else there is the potential of "rank sinks."

This implementation works with Multi(Di)Graphs. For multigraphs the weight between two nodes is set to be the sum of all edge weights between those nodes.

## 4.27.2 Hits

Hubs and authorities analysis of graph structure.

hits(G[, max_iter, tol, nstart, normalized])	Return HITS hubs and authorities values for nodes.
hits_numpy(G[, normalized])	Return HITS hubs and authorities values for nodes.
hits_scipy(G[, max_iter, tol, normalized])	Return HITS hubs and authorities values for nodes.
hub_matrix(G[, nodelist])	Return the HITS hub matrix.
$authority_matrix(G[, nodelist])$	Return the HITS authority matrix.

# hits

hits (G, max\_iter=100, tol=1e-08, nstart=None, normalized=True)

Return HITS hubs and authorities values for nodes.

The HITS algorithm computes two numbers for a node. Authorities estimates the node value based on the incoming links. Hubs estimates the node value based on outgoing links.

Parameters G: graph

A NetworkX graph

max\_iter: interger, optional

Maximum number of iterations in power method.

tol: float, optional

Error tolerance used to check convergence in power method iteration.

nstart: dictionary, optional

Starting value of each node for power method iteration.

normalized : bool (default=True)

Normalize results by the sum of all of the values.

**Returns** (hubs, authorities): two-tuple of dictionaries

Two dictionaries keyed by node containing the hub and authority values.

#### **Notes**

The eigenvector calculation is done by the power iteration method and has no guarantee of convergence. The iteration will stop after max\_iter iterations or an error tolerance of number\_of\_nodes(G)\*tol has been reached.

The HITS algorithm was designed for directed graphs but this algorithm does not check if the input graph is directed and will execute on undirected graphs.

#### References

[R293], [R294]

## **Examples**

```
>>> G=nx.path_graph(4)
>>> h,a=nx.hits(G)
```

## hits numpy

## hits\_numpy (G, normalized=True)

Return HITS hubs and authorities values for nodes.

The HITS algorithm computes two numbers for a node. Authorities estimates the node value based on the incoming links. Hubs estimates the node value based on outgoing links.

Parameters G: graph

A NetworkX graph

normalized : bool (default=True)

Normalize results by the sum of all of the values.

Returns (hubs, authorities): two-tuple of dictionaries

Two dictionaries keyed by node containing the hub and authority values.

#### **Notes**

The eigenvector calculation uses NumPy's interface to LAPACK.

The HITS algorithm was designed for directed graphs but this algorithm does not check if the input graph is directed and will execute on undirected graphs.

#### References

[R295], [R296]

# **Examples**

```
>>> G=nx.path_graph(4)
>>> h,a=nx.hits(G)
```

# hits\_scipy

hits\_scipy (G, max\_iter=100, tol=1e-06, normalized=True)

Return HITS hubs and authorities values for nodes.

The HITS algorithm computes two numbers for a node. Authorities estimates the node value based on the incoming links. Hubs estimates the node value based on outgoing links.

Parameters G: graph

A NetworkX graph

max\_iter: interger, optional

Maximum number of iterations in power method.

tol: float, optional

Error tolerance used to check convergence in power method iteration.

nstart: dictionary, optional

Starting value of each node for power method iteration.

normalized : bool (default=True)

Normalize results by the sum of all of the values.

Returns (hubs,authorities): two-tuple of dictionaries

Two dictionaries keyed by node containing the hub and authority values.

## **Notes**

This implementation uses SciPy sparse matrices.

The eigenvector calculation is done by the power iteration method and has no guarantee of convergence. The iteration will stop after max\_iter iterations or an error tolerance of number\_of\_nodes(G)\*tol has been reached.

The HITS algorithm was designed for directed graphs but this algorithm does not check if the input graph is directed and will execute on undirected graphs.

#### References

[R297], [R298]

## **Examples**

>>> G=nx.path\_graph(4)
>>> h,a=nx.hits(G)

# hub\_matrix

**hub\_matrix** (*G*, *nodelist=None*)

Return the HITS hub matrix.

# authority\_matrix

**authority\_matrix** (*G*, nodelist=None)
Return the HITS authority matrix.

# 4.28 Link Prediction

Link prediction algorithms.

$resource\_allocation\_index(G[,ebunch])$	Compute the resource allocation index of all node pairs in ebunch.
<pre>jaccard_coefficient(G[, ebunch])</pre>	Compute the Jaccard coefficient of all node pairs in ebunch.
$adamic\_adar\_index(G[,ebunch])$	Compute the Adamic-Adar index of all node pairs in ebunch.
$preferential\_attachment(G[,ebunch])$	Compute the preferential attachment score of all node pairs in ebunch.
$cn_soundarajan_hopcroft(G[, ebunch, community])$	Count the number of common neighbors of all node pairs in ebunch us
$ra_index_soundarajan_hopcroft(G[,ebunch,])$	Compute the resource allocation index of all node pairs in ebunch usin
within_inter_cluster( $G[$ , ebunch, delta,])	Compute the ratio of within- and inter-cluster common neighbors of al

# 4.28.1 resource\_allocation\_index

resource allocation index(G, ebunch=None)

Compute the resource allocation index of all node pairs in ebunch.

Resource allocation index of u and v is defined as

$$\sum_{w \in \Gamma(u) \cap \Gamma(v)} \frac{1}{|\Gamma(w)|}$$

where  $\Gamma(u)$  denotes the set of neighbors of u.

Parameters G: graph

A NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

Resource allocation index will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

Returns piter: iterator

4.28. Link Prediction 295

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their resource allocation index.

#### References

[R310]

## **Examples**

```
>>> import networkx as nx
>>> G = nx.complete_graph(5)
>>> preds = nx.resource_allocation_index(G, [(0, 1), (2, 3)])
>>> for u, v, p in preds:
... '(%d, %d) -> %.8f' % (u, v, p)
...
'(0, 1) -> 0.75000000'
'(2, 3) -> 0.75000000'
```

# 4.28.2 jaccard\_coefficient

```
jaccard_coefficient(G, ebunch=None)
```

Compute the Jaccard coefficient of all node pairs in ebunch.

Jaccard coefficient of nodes u and v is defined as

```
\frac{|\Gamma(u) \cap \Gamma(v)|}{|\Gamma(u) \cup \Gamma(v)|}
```

where  $\Gamma(u)$  denotes the set of neighbors of u.

Parameters G: graph

A NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

Jaccard coefficient will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

Returns piter: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their Jaccard coefficient.

#### References

[R307]

# **Examples**

```
>>> import networkx as nx
>>> G = nx.complete_graph(5)
>>> preds = nx.jaccard_coefficient(G, [(0, 1), (2, 3)])
>>> for u, v, p in preds:
... '(%d, %d) -> %.8f' % (u, v, p)
...
'(0, 1) -> 0.60000000'
'(2, 3) -> 0.60000000'
```

# 4.28.3 adamic\_adar\_index

#### adamic\_adar\_index(G, ebunch=None)

Compute the Adamic-Adar index of all node pairs in ebunch.

Adamic-Adar index of u and v is defined as

$$\sum_{w \in \Gamma(u) \cap \Gamma(v)} \frac{1}{\log |\Gamma(w)|}$$

where  $\Gamma(u)$  denotes the set of neighbors of u.

# Parameters G: graph

NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

Adamic-Adar index will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

## **Returns piter**: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their Adamic-Adar index.

#### References

[R305]

## **Examples**

```
>>> import networkx as nx
>>> G = nx.complete_graph(5)
>>> preds = nx.adamic_adar_index(G, [(0, 1), (2, 3)])
>>> for u, v, p in preds:
... '(%d, %d) -> %.8f' % (u, v, p)
...
'(0, 1) -> 2.16404256'
'(2, 3) -> 2.16404256'
```

# 4.28.4 preferential\_attachment

## preferential\_attachment(G, ebunch=None)

Compute the preferential attachment score of all node pairs in ebunch.

4.28. Link Prediction 297

Preferential attachment score of u and v is defined as

$$|\Gamma(u)||\Gamma(v)|$$

where  $\Gamma(u)$  denotes the set of neighbors of u.

Parameters G: graph

NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

Preferential attachment score will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

Returns piter: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their preferential attachment score.

#### References

[R308]

# **Examples**

```
>>> import networkx as nx
>>> G = nx.complete_graph(5)
>>> preds = nx.preferential_attachment(G, [(0, 1), (2, 3)])
>>> for u, v, p in preds:
... '(%d, %d) -> %d' % (u, v, p)
...
'(0, 1) -> 16'
'(2, 3) -> 16'
```

# 4.28.5 cn\_soundarajan\_hopcroft

cn soundarajan hopcroft(G, ebunch=None, community='community')

Count the number of common neighbors of all node pairs in ebunch using community information.

For two nodes u and v, this function computes the number of common neighbors and bonus one for each common neighbor belonging to the same community as u and v. Mathematically,

$$|\Gamma(u)\cap\Gamma(v)|+\sum_{w\in\Gamma(u)\cap\Gamma(v)}f(w)$$

where f(w) equals 1 if w belongs to the same community as u and v or 0 otherwise and  $\Gamma(u)$  denotes the set of neighbors of u.

Parameters G: graph

A NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

The score will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

```
community : string, optional (default = 'community')
```

Nodes attribute name containing the community information. G[u][community] identifies which community u belongs to. Each node belongs to at most one community. Default value: 'community'.

# Returns piter: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their score.

#### References

[R306]

## **Examples**

# 4.28.6 ra index soundarajan hopcroft

ra\_index\_soundarajan\_hopcroft (G, ebunch=None, community='community')

Compute the resource allocation index of all node pairs in ebunch using community information.

For two nodes u and v, this function computes the resource allocation index considering only common neighbors belonging to the same community as u and v. Mathematically,

$$\sum_{w \in \Gamma(u) \cap \Gamma(v)} \frac{f(w)}{|\Gamma(w)|}$$

where f(w) equals 1 if w belongs to the same community as u and v or 0 otherwise and  $\Gamma(u)$  denotes the set of neighbors of u.

Parameters G: graph

A NetworkX undirected graph.

**ebunch**: iterable of node pairs, optional (default = None)

The score will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

community : string, optional (default = 'community')

4.28. Link Prediction 299

Nodes attribute name containing the community information. G[u][community] identifies which community u belongs to. Each node belongs to at most one community. Default value: 'community'.

## Returns piter: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their score.

#### References

[R309]

#### **Examples**

# 4.28.7 within\_inter\_cluster

```
within_inter_cluster (G, ebunch=None, delta=0.001, community='community')

Compute the ratio of within- and inter-cluster common neighbors of all node pairs in ebunch.
```

For two nodes u and v, if a common neighbor w belongs to the same community as them, w is considered as within-cluster common neighbor of u and v. Otherwise, it is considered as inter-cluster common neighbor of u and v. The ratio between the size of the set of within- and inter-cluster common neighbors is defined as the WIC measure. [R311]

#### Parameters G: graph

A NetworkX undirected graph.

```
ebunch: iterable of node pairs, optional (default = None)
```

The WIC measure will be computed for each pair of nodes given in the iterable. The pairs must be given as 2-tuples (u, v) where u and v are nodes in the graph. If ebunch is None then all non-existent edges in the graph will be used. Default value: None.

```
delta: float, optional (default = 0.001)
```

Value to prevent division by zero in case there is no inter-cluster common neighbor between two nodes. See [R311] for details. Default value: 0.001.

```
community : string, optional (default = 'community')
```

Nodes attribute name containing the community information. G[u][community] identifies which community u belongs to. Each node belongs to at most one community. Default value: 'community'.

# Returns piter: iterator

An iterator of 3-tuples in the form (u, v, p) where (u, v) is a pair of nodes and p is their WIC measure.

#### References

[R311]

#### **Examples**

```
>>> import networkx as nx
>>> G = nx.Graph()
>>> G.add_edges_from([(0, 1), (0, 2), (0, 3), (1, 4), (2, 4), (3, 4)])
>>> G.node[0]['community'] = 0
>>> G.node[1]['community'] = 1
>>> G.node[2]['community'] = 0
>>> G.node[3]['community'] = 0
>>> G.node[4]['community'] = 0
>>> preds = nx.within_inter_cluster(G, [(0, 4)])
>>> for u, v, p in preds:
        '(%d, %d) -> %.8f' % (u, v, p)
'(0, 4) -> 1.99800200'
>>> preds = nx.within_inter_cluster(G, [(0, 4)], delta=0.5)
>>> for u, v, p in preds:
       '(%d, %d) -> %.8f' % (u, v, p)
'(0, 4) -> 1.333333333'
```

# 4.29 Matching

$ exttt{maximal\_matching}(G)$	Find a maximal cardinality matching in the graph.
<pre>max_weight_matching(G[, maxcardinality])</pre>	Compute a maximum-weighted matching of G.

# 4.29.1 maximal\_matching

#### maximal matching(G)

Find a maximal cardinality matching in the graph.

A matching is a subset of edges in which no node occurs more than once. The cardinality of a matching is the number of matched edges.

Parameters G: NetworkX graph

Undirected graph

**Returns matching**: set

4.29. Matching 301

A maximal matching of the graph.

#### **Notes**

The algorithm greedily selects a maximal matching M of the graph G (i.e. no superset of M exists). It runs in O(|E|) time.

# 4.29.2 max weight matching

 ${\tt max\_weight\_matching}$  (G, maxcardinality = False)

Compute a maximum-weighted matching of G.

A matching is a subset of edges in which no node occurs more than once. The cardinality of a matching is the number of matched edges. The weight of a matching is the sum of the weights of its edges.

**Parameters G**: NetworkX graph

Undirected graph

maxcardinality: bool, optional

If maxcardinality is True, compute the maximum-cardinality matching with maximum weight among all maximum-cardinality matchings.

Returns mate: dictionary

The matching is returned as a dictionary, mate, such that mate[v] == w if node v is matched to node w. Unmatched nodes do not occur as a key in mate.

#### **Notes**

If G has edges with 'weight' attribute the edge data are used as weight values else the weights are assumed to be 1.

This function takes time O(number\_of\_nodes \*\* 3).

If all edge weights are integers, the algorithm uses only integer computations. If floating point weights are used, the algorithm could return a slightly suboptimal matching due to numeric precision errors.

This method is based on the "blossom" method for finding augmenting paths and the "primal-dual" method for finding a matching of maximum weight, both methods invented by Jack Edmonds [R312].

#### References

[R312]

# 4.30 Maximal independent set

Algorithm to find a maximal (not maximum) independent set.

maximal\_independent\_set(G[, nodes]) Return a random maximal independent set guaranteed to contain a given set of node

# 4.30.1 maximal independent set

# maximal\_independent\_set (G, nodes=None)

Return a random maximal independent set guaranteed to contain a given set of nodes.

An independent set is a set of nodes such that the subgraph of G induced by these nodes contains no edges. A maximal independent set is an independent set such that it is not possible to add a new node and still get an independent set.

Parameters G: NetworkX graph

nodes: list or iterable

Nodes that must be part of the independent set. This set of nodes must be independent.

Returns indep nodes: list

List of nodes that are part of a maximal independent set.

## Raises NetworkXUnfeasible

If the nodes in the provided list are not part of the graph or do not form an independent set, an exception is raised.

#### **Notes**

This algorithm does not solve the maximum independent set problem.

## **Examples**

```
>>> G = nx.path_graph(5)
>>> nx.maximal_independent_set(G)
[4, 0, 2]
>>> nx.maximal_independent_set(G, [1])
[1, 3]
```

# 4.31 Minimum Spanning Tree

Computes minimum spanning tree of a weighted graph.

$minimum\_spanning\_tree(G[, weight])$	Return a minimum spanning tree or forest of an undirected weighted graph.
<pre>minimum_spanning_edges(G[, weight, data])</pre>	Generate edges in a minimum spanning forest of an undirected weighted graph

# 4.31.1 minimum spanning tree

```
minimum_spanning_tree(G, weight='weight')
```

Return a minimum spanning tree or forest of an undirected weighted graph.

A minimum spanning tree is a subgraph of the graph (a tree) with the minimum sum of edge weights.

If the graph is not connected a spanning forest is constructed. A spanning forest is a union of the spanning trees for each connected component of the graph.

Parameters G: NetworkX Graph

```
weight: string
```

Edge data key to use for weight (default 'weight').

Returns G: NetworkX Graph

A minimum spanning tree or forest.

#### **Notes**

Uses Kruskal's algorithm.

If the graph edges do not have a weight attribute a default weight of 1 will be used.

#### **Examples**

```
>>> G=nx.cycle_graph(4)
>>> G.add_edge(0,3,weight=2) # assign weight 2 to edge 0-3
>>> T=nx.minimum_spanning_tree(G)
>>> print(sorted(T.edges(data=True)))
[(0, 1, {}), (1, 2, {}), (2, 3, {})]
```

# 4.31.2 minimum spanning edges

```
minimum_spanning_edges (G, weight='weight', data=True)
```

Generate edges in a minimum spanning forest of an undirected weighted graph.

A minimum spanning tree is a subgraph of the graph (a tree) with the minimum sum of edge weights. A spanning forest is a union of the spanning trees for each connected component of the graph.

# Parameters G: NetworkX Graph

```
weight: string
```

Edge data key to use for weight (default 'weight').

data: bool, optional

If True yield the edge data along with the edge.

Returns edges: iterator

A generator that produces edges in the minimum spanning tree. The edges are three-tuples (u,v,w) where w is the weight.

## **Notes**

Uses Kruskal's algorithm.

If the graph edges do not have a weight attribute a default weight of 1 will be used.

Modified code from David Eppstein, April 2006 http://www.ics.uci.edu/~eppstein/PADS/

## **Examples**

```
>>> G=nx.cycle_graph(4)
>>> G.add_edge(0,3,weight=2) # assign weight 2 to edge 0-3
>>> mst=nx.minimum_spanning_edges(G,data=False) # a generator of MST edges
>>> edgelist=list(mst) # make a list of the edges
>>> print(sorted(edgelist))
[(0, 1), (1, 2), (2, 3)]
```

# 4.32 Operators

Unary operations on graphs

complement(G[, name])	Return the graph complement of G.
reverse(G[, copy])	Return the reverse directed graph of G.

# 4.32.1 complement

```
complement (G, name=None)
```

Return the graph complement of G.

Parameters G: graph

A NetworkX graph

name: string

Specify name for new graph

**Returns** GC : A new graph.

## Notes

Note that complement() does not create self-loops and also does not produce parallel edges for MultiGraphs.

Graph, node, and edge data are not propagated to the new graph.

## **4.32.2** reverse

```
reverse(G, copy=True)
```

Return the reverse directed graph of G.

Parameters G: directed graph

A NetworkX directed graph

copy: bool

If True, then a new graph is returned. If False, then the graph is reversed in place.

Returns H: directed graph

The reversed G.

Operations on graphs including union, intersection, difference.

4.32. Operators 305

compose(G, H[, name])	Return a new graph of G composed with H.	
union(G, H[, rename, name])	Return the union of graphs G and H.	
disjoint_union(G, H)	Return the disjoint union of graphs G and H.	
intersection(G, H)	Return a new graph that contains only the edges that exist in both G and H.	
difference(G, H)	Return a new graph that contains the edges that exist in G but not in H.	
$symmetric\_difference(G, H)$	Return new graph with edges that exist in either G or H but not both.	

# **4.32.3** compose

compose(G, H, name=None)

Return a new graph of G composed with H.

Composition is the simple union of the node sets and edge sets. The node sets of G and H do not need to be disjoint.

Parameters G,H: graph

A NetworkX graph

name: string

Specify name for new graph

**Returns** C: A new graph with the same type as G

## **Notes**

It is recommended that G and H be either both directed or both undirected. Attributes from H take precedent over attributes from G.

# 4.32.4 union

union (G, H, rename = (None, None), name = None)

Return the union of graphs G and H.

Graphs G and H must be disjoint, otherwise an exception is raised.

**Parameters G,H**: graph

A NetworkX graph

create\_using : NetworkX graph

Use specified graph for result. Otherwise

rename: bool, default=(None, None)

Node names of G and H can be changed by specifying the tuple rename=('G-','H-') (for example). Node "u" in G is then renamed "G-u" and "v" in H is renamed "H-v".

name: string

Specify the name for the union graph

**Returns** U: A union graph with the same type as G.

## See also:

disjoint\_union

#### **Notes**

To force a disjoint union with node relabeling, use disjoint\_union(G,H) or convert\_node\_labels\_to integers().

Graph, edge, and node attributes are propagated from G and H to the union graph. If a graph attribute is present in both G and H the value from H is used.

# 4.32.5 disjoint\_union

#### disjoint union (G, H)

Return the disjoint union of graphs G and H.

This algorithm forces distinct integer node labels.

**Parameters G,H**: graph
A NetworkX graph

**Returns** U: A union graph with the same type as G.

#### **Notes**

A new graph is created, of the same class as G. It is recommended that G and H be either both directed or both undirected.

The nodes of G are relabeled 0 to len(G)-1, and the nodes of H are relabeled len(G) to len(G)+len(H)-1.

Graph, edge, and node attributes are propagated from G and H to the union graph. If a graph attribute is present in both G and H the value from H is used.

# 4.32.6 intersection

#### intersection(G, H)

Return a new graph that contains only the edges that exist in both G and H.

The node sets of H and G must be the same.

Parameters G,H: graph

A NetworkX graph. G and H must have the same node sets.

**Returns GH**: A new graph with the same type as G.

#### **Notes**

Attributes from the graph, nodes, and edges are not copied to the new graph. If you want a new graph of the intersection of G and H with the attributes (including edge data) from G use remove\_nodes\_from() as follows

```
>>> G=nx.path_graph(3)
>>> H=nx.path_graph(5)
>>> R=G.copy()
>>> R.remove_nodes_from(n for n in G if n not in H)
```

4.32. Operators 307

# 4.32.7 difference

#### difference(G, H)

Return a new graph that contains the edges that exist in G but not in H.

The node sets of H and G must be the same.

Parameters G,H: graph

A NetworkX graph. G and H must have the same node sets.

**Returns D** : A new graph with the same type as G.

#### **Notes**

Attributes from the graph, nodes, and edges are not copied to the new graph. If you want a new graph of the difference of G and H with with the attributes (including edge data) from G use remove\_nodes\_from() as follows:

```
>>> G = nx.path_graph(3)
>>> H = nx.path_graph(5)
>>> R = G.copy()
>>> R.remove_nodes_from(n for n in G if n in H)
```

# 4.32.8 symmetric difference

## $symmetric\_difference(G, H)$

Return new graph with edges that exist in either G or H but not both.

The node sets of H and G must be the same.

Parameters G,H: graph

A NetworkX graph. G and H must have the same node sets.

**Returns D**: A new graph with the same type as G.

## Notes

Attributes from the graph, nodes, and edges are not copied to the new graph.

Operations on many graphs.

compose_all(graphs[, name])	Return the composition of all graphs.	
union_all(graphs[, rename, name])	Return the union of all graphs.	
disjoint_union_all(graphs)	Return the disjoint union of all graphs.	
intersection_all(graphs)	Return a new graph that contains only the edges that exist in all graphs.	

# 4.32.9 compose all

## compose\_all (graphs, name=None)

Return the composition of all graphs.

Composition is the simple union of the node sets and edge sets. The node sets of the supplied graphs need not be disjoint.

Parameters graphs: list

List of NetworkX graphs

name: string

Specify name for new graph

**Returns** C: A graph with the same type as the first graph in list

#### **Notes**

It is recommended that the supplied graphs be either all directed or all undirected.

Graph, edge, and node attributes are propagated to the union graph. If a graph attribute is present in multiple graphs, then the value from the last graph in the list with that attribute is used.

# 4.32.10 union all

union\_all (graphs, rename=(None, ), name=None)

Return the union of all graphs.

The graphs must be disjoint, otherwise an exception is raised.

Parameters graphs: list of graphs

List of NetworkX graphs

rename : bool , default=(None, None)

Node names of G and H can be changed by specifying the tuple rename=('G-','H-') (for example). Node "u" in G is then renamed "G-u" and "v" in H is renamed "H-v".

name: string

Specify the name for the union graph@not implemnted for('direct

**Returns** U: a graph with the same type as the first graph in list

#### See also:

```
union, disjoint_union_all
```

#### Notes

To force a disjoint union with node relabeling, use disjoint\_union\_all(G,H) or convert\_node\_labels\_to integers().

Graph, edge, and node attributes are propagated to the union graph. If a graph attribute is present in multiple graphs, then the value from the last graph in the list with that attribute is used.

# 4.32.11 disjoint union all

## disjoint\_union\_all (graphs)

Return the disjoint union of all graphs.

This operation forces distinct integer node labels starting with 0 for the first graph in the list and numbering consecutively.

Parameters graphs: list

4.32. Operators 309

List of NetworkX graphs

**Returns** U: A graph with the same type as the first graph in list

#### **Notes**

It is recommended that the graphs be either all directed or all undirected.

Graph, edge, and node attributes are propagated to the union graph. If a graph attribute is present in multiple graphs, then the value from the last graph in the list with that attribute is used.

# 4.32.12 intersection\_all

# intersection\_all (graphs)

Return a new graph that contains only the edges that exist in all graphs.

All supplied graphs must have the same node set.

# Parameters graphs\_list: list

List of NetworkX graphs

**Returns R**: A new graph with the same type as the first graph in list

#### **Notes**

Attributes from the graph, nodes, and edges are not copied to the new graph.

## Graph products.

cartesian_product(G, H)	Return the Cartesian product of G and H.
$lexicographic\_product(G, H)$	Return the lexicographic product of G and H.
$strong\_product(G, H)$	Return the strong product of G and H.
tensor_product(G, H)	Return the tensor product of G and H.

# 4.32.13 cartesian product

## $cartesian\_product(G, H)$

Return the Cartesian product of G and H.

The tensor product P of the graphs G and H has a node set that is the Cartesian product of the node sets, V(P)=V(G) imes V(H). P has an edge ((u,v),(x,y)) if and only if (u,v) is an edge in G and x==y or and (x,y) is an edge in H and u==v. and (x,y) is an edge in H.

## Parameters G, H: graphs

Networkx graphs.

## Returns P: NetworkX graph

The Cartesian product of G and H. P will be a multi-graph if either G or H is a multi-graph. Will be a directed if G and H are directed, and undirected if G and H are undirected.

## Raises NetworkXError

If G and H are not both directed or both undirected.

#### **Notes**

Node attributes in P are two-tuple of the G and H node attributes. Missing attributes are assigned None.

```
For example >>> G = nx.Graph() >>> H = nx.Graph() >>> G.add\_node(0,a1=True) >>> H.add\_node(`a',a2=`Spam') >>> P = nx.cartesian\_product(G,H) >>> P.nodes()[(0, `a')]
```

Edge attributes and edge keys (for multigraphs) are also copied to the new product graph

# 4.32.14 lexicographic\_product

## $lexicographic_product(G, H)$

Return the lexicographic product of G and H.

The lexicographical product P of the graphs G and H has a node set that is the Cartesian product of the node sets, V(P)=V(G) imes V(H). P has an edge ((u,v),(x,y)) if and only if (u,v) is an edge in G or u==v and (x,y) is an edge in H.

#### Parameters G, H: graphs

Networkx graphs.

## Returns P: NetworkX graph

The Cartesian product of G and H. P will be a multi-graph if either G or H is a multi-graph. Will be a directed if G and H are directed, and undirected if G and H are undirected.

#### Raises NetworkXError

If G and H are not both directed or both undirected.

#### **Notes**

Node attributes in P are two-tuple of the G and H node attributes. Missing attributes are assigned None.

```
For example >>> G = nx.Graph() >>> H = nx.Graph() >>> G.add\_node(0,a1=True) >>> H.add\_node(`a',a2=`Spam') >>> P = nx.lexicographic\_product(G,H) >>> P.nodes()[(0, `a')]
```

Edge attributes and edge keys (for multigraphs) are also copied to the new product graph

# 4.32.15 strong product

#### $strong_product(G, H)$

Return the strong product of G and H.

The strong product P of the graphs G and H has a node set that is the Cartesian product of the node sets, V(P)=V(G) imes V(H). P has an edge ((u,v),(x,y)) if and only if u==v and (x,y) is an edge in H, or x==y and (u,v) is an edge in G, or (u,v) is an edge in G and (x,y) is an edge in H.

## Parameters G, H: graphs

Networkx graphs.

## **Returns** P: NetworkX graph

The Cartesian product of G and H. P will be a multi-graph if either G or H is a multi-graph. Will be a directed if G and H are directed, and undirected if G and H are undirected.

4.32. Operators 311

#### Raises NetworkXError

If G and H are not both directed or both undirected.

#### **Notes**

Node attributes in P are two-tuple of the G and H node attributes. Missing attributes are assigned None.

```
For example >>> G = nx.Graph() >>> H = nx.Graph() >>> G.add_node(0,a1=True) >>> H.add_node('a',a2='Spam') >>> P = nx.strong_product(G,H) >>> P.nodes() [(0, 'a')]
```

Edge attributes and edge keys (for multigraphs) are also copied to the new product graph

# 4.32.16 tensor\_product

## $tensor\_product(G, H)$

Return the tensor product of G and H.

The tensor product P of the graphs G and H has a node set that is the Cartesian product of the node sets, V(P)=V(G) times V(H). P has an edge ((u,v),(x,y)) if and only if (u,v) is an edge in G and (x,y) is an edge in H

Sometimes referred to as the categorical product.

## Parameters G, H: graphs

Networkx graphs.

#### Returns P: NetworkX graph

The tensor product of G and H. P will be a multi-graph if either G or H is a multi-graph. Will be a directed if G and H are directed, and undirected if G and H are undirected.

#### Raises NetworkXError

If G and H are not both directed or both undirected.

#### **Notes**

Node attributes in P are two-tuple of the G and H node attributes. Missing attributes are assigned None.

```
For example >>> G = nx.Graph() >>> H = nx.Graph() >>> G.add\_node(0,a1=True) >>> H.add\_node(`a',a2=`Spam') >>> P = nx.tensor\_product(G,H) >>> P.nodes()[(0, `a')]
```

Edge attributes and edge keys (for multigraphs) are also copied to the new product graph

# 4.33 Rich Club

rich\_club\_coefficient(G[, normalized, Q]) Return the rich-club coefficient of the graph G.

# 4.33.1 rich club coefficient

```
rich_club_coefficient (G, normalized=True, Q=100)
```

Return the rich-club coefficient of the graph G.

The rich-club coefficient is the ratio, for every degree k, of the number of actual to the number of potential edges for nodes with degree greater than k:

$$\phi(k) = \frac{2Ek}{Nk(Nk - 1)}$$

where Nk is the number of nodes with degree larger than k, and Ek be the number of edges among those nodes.

Parameters G: NetworkX graph

normalized: bool (optional)

Normalize using randomized network (see [R313])

**Q**: float (optional, default=100)

If normalized=True build a random network by performing Q\*M double-edge swaps, where M is the number of edges in G, to use as a null-model for normalization.

Returns rc: dictionary

A dictionary, keyed by degree, with rich club coefficient values.

## **Notes**

The rich club definition and algorithm are found in [R313]. This algorithm ignores any edge weights and is not defined for directed graphs or graphs with parallel edges or self loops.

Estimates for appropriate values of Q are found in [R314].

## References

[R313], [R314]

## **Examples**

```
>>> G = nx.Graph([(0,1),(0,2),(1,2),(1,3),(1,4),(4,5)])
>>> rc = nx.rich_club_coefficient(G,normalized=False)
>>> rc[0]
0.4
```

# 4.34 Shortest Paths

Compute the shortest paths and path lengths between nodes in the graph.

These algorithms work with undirected and directed graphs.

<pre>shortest_path(G[, source, target, weight])</pre>	Compute shortest paths in the graph.
<pre>all_shortest_paths(G, source, target[, weight])</pre>	Compute all shortest paths in the graph.
shortest_path_length(G[, source, target, weight])	Compute shortest path lengths in the graph.
$average\_shortest\_path\_length(G[, weight])$	Return the average shortest path length.
has_path(G, source, target)	Return True if G has a path from source to target, False otherwise.

4.34. Shortest Paths 313

# 4.34.1 shortest path

**shortest\_path** (*G*, *source=None*, *target=None*, *weight=None*) Compute shortest paths in the graph.

# Parameters G: NetworkX graph

source: node, optional

Starting node for path. If not specified, compute shortest paths using all nodes as source nodes.

target: node, optional

Ending node for path. If not specified, compute shortest paths using all nodes as target nodes.

**weight**: None or string, optional (default = None)

If None, every edge has weight/distance/cost 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

## Returns path: list or dictionary

All returned paths include both the source and target in the path.

If the source and target are both specified, return a single list of nodes in a shortest path from the source to the target.

If only the source is specified, return a dictionary keyed by targets with a list of nodes in a shortest path from the source to one of the targets.

If only the target is specified, return a dictionary keyed by sources with a list of nodes in a shortest path from one of the sources to the target.

If neither the source nor target are specified return a dictionary of dictionaries with path[source][target]=[list of nodes in path].

#### See also:

```
all_pairs_shortest_path, all_pairs_dijkstra_path, single_source_shortest_path, single_source_dijkstra_path
```

## **Notes**

There may be more than one shortest path between a source and target. This returns only one of them.

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> print(nx.shortest_path(G,source=0,target=4))
[0, 1, 2, 3, 4]
>>> p=nx.shortest_path(G,source=0) # target not specified
>>> p[4]
[0, 1, 2, 3, 4]
>>> p=nx.shortest_path(G,target=4) # source not specified
>>> p[0]
[0, 1, 2, 3, 4]
>>> p=nx.shortest_path(G) # source,target not specified
```

```
>>> p[0][4]
[0, 1, 2, 3, 4]
```

# 4.34.2 all shortest paths

all\_shortest\_paths (G, source, target, weight=None)

Compute all shortest paths in the graph.

**Parameters G**: NetworkX graph

source: node

Starting node for path.

target: node

Ending node for path.

**weight**: None or string, optional (default = None)

If None, every edge has weight/distance/cost 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

Returns paths: generator of lists

A generator of all paths between source and target.

#### See also:

```
shortest_path, single_source_shortest_path, all_pairs_shortest_path
```

#### Notes

There may be many shortest paths between the source and target.

## **Examples**

```
>>> G=nx.Graph()
>>> G.add_path([0,1,2])
>>> G.add_path([0,10,2])
>>> print([p for p in nx.all_shortest_paths(G,source=0,target=2)])
[[0, 1, 2], [0, 10, 2]]
```

# 4.34.3 shortest path length

shortest\_path\_length(G, source=None, target=None, weight=None)

Compute shortest path lengths in the graph.

**Parameters G**: NetworkX graph

source: node, optional

Starting node for path. If not specified, compute shortest path lengths using all nodes as source nodes.

target: node, optional

4.34. Shortest Paths 315

Ending node for path. If not specified, compute shortest path lengths using all nodes as target nodes.

```
weight: None or string, optional (default = None)
```

If None, every edge has weight/distance/cost 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

## Returns length: int or dictionary

If the source and target are both specified, return the length of the shortest path from the source to the target.

If only the source is specified, return a dictionary keyed by targets whose values are the lengths of the shortest path from the source to one of the targets.

If only the target is specified, return a dictionary keyed by sources whose values are the lengths of the shortest path from one of the sources to the target.

If neither the source nor target are specified return a dictionary of dictionaries with path[source][target]=L, where L is the length of the shortest path from source to target.

#### Raises NetworkXNoPath

If no path exists between source and target.

#### See also:

```
all_pairs_shortest_path_length, all_pairs_dijkstra_path_length, single_source_dijkstra_path_length
```

#### **Notes**

The length of the path is always 1 less than the number of nodes involved in the path since the length measures the number of edges followed.

For digraphs this returns the shortest directed path length. To find path lengths in the reverse direction use G.reverse(copy=False) first to flip the edge orientation.

## **Examples**

```
>>> G=nx.path_graph(5)
>>> print(nx.shortest_path_length(G, source=0, target=4))
4
>>> p=nx.shortest_path_length(G, source=0) # target not specified
>>> p[4]
4
>>> p=nx.shortest_path_length(G, target=4) # source not specified
>>> p[0]
4
>>> p=nx.shortest_path_length(G) # source, target not specified
>>> p[0][4]
4
```

# 4.34.4 average\_shortest\_path\_length

```
average_shortest_path_length (G, weight=None)
Return the average shortest path length.
```

The average shortest path length is

$$a = \sum_{s,t \in V} \frac{d(s,t)}{n(n-1)}$$

where V is the set of nodes in G, d(s,t) is the shortest path from s to t, and n is the number of nodes in G.

Parameters G: NetworkX graph

**weight**: None or string, optional (default = None)

If None, every edge has weight/distance/cost 1. If a string, use this edge attribute as the edge weight. Any edge attribute not present defaults to 1.

## **Raises NetworkXError:**

if the graph is not connected.

# **Examples**

```
>>> G=nx.path_graph(5)
>>> print(nx.average_shortest_path_length(G))
2.0
```

For disconnected graphs you can compute the average shortest path length for each component: >>> G=nx.Graph([(1,2),(3,4)]) >>> for g in  $nx.connected\_component\_subgraphs(G)$ : ...  $print(nx.average\_shortest\_path\_length(g))$  1.0 1.0

# 4.34.5 has\_path

has\_path(G, source, target)

Return True if G has a path from source to target, False otherwise.

**Parameters G**: NetworkX graph

source: node

Starting node for path

target : node

Ending node for path

# 4.34.6 Advanced Interface

Shortest path algorithms for unweighted graphs.

$\verb single_source_shortest_path  (G, source[, cutoff]) \\$	Compute shortest path between source and all other nodes reachable
$single\_source\_shortest\_path\_length(G, source)$	Compute the shortest path lengths from source to all reachable nodes
${\tt all\_pairs\_shortest\_path}(G[,cutoff])$	Compute shortest paths between all nodes.
${ t all\_pairs\_shortest\_path\_length}(G[, cutoff])$	Compute the shortest path lengths between all nodes in G.
predecessor(G, source[, target, cutoff,])	Returns dictionary of predecessors for the path from source to all not

4.34. Shortest Paths 317

## single source shortest path

## single\_source\_shortest\_path(G, source, cutoff=None)

Compute shortest path between source and all other nodes reachable from source.

**Parameters G**: NetworkX graph

source: node label

Starting node for path **cutoff**: integer, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

**Returns lengths**: dictionary

Dictionary, keyed by target, of shortest paths.

## See also:

shortest\_path

#### **Notes**

The shortest path is not necessarily unique. So there can be multiple paths between the source and each target node, all of which have the same 'shortest' length. For each target node, this function returns only one of those paths.

## **Examples**

```
>>> G=nx.path_graph(5)
>>> path=nx.single_source_shortest_path(G,0)
>>> path[4]
[0, 1, 2, 3, 4]
```

# single\_source\_shortest\_path\_length

# $\verb|single_source_shortest_path_length| (\textit{G}, \textit{source}, \textit{cutoff=None})$

Compute the shortest path lengths from source to all reachable nodes.

**Parameters G**: NetworkX graph

source : node

Starting node for path

cutoff: integer, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns lengths: dictionary

Dictionary of shortest path lengths keyed by target.

# See also:

```
shortest_path_length
```

### **Examples**

```
>>> G=nx.path_graph(5)
>>> length=nx.single_source_shortest_path_length(G,0)
>>> length[4]
4
>>> print(length)
{0: 0, 1: 1, 2: 2, 3: 3, 4: 4}
```

### all\_pairs\_shortest\_path

### all\_pairs\_shortest\_path(G, cutoff=None)

Compute shortest paths between all nodes.

Parameters G: NetworkX graph

cutoff : integer, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns lengths: dictionary

Dictionary, keyed by source and target, of shortest paths.

#### See also:

```
floyd_warshall
```

### **Examples**

```
>>> G=nx.path_graph(5)
>>> path=nx.all_pairs_shortest_path(G)
>>> print(path[0][4])
[0, 1, 2, 3, 4]
```

### all\_pairs\_shortest\_path\_length

### all\_pairs\_shortest\_path\_length(G, cutoff=None)

Compute the shortest path lengths between all nodes in G.

**Parameters G**: NetworkX graph

cutoff: integer, optional

depth to stop the search. Only paths of length <= cutoff are returned.

Returns lengths: dictionary

Dictionary of shortest path lengths keyed by source and target.

#### **Notes**

The dictionary returned only has keys for reachable node pairs.

4.34. Shortest Paths 319

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> length=nx.all_pairs_shortest_path_length(G)
>>> print(length[1][4])
3
>>> length[1]
{0: 1, 1: 0, 2: 1, 3: 2, 4: 3}
```

### predecessor

**predecessor** (*G*, *source*, *target=None*, *cutoff=None*, *return\_seen=None*)

Returns dictionary of predecessors for the path from source to all nodes in G.

**Parameters G** : NetworkX graph

source: node label

Starting node for path

target: node label, optional

Ending node for path. If provided only predecessors between source and target are

returned

cutoff : integer, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns pred: dictionary

Dictionary, keyed by node, of predecessors in the shortest path.

### **Examples**

```
>>> G=nx.path_graph(4)
>>> print(G.nodes())
[0, 1, 2, 3]
>>> nx.predecessor(G,0)
{0: [], 1: [0], 2: [1], 3: [2]}
```

Shortest path algorithms for weighed graphs.

<pre>dijkstra_path(G, source, target[, weight])</pre>	Returns the shortest path from source to target in a weighted graph G
<pre>dijkstra_path_length(G, source, target[, weight])</pre>	Returns the shortest path length from source to target in a weighted g
$single\_source\_dijkstra\_path(G, source[,])$	Compute shortest path between source and all other reachable nodes
$single\_source\_dijkstra\_path\_length(G, source)$	Compute the shortest path length between source and all other reacha
$all\_pairs\_dijkstra\_path(G[, cutoff, weight])$	Compute shortest paths between all nodes in a weighted graph.
all_pairs_dijkstra_path_length( $G[, cutoff,]$ )	Compute shortest path lengths between all nodes in a weighted graph
single_source_dijkstra(G, source[, target,])	Compute shortest paths and lengths in a weighted graph G.
$\verb bidirectional_dijkstra  (G, source, target[,]) $	Dijkstra's algorithm for shortest paths using bidirectional search.
$ ext{dijkstra\_predecessor\_and\_distance}(G, source)$	Compute shortest path length and predecessors on shortest paths in w
bellman_ford(G, source[, weight])	Compute shortest path lengths and predecessors on shortest paths in
$\verb"negative_edge_cycle(G[, weight])"$	Return True if there exists a negative edge cycle anywhere in G.

### dijkstra\_path

```
dijkstra_path(G, source, target, weight='weight')
     Returns the shortest path from source to target in a weighted graph G.
          Parameters G: NetworkX graph
               source: node
                  Starting node
               target: node
                  Ending node
               weight: string, optional (default='weight')
                  Edge data key corresponding to the edge weight
          Returns path: list
                  List of nodes in a shortest path.
          Raises NetworkXNoPath
                  If no path exists between source and target.
     See also:
     bidirectional_dijkstra
     Notes
     Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.
     Examples
     >>> G=nx.path_graph(5)
     >>> print (nx.dijkstra_path(G,0,4))
     [0, 1, 2, 3, 4]
dijkstra_path_length
dijkstra_path_length(G, source, target, weight='weight')
     Returns the shortest path length from source to target in a weighted graph.
          Parameters G: NetworkX graph
               source: node label
                  starting node for path
```

Returns length: number

Edge data key corresponding to the edge weight

weight: string, optional (default='weight')

target: node label

ending node for path

4.34. Shortest Paths 321

Shortest path length.

#### Raises NetworkXNoPath

If no path exists between source and target.

#### See also:

```
bidirectional_dijkstra
```

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> print(nx.dijkstra_path_length(G,0,4))
```

### single\_source\_dijkstra\_path

```
single_source_dijkstra_path(G, source, cutoff=None, weight='weight')
```

Compute shortest path between source and all other reachable nodes for a weighted graph.

```
Parameters G: NetworkX graph
```

source: node

Starting node for path.

### weight: string, optional (default='weight')

Edge data key corresponding to the edge weight

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns paths: dictionary

Dictionary of shortest path lengths keyed by target.

### See also:

```
single_source_dijkstra
```

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

### **Examples**

```
>>> G=nx.path_graph(5)
>>> path=nx.single_source_dijkstra_path(G,0)
>>> path[4]
[0, 1, 2, 3, 4]
```

### single\_source\_dijkstra\_path\_length

```
single_source_dijkstra_path_length(G, source, cutoff=None, weight='weight')
```

Compute the shortest path length between source and all other reachable nodes for a weighted graph.

```
Parameters G: NetworkX graph
```

source: node label

Starting node for path

#### weight: string, optional (default='weight')

Edge data key corresponding to the edge weight.

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns length: dictionary

Dictionary of shortest lengths keyed by target.

#### See also:

```
single_source_dijkstra
```

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

### **Examples**

```
>>> G=nx.path_graph(5)
>>> length=nx.single_source_dijkstra_path_length(G,0)
>>> length[4]
4
>>> print(length)
{0: 0, 1: 1, 2: 2, 3: 3, 4: 4}
```

#### all\_pairs\_dijkstra\_path

```
all_pairs_dijkstra_path(G, cutoff=None, weight='weight')
```

Compute shortest paths between all nodes in a weighted graph.

**Parameters G**: NetworkX graph

weight: string, optional (default='weight')

Edge data key corresponding to the edge weight

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns distance: dictionary

Dictionary, keyed by source and target, of shortest paths.

### See also:

```
floyd_warshall
```

4.34. Shortest Paths 323

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

### **Examples**

```
>>> G=nx.path_graph(5)
>>> path=nx.all_pairs_dijkstra_path(G)
>>> print(path[0][4])
[0, 1, 2, 3, 4]
```

### all\_pairs\_dijkstra\_path\_length

```
\verb"all_pairs_dijkstra_path_length" (\textit{G}, \textit{cutoff=None}, \textit{weight='weight'})
```

Compute shortest path lengths between all nodes in a weighted graph.

Parameters G: NetworkX graph

```
weight: string, optional (default='weight')
```

Edge data key corresponding to the edge weight

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns distance: dictionary

Dictionary, keyed by source and target, of shortest path lengths.

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

The dictionary returned only has keys for reachable node pairs.

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> length=nx.all_pairs_dijkstra_path_length(G)
>>> print(length[1][4])
3
>>> length[1]
{0: 1, 1: 0, 2: 1, 3: 2, 4: 3}
```

## single\_source\_dijkstra

```
single_source_dijkstra(G, source, target=None, cutoff=None, weight='weight')
```

Compute shortest paths and lengths in a weighted graph G.

Uses Dijkstra's algorithm for shortest paths.

Parameters G: NetworkX graph

source: node label

```
Starting node for path
```

target : node label, optional

Ending node for path

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

### Returns distance, path: dictionaries

Returns a tuple of two dictionaries keyed by node. The first dictionary stores distance from the source. The second stores the path from the source to that node.

#### See also:

```
single_source_dijkstra_path, single_source_dijkstra_path_length
```

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

Based on the Python cookbook recipe (119466) at http://aspn.activestate.com/ASPN/Cookbook/Python/Recipe/119466

This algorithm is not guaranteed to work if edge weights are negative or are floating point numbers (overflows and roundoff errors can cause problems).

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> length,path=nx.single_source_dijkstra(G,0)
>>> print(length[4])
4
>>> print(length)
{0: 0, 1: 1, 2: 2, 3: 3, 4: 4}
>>> path[4]
[0, 1, 2, 3, 4]
```

### bidirectional\_dijkstra

```
bidirectional_dijkstra (G, source, target, weight='weight')

Dijkstra's algorithm for shortest paths using bidirectional search.
```

**Parameters G**: NetworkX graph

source: node

Starting node.

target: node

Ending node.

weight: string, optional (default='weight')

Edge data key corresponding to the edge weight

**Returns length**: number

Shortest path length.

4.34. Shortest Paths 325

Returns a tuple of two dictionaries keyed by node.

The first dictionary stores distance from the source.

The second stores the path from the source to that node.

#### Raises NetworkXNoPath

If no path exists between source and target.

#### See also:

```
shortest_path, shortest_path_length
```

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

In practice bidirectional Dijkstra is much more than twice as fast as ordinary Dijkstra.

Ordinary Dijkstra expands nodes in a sphere-like manner from the source. The radius of this sphere will eventually be the length of the shortest path. Bidirectional Dijkstra will expand nodes from both the source and the target, making two spheres of half this radius. Volume of the first sphere is pi\*r\*r while the others are 2\*pi\*r/2\*r/2, making up half the volume.

This algorithm is not guaranteed to work if edge weights are negative or are floating point numbers (overflows and roundoff errors can cause problems).

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> length,path=nx.bidirectional_dijkstra(G,0,4)
>>> print(length)
4
>>> print(path)
[0, 1, 2, 3, 4]
```

#### dijkstra predecessor and distance

```
\label{linear_distance} \begin{tabular}{ll} \bf dijkstra\_predecessor\_and\_distance (\it{G}, source, cutoff=None, weight='weight') \\ \end{tabular}
```

Compute shortest path length and predecessors on shortest paths in weighted graphs.

```
Parameters G: NetworkX graph
source: node label
Starting node for path
```

weight: string, optional (default='weight')

Edge data key corresponding to the edge weight

cutoff: integer or float, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

Returns pred, distance: dictionaries

Returns two dictionaries representing a list of predecessors of a node and the distance to each node.

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

The list of predecessors contains more than one element only when there are more than one shortest paths to the key node.

#### bellman ford

```
bellman_ford(G, source, weight='weight')
```

Compute shortest path lengths and predecessors on shortest paths in weighted graphs.

The algorithm has a running time of O(mn) where n is the number of nodes and m is the number of edges. It is slower than Dijkstra but can handle negative edge weights.

#### **Parameters G**: NetworkX graph

The algorithm works for all types of graphs, including directed graphs and multigraphs.

source: node label

Starting node for path

```
weight: string, optional (default='weight')
```

Edge data key corresponding to the edge weight

#### **Returns pred, dist**: dictionaries

Returns two dictionaries keyed by node to predecessor in the path and to the distance from the source respectively.

### Raises NetworkXUnbounded

If the (di)graph contains a negative cost (di)cycle, the algorithm raises an exception to indicate the presence of the negative cost (di)cycle. Note: any negative weight edge in an undirected graph is a negative cost cycle.

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

The dictionaries returned only have keys for nodes reachable from the source.

In the case where the (di)graph is not connected, if a component not containing the source contains a negative cost (di)cycle, it will not be detected.

### **Examples**

```
>>> import networkx as nx
>>> G = nx.path_graph(5, create_using = nx.DiGraph())
>>> pred, dist = nx.bellman_ford(G, 0)
>>> sorted(pred.items())
[(0, None), (1, 0), (2, 1), (3, 2), (4, 3)]
>>> sorted(dist.items())
[(0, 0), (1, 1), (2, 2), (3, 3), (4, 4)]
```

4.34. Shortest Paths 327

```
>>> from nose.tools import assert_raises
>>> G = nx.cycle_graph(5, create_using = nx.DiGraph())
>>> G[1][2]['weight'] = -7
>>> assert_raises(nx.NetworkXUnbounded, nx.bellman_ford, G, 0)
```

#### negative\_edge\_cycle

```
negative_edge_cycle (G, weight='weight')
```

Return True if there exists a negative edge cycle anywhere in G.

**Parameters G**: NetworkX graph

weight: string, optional (default='weight')

Edge data key corresponding to the edge weight

Returns negative\_cycle: bool

True if a negative edge cycle exists, otherwise False.

#### **Notes**

Edge weight attributes must be numerical. Distances are calculated as sums of weighted edges traversed.

This algorithm uses bellman\_ford() but finds negative cycles on any component by first adding a new node connected to every node, and starting bellman\_ford on that node. It then removes that extra node.

#### **Examples**

```
>>> import networkx as nx
>>> G = nx.cycle_graph(5, create_using = nx.DiGraph())
>>> print(nx.negative_edge_cycle(G))
False
>>> G[1][2]['weight'] = -7
>>> print(nx.negative_edge_cycle(G))
True
```

## 4.34.7 Dense Graphs

Floyd-Warshall algorithm for shortest paths.

floyd_warshall(G[, weight])	Find all-pairs shortest path lengths using Floyd's algorithm.
floyd_warshall_predecessor_and_distance( $G[,]$ )	Find all-pairs shortest path lengths using Floyd's algorithm.
floyd_warshall_numpy(G[, nodelist, weight])	Find all-pairs shortest path lengths using Floyd's algorithm.

### floyd\_warshall

```
floyd_warshall (G, weight='weight')
```

Find all-pairs shortest path lengths using Floyd's algorithm.

Parameters G: NetworkX graph

weight: string, optional (default= 'weight')

Edge data key corresponding to the edge weight.

Returns distance: dict

A dictionary, keyed by source and target, of shortest paths distances between nodes.

#### See also:

#### **Notes**

Floyd's algorithm is appropriate for finding shortest paths in dense graphs or graphs with negative weights when Dijkstra's algorithm fails. This algorithm can still fail if there are negative cycles. It has running time  $O(n^3)$  with running space of  $O(n^2)$ .

### floyd\_warshall\_predecessor\_and\_distance

```
floyd_warshall_predecessor_and_distance(G, weight='weight')
```

Find all-pairs shortest path lengths using Floyd's algorithm.

**Parameters G**: NetworkX graph

weight: string, optional (default= 'weight')

Edge data key corresponding to the edge weight.

Returns predecessor, distance: dictionaries

Dictionaries, keyed by source and target, of predecessors and distances in the shortest path.

#### See also:

```
floyd_warshall, floyd_warshall_numpy, all_pairs_shortest_path, all_pairs_shortest_path_length
```

#### **Notes**

Floyd's algorithm is appropriate for finding shortest paths in dense graphs or graphs with negative weights when Dijkstra's algorithm fails. This algorithm can still fail if there are negative cycles. It has running time  $O(n^3)$  with running space of  $O(n^2)$ .

### floyd\_warshall\_numpy

### floyd\_warshall\_numpy (G, nodelist=None, weight='weight')

Find all-pairs shortest path lengths using Floyd's algorithm.

Parameters G: NetworkX graph

**nodelist**: list, optional

The rows and columns are ordered by the nodes in nodelist. If nodelist is None then the ordering is produced by G.nodes().

weight: string, optional (default= 'weight')

Edge data key corresponding to the edge weight.

4.34. Shortest Paths 329

### Returns distance: NumPy matrix

A matrix of shortest path distances between nodes. If there is no path between to nodes the corresponding matrix entry will be Inf.

#### **Notes**

Floyd's algorithm is appropriate for finding shortest paths in dense graphs or graphs with negative weights when Dijkstra's algorithm fails. This algorithm can still fail if there are negative cycles. It has running time  $O(n^3)$  with running space of  $O(n^2)$ .

## 4.34.8 A\* Algorithm

Shortest paths and path lengths using A\* ("A star") algorithm.

```
astar_path(G, source, target[, heuristic, ...]) Return a list of nodes in a shortest path between source and target using the A* ("A-star_path_length(G, source, target[, ...]) Return the length of the shortest path between source and target using the A* ("A-star_path_length(G, source, target[, ...])
```

### astar\_path

```
astar_path (G, source, target, heuristic=None, weight='weight')
```

Return a list of nodes in a shortest path between source and target using the A\* ("A-star") algorithm.

There may be more than one shortest path. This returns only one.

#### **Parameters G**: NetworkX graph

source: node

Starting node for path

target: node

Ending node for path

heuristic: function

A function to evaluate the estimate of the distance from the a node to the target. The function takes two nodes arguments and must return a number.

### weight: string, optional (default='weight')

Edge data key corresponding to the edge weight.

#### Raises NetworkXNoPath

If no path exists between source and target.

### See also:

```
shortest_path, dijkstra_path
```

#### **Examples**

```
>>> G=nx.path_graph(5)
>>> print(nx.astar_path(G,0,4))
[0, 1, 2, 3, 4]
>>> G=nx.grid_graph(dim=[3,3]) # nodes are two-tuples (x,y)
```

```
>>> def dist(a, b):
... (x1, y1) = a
... (x2, y2) = b
... return ((x1 - x2) ** 2 + (y1 - y2) ** 2) ** 0.5
>>> print(nx.astar_path(G, (0, 0), (2, 2), dist))
[(0, 0), (0, 1), (1, 1), (1, 2), (2, 2)]
```

### astar\_path\_length

astar\_path\_length(G, source, target, heuristic=None, weight='weight')

Return the length of the shortest path between source and target using the A\* ("A-star") algorithm.

**Parameters G**: NetworkX graph

source: node

Starting node for path

target: node

Ending node for path

heuristic: function

A function to evaluate the estimate of the distance from the a node to the target. The function takes two nodes arguments and must return a number.

#### Raises NetworkXNoPath

If no path exists between source and target.

### See also:

astar\_path

# 4.35 Simple Paths

all\_simple\_paths(G, source, target[, cutoff]) Generate all simple paths in the graph G from source to target.

## 4.35.1 all\_simple\_paths

```
all_simple_paths (G, source, target, cutoff=None)
```

Generate all simple paths in the graph G from source to target.

A simple path is a path with no repeated nodes.

**Parameters G**: NetworkX graph

source: node

Starting node for path

target: node

Ending node for path

cutoff : integer, optional

Depth to stop the search. Only paths of length <= cutoff are returned.

4.35. Simple Paths 331

**Returns** path\_generator: generator

A generator that produces lists of simple paths. If there are no paths between the source and target within the given cutoff the generator produces no output.

#### See also:

```
all_shortest_paths, shortest_path
```

#### **Notes**

This algorithm uses a modified depth-first search to generate the paths [R315]. A single path can be found in O(V+E) time but the number of simple paths in a graph can be very large, e.g. O(n!) in the complete graph of order n.

#### References

[R315]

#### **Examples**

```
>>> G = nx.complete_graph(4)
>>> for path in nx.all_simple_paths(G, source=0, target=3):
...     print(path)
...
[0, 1, 2, 3]
[0, 1, 3]
[0, 2, 1, 3]
[0, 2, 3]
[0, 3]
>>> paths = nx.all_simple_paths(G, source=0, target=3, cutoff=2)
>>> print(list(paths))
[[0, 1, 3], [0, 2, 3], [0, 3]]
```

# 4.36 Swap

Swap edges in a graph.

double_edge_swap(G[, nswap, max_tries])	Swap two edges in the graph while keeping the node degrees fixed.
$connected\_double\_edge\_swap(G[, nswap])$	Attempt nswap double-edge swaps in the graph G.

## 4.36.1 double\_edge\_swap

```
double\_edge\_swap(G, nswap=1, max\_tries=100)
```

Swap two edges in the graph while keeping the node degrees fixed.

A double-edge swap removes two randomly chosen edges u-v and x-y and creates the new edges u-x and v-y:

If either the edge u-x or v-y already exist no swap is performed and another attempt is made to find a suitable edge pair.

**Parameters G**: graph

An undirected graph

nswap : integer (optional, default=1)

Number of double-edge swaps to perform

max\_tries : integer (optional)

Maximum number of attempts to swap edges

Returns G: graph

The graph after double edge swaps.

#### **Notes**

Does not enforce any connectivity constraints.

The graph G is modified in place.

## 4.36.2 connected double edge swap

### connected\_double\_edge\_swap(G, nswap=1)

Attempt nswap double-edge swaps in the graph G.

A double-edge swap removes two randomly chosen edges u-v and x-y and creates the new edges u-x and v-y:

If either the edge u-x or v-y already exist no swap is performed so the actual count of swapped edges is always <= nswap

Parameters G: graph

An undirected graph

nswap : integer (optional, default=1)

Number of double-edge swaps to perform

Returns G: int

The number of successful swaps

#### **Notes**

The initial graph G must be connected, and the resulting graph is connected. The graph G is modified in place.

#### References

[R316]

4.36. Swap 333

## 4.37 Traversal

## 4.37.1 Depth First Search

Basic algorithms for depth-first searching the nodes of a graph.

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

dfs_edges(G[, source])	Produce edges in a depth-first-search (DFS).
dfs_tree(G, source)	Return oriented tree constructed from a depth-first-search from source.
$dfs\_predecessors(G[, source])$	Return dictionary of predecessors in depth-first-search from source.
$dfs\_successors(G[, source])$	Return dictionary of successors in depth-first-search from source.
$dfs\_preorder\_nodes(G[, source])$	Produce nodes in a depth-first-search pre-ordering starting from source.
$dfs_postorder_nodes(G[, source])$	Produce nodes in a depth-first-search post-ordering starting from source.
$dfs_{abeled\_edges}(G[, source])$	Produce edges in a depth-first-search (DFS) labeled by type.

### dfs\_edges

### dfs\_edges (G, source=None)

Produce edges in a depth-first-search (DFS).

Parameters G: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

Returns edges: generator

A generator of edges in the depth-first-search.

### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(list(nx.dfs_edges(G,0)))
[(0, 1), (1, 2)]
```

### dfs\_tree

### dfs\_tree(G, source)

Return oriented tree constructed from a depth-first-search from source.

Parameters G: NetworkX graph

source: node, optional

Specify starting node for depth-first search.

### Returns T: NetworkX DiGraph

An oriented tree

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> T = nx.dfs_tree(G,0)
>>> print(T.edges())
[(0, 1), (1, 2)]
```

### dfs predecessors

### dfs\_predecessors(G, source=None)

Return dictionary of predecessors in depth-first-search from source.

**Parameters G**: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

Returns pred: dict

A dictionary with nodes as keys and predecessor nodes as values.

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(nx.dfs_predecessors(G,0))
{1: 0, 2: 1}
```

### dfs\_successors

### dfs\_successors(G, source=None)

Return dictionary of successors in depth-first-search from source.

Parameters G: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

4.37. Traversal 335

#### Returns succ: dict

A dictionary with nodes as keys and list of successor nodes as values.

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(nx.dfs_successors(G,0))
{0: [1], 1: [2]}
```

### dfs\_preorder\_nodes

```
dfs preorder nodes(G, source=None)
```

Produce nodes in a depth-first-search pre-ordering starting from source.

Parameters G: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

Returns nodes: generator

A generator of nodes in a depth-first-search pre-ordering.

### Notes

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(list(nx.dfs_preorder_nodes(G,0)))
[0, 1, 2]
```

#### dfs\_postorder\_nodes

```
dfs_postorder_nodes (G, source=None)
```

Produce nodes in a depth-first-search post-ordering starting from source.

**Parameters G**: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

Returns nodes: generator

A generator of nodes in a depth-first-search post-ordering.

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(list(nx.dfs_postorder_nodes(G,0)))
[2, 1, 0]
```

### dfs\_labeled\_edges

### dfs\_labeled\_edges(G, source=None)

Produce edges in a depth-first-search (DFS) labeled by type.

Parameters G: NetworkX graph

source: node, optional

Specify starting node for depth-first search and return edges in the component reachable from source.

Returns edges: generator

A generator of edges in the depth-first-search labeled with 'forward', 'nontree', and 'reverse'.

### Notes

Based on http://www.ics.uci.edu/~eppstein/PADS/DFS.py by D. Eppstein, July 2004.

If a source is not specified then a source is chosen arbitrarily and repeatedly until all components in the graph are searched.

### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> edges = (list(nx.dfs_labeled_edges(G,0)))
```

4.37. Traversal 337

### 4.37.2 Breadth First Search

Basic algorithms for breadth-first searching the nodes of a graph.

<pre>bfs_edges(G, source[, reverse])</pre>	) Produce edges in a breadth-first-search starting at source.	
<pre>bfs_tree(G, source[, reverse])</pre>	Return an oriented tree constructed from of a breadth-first-search starting at source.	
bfs_predecessors(G, source)	Return dictionary of predecessors in breadth-first-search from source.	
bfs_successors(G, source)	Return dictionary of successors in breadth-first-search from source.	

### bfs edges

bfs\_edges (G, source, reverse=False)

Produce edges in a breadth-first-search starting at source.

Parameters G: NetworkX graph

source: node

Specify starting node for breadth-first search and return edges in the component reachable from source.

reverse: bool, optional

If True traverse a directed graph in the reverse direction

Returns edges: generator

A generator of edges in the breadth-first-search.

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/BFS.py by D. Eppstein, July 2004.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(list(nx.bfs_edges(G,0)))
[(0, 1), (1, 2)]
```

### bfs tree

bfs\_tree (G, source, reverse=False)

Return an oriented tree constructed from of a breadth-first-search starting at source.

Parameters G: NetworkX graph

source: node

Specify starting node for breadth-first search and return edges in the component reachable from source.

reverse: bool, optional

If True traverse a directed graph in the reverse direction

Returns T: NetworkX DiGraph

An oriented tree

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/BFS.py by D. Eppstein, July 2004.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(list(nx.bfs_edges(G,0)))
[(0, 1), (1, 2)]
```

### bfs predecessors

### bfs\_predecessors(G, source)

Return dictionary of predecessors in breadth-first-search from source.

Parameters G: NetworkX graph

source: node

Specify starting node for breadth-first search and return edges in the component reachable from source.

Returns pred: dict

A dictionary with nodes as keys and predecessor nodes as values.

### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/BFS.py by D. Eppstein, July 2004.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(nx.bfs_predecessors(G,0))
{1: 0, 2: 1}
```

### bfs\_successors

### bfs\_successors(G, source)

Return dictionary of successors in breadth-first-search from source.

**Parameters G**: NetworkX graph

source: node

Specify starting node for breadth-first search and return edges in the component reachable from source.

Returns succ: dict

4.37. Traversal 339

A dictionary with nodes as keys and list of successors nodes as values.

#### **Notes**

Based on http://www.ics.uci.edu/~eppstein/PADS/BFS.py by D. Eppstein, July 2004.

#### **Examples**

```
>>> G = nx.Graph()
>>> G.add_path([0,1,2])
>>> print(nx.bfs_successors(G,0))
{0: [1], 1: [2]}
```

## 4.37.3 Depth First Search on Edges

Algorithms for a depth-first traversal of edges in a graph.

edge\_dfs(G[, source, orientation]) A directed, depth-first traversal of edges in G, beginning at source.

### edge\_dfs

```
edge_dfs (G, source=None, orientation='original')
```

A directed, depth-first traversal of edges in G, beginning at source.

#### Parameters G: graph

A directed/undirected graph/multigraph.

```
source: node, list of nodes
```

The node from which the traversal begins. If None, then a source is chosen arbitrarily and repeatedly until all edges from each node in the graph are searched.

```
orientation: 'original' | 'reverse' | 'ignore'
```

For directed graphs and directed multigraphs, edge traversals need not respect the original orientation of the edges. When set to 'reverse', then every edge will be traversed in the reverse direction. When set to 'ignore', then each directed edge is treated as a single undirected edge that can be traversed in either direction. For undirected graphs and undirected multigraphs, this parameter is meaningless and is not consulted by the algorithm.

### See also:

```
dfs_edges
```

### **Notes**

The goal of this function is to visit edges. It differs from the more familiar depth-first traversal of nodes, as provided by  $networkx.algorithms.traversal.depth_first_search.dfs_edges(), in that it does not stop once every node has been visited. In a directed graph with edges [(0, 1), (1, 2), (2, 1)], the edge (2, 1) would not be visited if not for the functionality provided by this function.$ 

#### **Examples**

```
>>> import networkx as nx
>>> nodes = [0, 1, 2, 3]
>>> edges = [(0, 1), (1, 0), (1, 0), (2, 1), (3, 1)]
>>> list(nx.edge_dfs(nx.Graph(edges), nodes))
[(0, 1), (1, 2), (1, 3)]
>>> list(nx.edge_dfs(nx.DiGraph(edges), nodes))
[(0, 1), (1, 0), (2, 1), (3, 1)]
>>> list(nx.edge_dfs(nx.MultiGraph(edges), nodes))
[(0, 1, 0), (1, 0, 1), (0, 1, 2), (1, 2, 0), (1, 3, 0)]
>>> list(nx.edge_dfs(nx.MultiDiGraph(edges), nodes))
[(0, 1, 0), (1, 0, 0), (1, 0, 1), (2, 1, 0), (3, 1, 0)]
>>> list(nx.edge_dfs(nx.DiGraph(edges), nodes, orientation='ignore'))
[(0, 1, 'forward'), (1, 0, 'forward'), (2, 1, 'reverse'), (3, 1, 'reverse')]
>>> list(nx.edge_dfs(nx.MultiDiGraph(edges), nodes, orientation='ignore'))
[(0, 1, 0, 'forward'), (1, 0, 0, 'forward'), (1, 0, 1, 'reverse'), (2, 1, 0, 'reverse'), (3, 1, 'reverse'), (3,
```

## 4.38 Tree

## 4.38.1 Recognition

#### **Recognition Tests**

A *forest* is an acyclic, undirected graph, and a *tree* is a connected forest. Depending on the subfield, there are various conventions for generalizing these definitions to directed graphs.

In one convention, directed variants of forest and tree are defined in an identical manner, except that the direction of the edges is ignored. In effect, each directed edge is treated as a single undirected edge. Then, additional restrictions are imposed to define *branchings* and *arborescences*.

In another convention, directed variants of forest and tree correspond to the previous convention's branchings and arborescences, respectively. Then two new terms, *polyforest* and *polytree*, are defined to correspond to the other convention's forest and tree.

#### Summarizing:

Each convention has its reasons. The first convention emphasizes definitional similarity in that directed forests and trees are only concerned with acyclicity and do not have an in-degree constraint, just as their undirected counterparts do not. The second convention emphasizes functional similarity in the sense that the directed analog of a spanning tree is

4.38. Tree 341

a spanning arborescence. That is, take any spanning tree and choose one node as the root. Then every edge is assigned a direction such there is a directed path from the root to every other node. The result is a spanning arborescence.

NetworkX follows convention "A". Explicitly, these are:

undirected forest An undirected graph with no undirected cycles.

undirected tree A connected, undirected forest.

**directed forest** A directed graph with no undirected cycles. Equivalently, the underlying graph structure (which ignores edge orientations) is an undirected forest. In convention B, this is known as a polyforest.

**directed tree** A weakly connected, directed forest. Equivalently, the underlying graph structure (which ignores edge orientations) is an undirected tree. In convention B, this is known as a polytree.

**branching** A directed forest with each node having, at most, one parent. So the maximum in-degree is equal to 1. In convention B, this is known as a forest.

**arborescence** A directed tree with each node having, at most, one parent. So the maximum in-degree is equal to 1. In convention B, this is known as a tree.

For trees and arborescences, the adjective "spanning" may be added to designate that the graph, when considered as a forest/branching, consists of a single tree/arborescence that includes all nodes in the graph. It is true, by definition, that every tree/arborescence is spanning with respect to the nodes that define the tree/arborescence and so, it might seem redundant to introduce the notion of "spanning". However, the nodes may represent a subset of nodes from a larger graph, and it is in this context that the term "spanning" becomes a useful notion.

is_tree(G)	Returns True if G is a tree.
is_forest(G)	Returns True if G is a forest.
is_arborescence( $G$ )	Returns True if G is an arborescence.
is_branching( <b>G</b> )	Returns True if G is a branching.

### is\_tree

### $is\_tree(G)$

Returns True if G is a tree.

A tree is a connected graph with no undirected cycles.

For directed graphs, G is a tree if the underlying graph is a tree. The underlying graph is obtained by treating each directed edge as a single undirected edge in a multigraph.

Parameters G: graph

The graph to test.

Returns b: bool

A boolean that is True if G is a tree.

### See also:

is\_arborescence

#### **Notes**

In another convention, a directed tree is known as a polytree and then tree corresponds to an arborescence.

### is forest

### $is\_forest(G)$

Returns True if G is a forest.

A forest is a graph with no undirected cycles.

For directed graphs, G is a forest if the underlying graph is a forest. The underlying graph is obtained by treating each directed edge as a single undirected edge in a multigraph.

Parameters G: graph

The graph to test.

Returns b: bool

A boolean that is True if G is a forest.

#### See also:

is branching

#### **Notes**

In another convention, a directed forest is known as a polyforest and then forest corresponds to a branching.

### is\_arborescence

#### $is\_arborescence(G)$

Returns True if G is an arborescence.

An arborescence is a directed tree with maximum in-degree equal to 1.

Parameters G: graph

The graph to test.

Returns b : bool

A boolean that is True if G is an arborescence.

#### See also:

is\_tree

### Notes

In another convention, an arborescence is known as a tree.

### is\_branching

#### $is\_branching(G)$

Returns True if G is a branching.

A branching is a directed forest with maximum in-degree equal to 1.

Parameters G: directed graph

The directed graph to test.

4.38. Tree 343

#### Returns b: bool

A boolean that is True if G is a branching.

#### See also:

```
is_forest
```

#### **Notes**

In another convention, a branching is also known as a *forest*.

## 4.38.2 Branchings and Spanning Arborescences

Algorithms for finding optimum branchings and spanning arborescences.

This implementation is based on:

J. Edmonds, Optimum branchings, J. Res. Natl. Bur. Standards 71B (1967), 233–240. URL: http://archive.org/details/jresv71Bn4p233

branching_weight(G[, attr, default])	Returns the total weight of a branching.
<pre>greedy_branching(G[, attr, default, kind])</pre>	Returns a branching obtained through a greedy algorithm.
$maximum_branching(G[, attr, default])$	Returns a maximum branching from G.
minimum_branching(G[, attr, default])	Returns a minimum branching from G.
$\verb  maximum\_spanning\_arborescence  (G[, attr, default])  $	Returns a maximum spanning arborescence from G.
$\verb minimum_spanning_arborescence  (G[, attr, default]) $	Returns a minimum spanning arborescence from G.
Edmonds(G[, seed])	Edmonds algorithm for finding optimal branchings and spanning arbo

#### branching weight

```
branching_weight (G, attr='weight', default=1)
```

Returns the total weight of a branching.

#### greedy\_branching

```
greedy_branching (G, attr='weight', default=1, kind='max')
```

Returns a branching obtained through a greedy algorithm.

This algorithm is wrong, and cannot give a proper optimal branching. However, we include it for pedagogical reasons, as it can be helpful to see what its outputs are.

The output is a branching, and possibly, a spanning arborescence. However, it is not guaranteed to be optimal in either case.

#### Parameters G: DiGraph

The directed graph to scan.

attr : str

The attribute to use as weights. If None, then each edge will be treated equally with a weight of 1.

default: float

When attr is not None, then if an edge does not have that attribute, default specifies

```
what value it should take.
               kind: str
                   The type of optimum to search for: 'min' or 'max' greedy branching.
           Returns B: directed graph
                   The greedily obtained branching.
maximum branching
maximum_branching(G, attr='weight', default=1)
     Returns a maximum branching from G.
           Parameters G: (multi)digraph-like
                   The graph to be searched.
               attr: str
                   The edge attribute used to in determining optimality.
               default: float
                   The value of the edge attribute used if an edge does not have the attribute attr.
           Returns B: (multi)digraph-like
                   A maximum branching.
minimum branching
minimum branching (G, attr='weight', default=1)
     Returns a minimum branching from G.
           Parameters G: (multi)digraph-like
                   The graph to be searched.
               attr: str
                   The edge attribute used to in determining optimality.
               default: float
                   The value of the edge attribute used if an edge does not have the attribute attr.
           Returns B: (multi)digraph-like
                   A minimum branching.
maximum spanning arborescence
maximum\_spanning\_arborescence(G, attr='weight', default=1)
     Returns a maximum spanning arborescence from G.
           Parameters G: (multi)digraph-like
                   The graph to be searched.
               attr: str
```

4.38. Tree 345

The edge attribute used to in determining optimality.

default: float

The value of the edge attribute used if an edge does not have the attribute attr.

Returns B: (multi)digraph-like

A maximum spanning arborescence.

### Raises NetworkXException

If the graph does not contain a maximum spanning arborescence.

### minimum\_spanning\_arborescence

```
minimum_spanning_arborescence(G, attr='weight', default=1)
```

Returns a minimum spanning arborescence from G.

Parameters G: (multi)digraph-like

The graph to be searched.

attr: str

The edge attribute used to in determining optimality.

default: float

The value of the edge attribute used if an edge does not have the attribute attr.

Returns B: (multi)digraph-like

A minimum spanning arborescence.

### Raises NetworkXException

If the graph does not contain a minimum spanning arborescence.

#### **Edmonds**

```
{f class} \ {f Edmonds} \ (G, seed = None)
```

Edmonds algorithm for finding optimal branchings and spanning arborescences.

```
___init__(G, seed=None)
```

#### **Methods**

```
__init__(G[, seed])
find_optimum([attr, default, kind, style]) Returns a branching from G.
```

# 4.39 Vitality

Vitality measures.

 $closeness\_vitality(G[, weight])$  Compute closeness vitality for nodes.

## 4.39.1 closeness\_vitality

### closeness\_vitality(G, weight=None)

Compute closeness vitality for nodes.

Closeness vitality of a node is the change in the sum of distances between all node pairs when excluding that node.

Parameters G: graph

weight: None or string (optional)

The name of the edge attribute used as weight. If None the edge weights are ignored.

Returns nodes: dictionary

Dictionary with nodes as keys and closeness vitality as the value.

#### See also:

```
closeness_centrality
```

#### References

[R317]

### **Examples**

```
>>> G=nx.cycle_graph(3)
>>> nx.closeness_vitality(G)
{0: 4.0, 1: 4.0, 2: 4.0}
```

4.39. Vitality 347

**CHAPTER** 

**FIVE** 

## **FUNCTIONS**

Functional interface to graph methods and assorted utilities.

# 5.1 Graph

degree(G[, nbunch, weight])	Return degree of single node or of nbunch of nodes.
$ ext{degree\_histogram}(G)$	Return a list of the frequency of each degree value.
density(G)	Return the density of a graph.
info(G[, n])	Print short summary of information for the graph G or the node n.
<pre>create_empty_copy(G[, with_nodes])</pre>	Return a copy of the graph G with all of the edges removed.
is_directed(G)	Return True if graph is directed.

## **5.1.1 degree**

degree (G, nbunch=None, weight=None)

Return degree of single node or of nbunch of nodes. If nbunch is ommitted, then return degrees of *all* nodes.

# 5.1.2 degree\_histogram

### ${\tt degree\_histogram}\,(G)$

Return a list of the frequency of each degree value.

Parameters G: Networkx graph

A graph

Returns hist: list

A list of frequencies of degrees. The degree values are the index in the list.

### **Notes**

Note: the bins are width one, hence len(list) can be large (Order(number\_of\_edges))

# 5.1.3 density

### $\mathtt{density}\,(G)$

Return the density of a graph.

The density for undirected graphs is

$$d = \frac{2m}{n(n-1)},$$

and for directed graphs is

$$d = \frac{m}{n(n-1)},$$

where n is the number of nodes and m is the number of edges in G.

#### **Notes**

The density is 0 for a graph without edges and 1 for a complete graph. The density of multigraphs can be higher than 1.

Self loops are counted in the total number of edges so graphs with self loops can have density higher than 1.

### 5.1.4 info

info(G, n=None)

Print short summary of information for the graph G or the node n.

Parameters G: Networkx graph

A graph

n: node (any hashable)

A node in the graph G

## 5.1.5 create\_empty\_copy

create\_empty\_copy(G, with\_nodes=True)

Return a copy of the graph G with all of the edges removed.

Parameters G: graph

A NetworkX graph

with\_nodes : bool (default=True)

Include nodes.

#### Notes

Graph, node, and edge data is not propagated to the new graph.

## 5.1.6 is directed

### $is\_directed(G)$

Return True if graph is directed.

# 5.2 Nodes

5.2. Nodes 351

nodes(G)	Return a copy of the graph nodes in a list.
$number\_of\_nodes(G)$	Return the number of nodes in the graph.
$nodes\_iter(G)$	Return an iterator over the graph nodes.
all_neighbors(graph, node)	Returns all of the neighbors of a node in the graph.
non_neighbors(graph, node)	Returns the non-neighbors of the node in the graph.
$common_neighbors(G, u, v)$	Return the common neighbors of two nodes in a graph.

### 5.2.1 nodes

### nodes(G)

Return a copy of the graph nodes in a list.

### 5.2.2 number of nodes

#### $number_of_nodes(G)$

Return the number of nodes in the graph.

## 5.2.3 nodes iter

#### $nodes\_iter(G)$

Return an iterator over the graph nodes.

## 5.2.4 all\_neighbors

### all\_neighbors (graph, node)

Returns all of the neighbors of a node in the graph.

If the graph is directed returns predecessors as well as successors.

### Parameters graph: NetworkX graph

Graph to find neighbors.

node: node

The node whose neighbors will be returned.

Returns neighbors: iterator

Iterator of neighbors

## 5.2.5 non\_neighbors

### non\_neighbors (graph, node)

Returns the non-neighbors of the node in the graph.

Parameters graph: NetworkX graph

Graph to find neighbors.

node: node

The node whose neighbors will be returned.

### Returns non\_neighbors: iterator

Iterator of nodes in the graph that are not neighbors of the node.

## 5.2.6 common\_neighbors

```
common neighbors (G, u, v)
```

Return the common neighbors of two nodes in a graph.

Parameters G: graph

A NetworkX undirected graph.

**u**, **v** : nodes

Nodes in the graph.

Returns cnbors: iterator

Iterator of common neighbors of u and v in the graph.

#### Raises NetworkXError

If u or v is not a node in the graph.

### **Examples**

```
>>> G = nx.complete_graph(5)
>>> sorted(nx.common_neighbors(G, 0, 1))
[2, 3, 4]
```

# 5.3 Edges

edges(G[, nbunch])	Return list of edges incident to nodes in nbunch.
$number_of_edges(G)$	Return the number of edges in the graph.
edges_iter(G[, nbunch])	Return iterator over edges incident to nodes in nbunch.
non_edges(graph)	Returns the non-existent edges in the graph.

## 5.3.1 edges

```
edges (G, nbunch=None)
```

Return list of edges incident to nodes in nbunch.

Return all edges if nbunch is unspecified or nbunch=None.

For digraphs, edges=out\_edges

## 5.3.2 number\_of\_edges

### ${\tt number\_of\_edges}\,(G)$

Return the number of edges in the graph.

5.3. Edges 353

### 5.3.3 edges iter

```
edges_iter(G, nbunch=None)
```

Return iterator over edges incident to nodes in nbunch.

Return all edges if nbunch is unspecified or nbunch=None.

For digraphs, edges=out\_edges

## 5.3.4 non\_edges

```
non_edges (graph)
```

Returns the non-existent edges in the graph.

Parameters graph: NetworkX graph.

Graph to find non-existent edges.

Returns non\_edges: iterator

Iterator of edges that are not in the graph.

## 5.4 Attributes

set_node_attributes(G, name, values)	Set node attributes from dictionary of nodes and values
<pre>get_node_attributes(G, name)</pre>	Get node attributes from graph
set_edge_attributes(G, name, values)	Set edge attributes from dictionary of edge tuples and values.
<pre>get_edge_attributes(G, name)</pre>	Get edge attributes from graph

### 5.4.1 set node attributes

```
set_node_attributes(G, name, values)
```

Set node attributes from dictionary of nodes and values

**Parameters G**: NetworkX Graph

**name**: string
Attribute name

values: dict

Dictionary of attribute values keyed by node. If values is not a dictionary, then it is treated as a single attribute value that is then applied to every node in G.

### **Examples**

```
>>> G = nx.path_graph(3)
>>> bb = nx.betweenness_centrality(G)
>>> nx.set_node_attributes(G, 'betweenness', bb)
>>> G.node[1]['betweenness']
1.0
```

## 5.4.2 get node attributes

### get\_node\_attributes (G, name)

Get node attributes from graph

Parameters G: NetworkX Graph

name: string

Attribute name

**Returns** Dictionary of attributes keyed by node.

## **Examples**

```
>>> G=nx.Graph()
>>> G.add_nodes_from([1,2,3],color='red')
>>> color=nx.get_node_attributes(G,'color')
>>> color[1]
'red'
```

## 5.4.3 set\_edge\_attributes

### set\_edge\_attributes (G, name, values)

Set edge attributes from dictionary of edge tuples and values.

Parameters G: NetworkX Graph

name: string

Attribute name

values : dict

Dictionary of attribute values keyed by edge (tuple). For multigraphs, the keys tuples must be of the form (u, v, key). For non-multigraphs, the keys must be tuples of the form (u, v). If values is not a dictionary, then it is treated as a single attribute value that is then applied to every edge in G.

### **Examples**

```
>>> G = nx.path_graph(3)
>>> bb = nx.edge_betweenness_centrality(G, normalized=False)
>>> nx.set_edge_attributes(G, 'betweenness', bb)
>>> G[1][2]['betweenness']
2.0
```

## 5.4.4 get edge attributes

## get\_edge\_attributes(G, name)

Get edge attributes from graph

**Parameters G**: NetworkX Graph

name: string

5.4. Attributes 355

Attribute name

**Returns** Dictionary of attributes keyed by edge. For (di)graphs, the keys are 2-tuples of the form: (u,v). For multi(di)graphs, the keys are 3-tuples of the form: (u, v, key).

### **Examples**

```
>>> G=nx.Graph()
>>> G.add_path([1,2,3],color='red')
>>> color=nx.get_edge_attributes(G,'color')
>>> color[(1,2)]
'red'
```

# 5.5 Freezing graph structure

freeze(G)	Modify graph to prevent further change by adding or removing nodes or edges.
is_frozen(G)	Return True if graph is frozen.

### **5.5.1** freeze

### freeze(G)

Modify graph to prevent further change by adding or removing nodes or edges.

Node and edge data can still be modified.

**Parameters G**: graph

A NetworkX graph

#### See also:

```
is frozen
```

### **Notes**

To "unfreeze" a graph you must make a copy by creating a new graph object:

```
>>> graph = nx.path_graph(4)
>>> frozen_graph = nx.freeze(graph)
>>> unfrozen_graph = nx.Graph(frozen_graph)
>>> nx.is_frozen(unfrozen_graph)
False
```

### **Examples**

```
>>> G=nx.Graph()
>>> G.add_path([0,1,2,3])
>>> G=nx.freeze(G)
>>> try:
... G.add_edge(4,5)
```

```
... except nx.NetworkXError as e:
... print(str(e))
Frozen graph can't be modified
```

# 5.5.2 is\_frozen

## $is\_frozen(G)$

Return True if graph is frozen.

 $Parameters \ G: graph$ 

A NetworkX graph

See also:

freeze

**CHAPTER** 

SIX

## **GRAPH GENERATORS**

## 6.1 Atlas

Generators for the small graph atlas.

See "An Atlas of Graphs" by Ronald C. Read and Robin J. Wilson, Oxford University Press, 1998.

Because of its size, this module is not imported by default.

graph\_atlas\_g() Return the list [G0,G1,...,G1252] of graphs as named in the Graph Atlas.

## 6.1.1 graph\_atlas\_g

### graph\_atlas\_g()

Return the list [G0,G1,...,G1252] of graphs as named in the Graph Atlas. G0,G1,...,G1252 are all graphs with up to 7 nodes.

## The graphs are listed:

- 1. in increasing order of number of nodes;
- 2. for a fixed number of nodes, in increasing order of the number of edges;
- 3. for fixed numbers of nodes and edges, in increasing order of the degree sequence, for example 111223 < 112222;
- 4. for fixed degree sequence, in increasing number of automorphisms.

Note that indexing is set up so that for GAG=graph\_atlas\_g(), then G123=GAG[123] and G[0]=empty\_graph(0)

## 6.2 Classic

Generators for some classic graphs.

The typical graph generator is called as follows:

```
>>> G=nx.complete_graph(100)
```

returning the complete graph on n nodes labeled 0,...,99 as a simple graph. Except for empty\_graph, all the generators in this module return a Graph class (i.e. a simple, undirected graph).

balanced\_tree(r, h[, create\_using])

Return the perfectly balanced r-tree of height h.

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<pre>barbell_graph(m1, m2[, create_using])</pre>	Return the Barbell Graph: two complete graphs connected by a path.
<pre>complete_graph(n[, create_using])</pre>	Return the complete graph K_n with n nodes.
circular_ladder_graph(n[, create_using])	Return the circular ladder graph CL_n of length n.
cycle_graph(n[, create_using])	Return the cycle graph C_n over n nodes.
${\tt dorogovtsev\_goltsev\_mendes\_graph(n[,])}$	Return the hierarchically constructed Dorogovtsev-Goltsev-Mendes graph.
<pre>empty_graph([n, create_using])</pre>	Return the empty graph with n nodes and zero edges.
<pre>grid_2d_graph(m, n[, periodic, create_using])</pre>	Return the 2d grid graph of mxn nodes, each connected to its nearest neight
grid_graph(dim[, periodic])	Return the n-dimensional grid graph.
hypercube_graph(n)	Return the n-dimensional hypercube.
hypercube_graph(n) ladder_graph(n[, create_using])	Return the Ladder graph of length n.
	Return the Ladder graph of length n.  Return the Lollipop Graph; $K_m$ connected to $P_n$ .
<pre>ladder_graph(n[, create_using]) lollipop_graph(m, n[, create_using]) null_graph([create_using])</pre>	Return the Ladder graph of length n.  Return the Lollipop Graph; $K_m$ connected to $P_n$ .  Return the Null graph with no nodes or edges.
<pre>ladder_graph(n[, create_using]) lollipop_graph(m, n[, create_using])</pre>	Return the Ladder graph of length n.  Return the Lollipop Graph; $K_m$ connected to $P_n$ .  Return the Null graph with no nodes or edges.  Return the Path graph P_n of n nodes linearly connected by n-1 edges.
<pre>ladder_graph(n[, create_using]) lollipop_graph(m, n[, create_using]) null_graph([create_using]) path_graph(n[, create_using]) star_graph(n[, create_using])</pre>	Return the Ladder graph of length n.  Return the Lollipop Graph; $K_m$ connected to $P_n$ .  Return the Null graph with no nodes or edges.  Return the Path graph P_n of n nodes linearly connected by n-1 edges.  Return the Star graph with n+1 nodes: one center node, connected to n outer.
<pre>ladder_graph(n[, create_using]) lollipop_graph(m, n[, create_using]) null_graph([create_using]) path_graph(n[, create_using])</pre>	Return the Ladder graph of length n.  Return the Lollipop Graph; $K_m$ connected to $P_n$ .  Return the Null graph with no nodes or edges.  Return the Path graph P_n of n nodes linearly connected by n-1 edges.

## 6.2.1 balanced\_tree

balanced\_tree (r, h, create\_using=None)

Return the perfectly balanced r-tree of height h.

Parameters r: int

Branching factor of the tree

h: int

Height of the tree

create\_using : NetworkX graph type, optional

Use specified type to construct graph (default = networkx.Graph)

**Returns** G: networkx Graph

A tree with n nodes

### **Notes**

This is the rooted tree where all leaves are at distance h from the root. The root has degree r and all other internal nodes have degree r+1.

Node labels are the integers 0 (the root) up to number\_of\_nodes - 1.

Also refered to as a complete r-ary tree.

## 6.2.2 barbell graph

barbell\_graph (m1, m2, create\_using=None)

Return the Barbell Graph: two complete graphs connected by a path.

For m1 > 1 and m2 >= 0.

Two identical complete graphs  $K_{m1}$  form the left and right bells, and are connected by a path  $P_{m2}$ .

**The 2\*m1+m2 nodes are numbered** 0,...,m1-1 for the left barbell, m1,...,m1+m2-1 for the path, and m1+m2,...,2\*m1+m2-1 for the right barbell.

The 3 subgraphs are joined via the edges (m1-1,m1) and (m1+m2-1,m1+m2). If m2=0, this is merely two complete graphs joined together.

This graph is an extremal example in David Aldous and Jim Fill's etext on Random Walks on Graphs.

## 6.2.3 complete graph

### complete\_graph (n, create\_using=None)

Return the complete graph K\_n with n nodes.

Node labels are the integers 0 to n-1.

## 6.2.4 circular\_ladder\_graph

### circular\_ladder\_graph (n, create\_using=None)

Return the circular ladder graph CL\_n of length n.

CL\_n consists of two concentric n-cycles in which each of the n pairs of concentric nodes are joined by an edge.

Node labels are the integers 0 to n-1

## 6.2.5 cycle graph

### cycle\_graph (n, create\_using=None)

Return the cycle graph C\_n over n nodes.

C\_n is the n-path with two end-nodes connected.

Node labels are the integers 0 to n-1 If create\_using is a DiGraph, the direction is in increasing order.

## 6.2.6 dorogovtsev\_goltsev\_mendes\_graph

### dorogovtsev\_goltsev\_mendes\_graph (n, create\_using=None)

Return the hierarchically constructed Dorogovtsev-Goltsev-Mendes graph.

n is the generation. See: arXiv:/cond-mat/0112143 by Dorogovtsev, Goltsev and Mendes.

## 6.2.7 empty\_graph

#### empty\_graph (n=0, create\_using=None)

Return the empty graph with n nodes and zero edges.

Node labels are the integers 0 to n-1

For example: >>> G=nx.empty\_graph(10) >>> G.number\_of\_nodes() 10 >>> G.number\_of\_edges() 0

The variable create\_using should point to a "graph"-like object that will be cleaned (nodes and edges will be removed) and refitted as an empty "graph" with n nodes with integer labels. This capability is useful for specifying the class-nature of the resulting empty "graph" (i.e. Graph, DiGraph, MyWeirdGraphClass, etc.).

The variable create\_using has two main uses: Firstly, the variable create\_using can be used to create an empty digraph, network,etc. For example,

6.2. Classic 361

```
>>> n=10
>>> G=nx.empty_graph(n,create_using=nx.DiGraph())
```

will create an empty digraph on n nodes.

Secondly, one can pass an existing graph (digraph, pseudograph, etc.) via create\_using. For example, if G is an existing graph (resp. digraph, pseudograph, etc.), then empty\_graph(n,create\_using=G) will empty G (i.e. delete all nodes and edges using G.clear() in base) and then add n nodes and zero edges, and return the modified graph (resp. digraph, pseudograph, etc.).

See also create\_empty\_copy(G).

## 6.2.8 grid 2d graph

### grid\_2d\_graph (m, n, periodic=False, create\_using=None)

Return the 2d grid graph of mxn nodes, each connected to its nearest neighbors. Optional argument periodic=True will connect boundary nodes via periodic boundary conditions.

## 6.2.9 grid\_graph

### grid\_graph (dim, periodic=False)

Return the n-dimensional grid graph.

The dimension is the length of the list 'dim' and the size in each dimension is the value of the list element.

E.g. G=grid\_graph(dim=[2,3]) produces a 2x3 grid graph.

If periodic=True then join grid edges with periodic boundary conditions.

## 6.2.10 hypercube\_graph

### $hypercube_graph(n)$

Return the n-dimensional hypercube.

Node labels are the integers 0 to 2\*\*n - 1.

## 6.2.11 ladder graph

### ladder\_graph (n, create\_using=None)

Return the Ladder graph of length n.

This is two rows of n nodes, with each pair connected by a single edge.

Node labels are the integers 0 to 2\*n - 1.

## 6.2.12 lollipop graph

### lollipop\_graph (m, n, create\_using=None)

Return the Lollipop Graph;  $K_m$  connected to  $P_n$ .

This is the Barbell Graph without the right barbell.

For m>1 and n>=0, the complete graph  $K_m$  is connected to the path  $P_n$ . The resulting m+n nodes are labelled 0,...,m-1 for the complete graph and m,...,m+n-1 for the path. The 2 subgraphs are joined via the edge (m-1,m). If n=0, this is merely a complete graph.

Node labels are the integers 0 to number\_of\_nodes - 1.

(This graph is an extremal example in David Aldous and Jim Fill's etext on Random Walks on Graphs.)

## 6.2.13 null\_graph

### null graph(create using=None)

Return the Null graph with no nodes or edges.

See empty\_graph for the use of create\_using.

## 6.2.14 path\_graph

### path\_graph (n, create\_using=None)

Return the Path graph P\_n of n nodes linearly connected by n-1 edges.

Node labels are the integers 0 to n - 1. If create\_using is a DiGraph then the edges are directed in increasing order.

## 6.2.15 star\_graph

### star\_graph (n, create\_using=None)

Return the Star graph with n+1 nodes: one center node, connected to n outer nodes.

Node labels are the integers 0 to n.

# 6.2.16 trivial\_graph

### trivial\_graph (create\_using=None)

Return the Trivial graph with one node (with integer label 0) and no edges.

## 6.2.17 wheel\_graph

#### wheel graph (n, create using=None)

Return the wheel graph: a single hub node connected to each node of the (n-1)-node cycle graph.

Node labels are the integers 0 to n - 1.

# 6.3 Expanders

Provides explicit constructions of expander graphs.

margulis\_gabber\_galil\_graph(n[, create\_using]) Return the Margulis-Gabber-Galil undirected MultiGraph on  $n^2$  nodes. chordal\_cycle\_graph(p[, create\_using]) Return the chordal cycle graph on p nodes.

## 6.4 Small

Various small and named graphs, together with some compact generators.

6.4. Small 363

make_small_graph(graph_description[,])	Return the small graph described by graph_description.
LCF_graph(n, shift_list, repeats[, create_using])	Return the cubic graph specified in LCF notation.
bull_graph([create_using])	Return the Bull graph.
chvatal_graph([create_using])	Return the Chvátal graph.
cubical_graph([create_using])	Return the 3-regular Platonic Cubical graph.
desargues_graph([create_using])	Return the Desargues graph.
diamond_graph([create_using])	Return the Diamond graph.
dodecahedral_graph([create_using])	Return the Platonic Dodecahedral graph.
<pre>frucht_graph([create_using])</pre>	Return the Frucht Graph.
heawood_graph([create_using])	Return the Heawood graph, a (3,6) cage.
house_graph([create_using])	Return the House graph (square with triangle on top).
house_x_graph([create_using])	Return the House graph with a cross inside the house square.
<pre>icosahedral_graph([create_using])</pre>	Return the Platonic Icosahedral graph.
<pre>krackhardt_kite_graph([create_using])</pre>	Return the Krackhardt Kite Social Network.
<pre>moebius_kantor_graph([create_using])</pre>	Return the Moebius-Kantor graph.
octahedral_graph([create_using])	Return the Platonic Octahedral graph.
pappus_graph()	Return the Pappus graph.
<pre>petersen_graph([create_using])</pre>	Return the Petersen graph.
sedgewick_maze_graph([create_using])	Return a small maze with a cycle.
tetrahedral_graph([create_using])	Return the 3-regular Platonic Tetrahedral graph.
truncated_cube_graph([create_using])	Return the skeleton of the truncated cube.
<pre>truncated_tetrahedron_graph([create_using])</pre>	Return the skeleton of the truncated Platonic tetrahedron.
tutte_graph([create_using])	Return the Tutte graph.

## 6.4.1 make\_small\_graph

 $\verb|make_small_graph| (graph_description, create\_using=None)|$ 

Return the small graph described by graph\_description.

graph\_description is a list of the form [ltype,name,n,xlist]

Here ltype is one of "adjacencylist" or "edgelist", name is the name of the graph and n the number of nodes. This constructs a graph of n nodes with integer labels 0,..,n-1.

If ltype="adjacencylist" then xlist is an adjacency list with exactly n entries, in with the j'th entry (which can be empty) specifies the nodes connected to vertex j. e.g. the "square" graph C\_4 can be obtained by

```
>>> G=nx.make_small_graph(["adjacencylist", "C_4", 4, [[2,4],[1,3],[2,4],[1,3]]])
```

or, since we do not need to add edges twice,

```
>>> G=nx.make_small_graph(["adjacencylist", "C_4", 4, [[2, 4], [3], [4], []]])
```

If ltype="edgelist" then xlist is an edge list written as [[v1,w2],[v2,w2],...,[vk,wk]], where vj and wj integers in the range 1,..,n e.g. the "square" graph C\_4 can be obtained by

```
>>> G=nx.make_small_graph(["edgelist", "C_4", 4, [[1,2], [3,4], [2,3], [4,1]]])
```

Use the create\_using argument to choose the graph class/type.

## 6.4.2 LCF\_graph

**LCF\_graph** (*n*, *shift\_list*, *repeats*, *create\_using=None*) Return the cubic graph specified in LCF notation.

LCF notation (LCF=Lederberg-Coxeter-Fruchte) is a compressed notation used in the generation of various cubic Hamiltonian graphs of high symmetry. See, for example, dodecahedral\_graph, desargues\_graph, hea-wood\_graph and pappus\_graph below.

**n (number of nodes)** The starting graph is the n-cycle with nodes 0,...,n-1. (The null graph is returned if n < 0.)

```
shift list = [s1,s2,...,sk], a list of integer shifts mod n,
```

**repeats** integer specifying the number of times that shifts in shift\_list are successively applied to each v\_current in the n-cycle to generate an edge between v\_current and v\_current+shift mod n.

For v1 cycling through the n-cycle a total of k\*repeats with shift cycling through shiftlist repeats times connect v1 with v1+shift mod n

```
The utility graph K_{3,3}

>>> G=nx.LCF_graph(6,[3,-3],3)

The Heawood graph

>>> G=nx.LCF_graph(14,[5,-5],7)
```

See http://mathworld.wolfram.com/LCFNotation.html for a description and references.

## 6.4.3 bull graph

```
bull_graph (create_using=None)
Return the Bull graph.
```

# 6.4.4 chvatal\_graph

```
chvatal_graph (create_using=None)
Return the Chvátal graph.
```

## 6.4.5 cubical\_graph

```
cubical_graph (create_using=None)
Return the 3-regular Platonic Cubical graph.
```

# 6.4.6 desargues\_graph

```
desargues_graph (create_using=None)
Return the Desargues graph.
```

# 6.4.7 diamond\_graph

```
diamond_graph (create_using=None)
Return the Diamond graph.
```

6.4. Small 365

## 6.4.8 dodecahedral\_graph

### dodecahedral\_graph (create\_using=None)

Return the Platonic Dodecahedral graph.

## 6.4.9 frucht graph

### frucht\_graph (create\_using=None)

Return the Frucht Graph.

The Frucht Graph is the smallest cubical graph whose automorphism group consists only of the identity element.

## 6.4.10 heawood\_graph

### heawood\_graph (create\_using=None)

Return the Heawood graph, a (3,6) cage.

## 6.4.11 house graph

### house\_graph (create\_using=None)

Return the House graph (square with triangle on top).

## 6.4.12 house\_x\_graph

### house\_x\_graph (create\_using=None)

Return the House graph with a cross inside the house square.

## 6.4.13 icosahedral\_graph

### icosahedral\_graph (create\_using=None)

Return the Platonic Icosahedral graph.

## 6.4.14 krackhardt\_kite\_graph

### krackhardt\_kite\_graph (create\_using=None)

Return the Krackhardt Kite Social Network.

A 10 actor social network introduced by David Krackhardt to illustrate: degree, betweenness, centrality, closeness, etc. The traditional labeling is: Andre=1, Beverley=2, Carol=3, Diane=4, Ed=5, Fernando=6, Garth=7, Heather=8, Ike=9, Jane=10.

## 6.4.15 moebius kantor graph

## moebius\_kantor\_graph(create\_using=None)

Return the Moebius-Kantor graph.

## 6.4.16 octahedral\_graph

octahedral\_graph (create\_using=None)
Return the Platonic Octahedral graph.

## 6.4.17 pappus\_graph

pappus\_graph ()
 Return the Pappus graph.

## 6.4.18 petersen graph

petersen\_graph (create\_using=None)
Return the Petersen graph.

## 6.4.19 sedgewick maze graph

sedgewick\_maze\_graph (create\_using=None)

Return a small maze with a cycle.

This is the maze used in Sedgewick,3rd Edition, Part 5, Graph Algorithms, Chapter 18, e.g. Figure 18.2 and following. Nodes are numbered 0,...,7

## 6.4.20 tetrahedral\_graph

tetrahedral\_graph (create\_using=None)
Return the 3-regular Platonic Tetrahedral graph.

## 6.4.21 truncated cube graph

truncated\_cube\_graph (create\_using=None)
Return the skeleton of the truncated cube.

## 6.4.22 truncated tetrahedron graph

truncated\_tetrahedron\_graph (create\_using=None)
Return the skeleton of the truncated Platonic tetrahedron.

## 6.4.23 tutte graph

**tutte\_graph** (*create\_using=None*)
Return the Tutte graph.

# 6.5 Random Graphs

Generators for random graphs.

<pre>fast_gnp_random_graph(n, p[, seed, directed])</pre>	Return a random graph G <sub>{n,p}</sub> (Erdős-Rényi graph, binomial graph).
<pre>gnp_random_graph(n, p[, seed, directed])</pre>	Return a random graph $G_{n,p}$ (Erdős-Rényi graph, binomial graph).
${\tt dense\_gnm\_random\_graph(n,m[,seed])}$	Return the random graph $G_{n,m}$ .
<pre>gnm_random_graph(n, m[, seed, directed])</pre>	Return the random graph $G_{n,m}$ .
<pre>erdos_renyi_graph(n, p[, seed, directed])</pre>	Return a random graph $G_{n,p}$ (Erdős-Rényi graph, binomial graph).
<pre>binomial_graph(n, p[, seed, directed])</pre>	Return a random graph $G_{n,p}$ (Erdős-Rényi graph, binomial graph).
$newman_watts_strogatz_graph(n, k, p[, seed])$	Return a Newman-Watts-Strogatz small world graph.
watts_strogatz_graph(n, k, p[, seed])	Return a Watts-Strogatz small-world graph.
$\verb connected_watts_strogatz_graph (n,k,p[,]) $	Return a connected Watts-Strogatz small-world graph.
$random\_regular\_graph(d, n[, seed])$	Return a random regular graph of n nodes each with degree d.
barabasi_albert_graph(n, m[, seed])	Return random graph using Barabási-Albert preferential attachment mo
<pre>powerlaw_cluster_graph(n, m, p[, seed])</pre>	Holme and Kim algorithm for growing graphs with powerlaw degree di
random_lobster(n, p1, p2[, seed])	Return a random lobster.
random_shell_graph(constructor[, seed])	Return a random shell graph for the constructor given.
<pre>random_powerlaw_tree(n[, gamma, seed, tries])</pre>	Return a tree with a powerlaw degree distribution.
random_powerlaw_tree_sequence(n[, gamma,])	Return a degree sequence for a tree with a powerlaw distribution.

## 6.5.1 fast gnp random graph

**fast\_gnp\_random\_graph** (*n*, *p*, *seed=None*, *directed=False*) Return a random graph G\_{n,p} (Erdős-Rényi graph, binomial graph).

Parameters n: int

The number of nodes.

**p**: float

Probability for edge creation.

seed: int, optional

Seed for random number generator (default=None).

directed : bool, optional (default=False)

If True return a directed graph

### See also:

gnp\_random\_graph

### **Notes**

The  $G_{n,p}$  graph algorithm chooses each of the [n(n-1)]/2 (undirected) or n(n-1) (directed) possible edges with probability p.

This algorithm is O(n+m) where m is the expected number of edges m=p\*n\*(n-1)/2.

It should be faster than gnp\_random\_graph when p is small and the expected number of edges is small (sparse graph).

### References

[R349]

## 6.5.2 gnp random graph

```
gnp_random_graph (n, p, seed=None, directed=False)
    Return a random graph G_{n,p} (Erdős-Rényi graph, binomial graph).
    Chooses each of the possible edges with probability p.
    This is also called binomial_graph and erdos_renyi_graph.
    Parameters n: int
        The number of nodes.
    p: float
        Probability for edge creation.
    seed: int, optional
        Seed for random number generator (default=None).
    directed: bool, optional (default=False)
        If True return a directed graph
    See also:
    fast_gnp_random_graph
    Notes
```

This is an O(n^2) algorithm. For sparse graphs (small p) see fast\_gnp\_random\_graph for a faster algorithm.

### References

[R350], [R351]

## 6.5.3 dense\_gnm\_random\_graph

```
dense_gnm_random_graph (n, m, seed=None)
Return the random graph G_{n,m}.
```

Gives a graph picked randomly out of the set of all graphs with n nodes and m edges. This algorithm should be faster than gnm\_random\_graph for dense graphs.

```
Parameters n: int

The number of nodes.

m: int

The number of edges.

seed: int, optional

Seed for random number generator (default=None).

See also:

gnm_random_graph
```

#### **Notes**

Algorithm by Keith M. Briggs Mar 31, 2006. Inspired by Knuth's Algorithm S (Selection sampling technique), in section 3.4.2 of [R346].

#### References

[R346]

## 6.5.4 gnm random graph

```
gnm\_random\_graph(n, m, seed=None, directed=False)
```

Return the random graph  $G_{n,m}$ .

Produces a graph picked randomly out of the set of all graphs with n nodes and m edges.

```
Parameters n: int
```

The number of nodes.

m: int

The number of edges.

seed: int, optional

Seed for random number generator (default=None).

directed : bool, optional (default=False)

If True return a directed graph

# 6.5.5 erdos\_renyi\_graph

```
erdos_renyi_graph(n, p, seed=None, directed=False)
```

Return a random graph G\_{n,p} (Erdős-Rényi graph, binomial graph).

Chooses each of the possible edges with probability p.

This is also called binomial\_graph and erdos\_renyi\_graph.

### Parameters n: int

The number of nodes.

p: float

Probability for edge creation.

seed: int, optional

Seed for random number generator (default=None).

**directed**: bool, optional (default=False)

If True return a directed graph

## See also:

```
fast_gnp_random_graph
```

#### **Notes**

This is an O(n^2) algorithm. For sparse graphs (small p) see fast\_gnp\_random\_graph for a faster algorithm.

### References

[R347], [R348]

## 6.5.6 binomial graph

```
binomial_graph (n, p, seed=None, directed=False)
```

Return a random graph G\_{n,p} (Erdős-Rényi graph, binomial graph).

Chooses each of the possible edges with probability p.

This is also called binomial\_graph and erdos\_renyi\_graph.

```
Parameters n: int
```

The number of nodes.

**p**: float

Probability for edge creation.

seed: int, optional

Seed for random number generator (default=None).

directed: bool, optional (default=False)

If True return a directed graph

#### See also:

```
fast_gnp_random_graph
```

## **Notes**

This is an O(n^2) algorithm. For sparse graphs (small p) see fast\_gnp\_random\_graph for a faster algorithm.

## References

[R344], [R345]

## 6.5.7 newman watts strogatz graph

```
\verb"newman_watts_strogatz_graph" (n, k, p, seed=None)
```

Return a Newman-Watts-Strogatz small world graph.

Parameters n: int

The number of nodes

 $\mathbf{k}$ : int

Each node is connected to k nearest neighbors in ring topology

**p**: float

The probability of adding a new edge for each edge

seed: int, optional

seed for random number generator (default=None)

#### See also:

```
watts strogatz graph
```

#### **Notes**

First create a ring over n nodes. Then each node in the ring is connected with its k nearest neighbors (k-1 neighbors if k is odd). Then shortcuts are created by adding new edges as follows: for each edge u-v in the underlying "n-ring with k nearest neighbors" with probability p add a new edge u-w with randomly-chosen existing node w. In contrast with watts\_strogatz\_graph(), no edges are removed.

#### References

[R352]

## 6.5.8 watts strogatz graph

```
watts\_strogatz\_graph(n, k, p, seed=None)
```

Return a Watts-Strogatz small-world graph.

Parameters n: int

The number of nodes

k: int

Each node is connected to k nearest neighbors in ring topology

**p**: float

The probability of rewiring each edge

seed: int, optional

Seed for random number generator (default=None)

## See also:

```
newman_watts_strogatz_graph, connected_watts_strogatz_graph
```

### **Notes**

First create a ring over n nodes. Then each node in the ring is connected with its k nearest neighbors (k-1 neighbors if k is odd). Then shortcuts are created by replacing some edges as follows: for each edge u-v in the underlying "n-ring with k nearest neighbors" with probability p replace it with a new edge u-w with uniformly random choice of existing node w.

In contrast with newman\_watts\_strogatz\_graph(), the random rewiring does not increase the number of edges. The rewired graph is not guaranteed to be connected as in connected\_watts\_strogatz\_graph().

#### References

[R356]

## 6.5.9 connected watts strogatz graph

```
connected_watts_strogatz_graph (n, k, p, tries=100, seed=None)
```

Return a connected Watts-Strogatz small-world graph.

Attempt to generate a connected realization by repeated generation of Watts-Strogatz small-world graphs. An exception is raised if the maximum number of tries is exceeded.

#### Parameters n: int

The number of nodes

k: int

Each node is connected to k nearest neighbors in ring topology

p: float

The probability of rewiring each edge

tries: int

Number of attempts to generate a connected graph.

seed: int, optional

The seed for random number generator.

### See also:

```
newman_watts_strogatz_graph, watts_strogatz_graph
```

## 6.5.10 random\_regular\_graph

```
random_regular_graph (d, n, seed=None)
```

Return a random regular graph of n nodes each with degree d.

The resulting graph G has no self-loops or parallel edges.

### Parameters d: int

Degree

n: integer

Number of nodes. The value of n\*d must be even.

seed: hashable object

The seed for random number generator.

### Notes

The nodes are numbered form 0 to n-1.

Kim and Vu's paper [R355] shows that this algorithm samples in an asymptotically uniform way from the space of random graphs when  $d = O(n^{**}(1/3-epsilon))$ .

#### References

[R354], [R355]

## 6.5.11 barabasi\_albert\_graph

## barabasi\_albert\_graph (n, m, seed=None)

Return random graph using Barabási-Albert preferential attachment model.

A graph of n nodes is grown by attaching new nodes each with m edges that are preferentially attached to existing nodes with high degree.

#### Parameters n: int

Number of nodes

m: int

Number of edges to attach from a new node to existing nodes

seed: int, optional

Seed for random number generator (default=None).

#### **Returns G**: Graph

### **Notes**

The initialization is a graph with with m nodes and no edges.

#### References

[R343]

## 6.5.12 powerlaw\_cluster\_graph

## powerlaw\_cluster\_graph (n, m, p, seed=None)

Holme and Kim algorithm for growing graphs with powerlaw degree distribution and approximate average clustering.

### Parameters n: int

the number of nodes

m: int

the number of random edges to add for each new node

p: float,

Probability of adding a triangle after adding a random edge

seed: int, optional

Seed for random number generator (default=None).

#### **Notes**

The average clustering has a hard time getting above a certain cutoff that depends on m. This cutoff is often quite low. Note that the transitivity (fraction of triangles to possible triangles) seems to go down with network size.

It is essentially the Barabási-Albert (B-A) growth model with an extra step that each random edge is followed by a chance of making an edge to one of its neighbors too (and thus a triangle).

This algorithm improves on B-A in the sense that it enables a higher average clustering to be attained if desired.

It seems possible to have a disconnected graph with this algorithm since the initial m nodes may not be all linked to a new node on the first iteration like the B-A model.

#### References

[R353]

## 6.5.13 random lobster

```
random_lobster (n, p1, p2, seed=None)
```

Return a random lobster.

A lobster is a tree that reduces to a caterpillar when pruning all leaf nodes.

A caterpillar is a tree that reduces to a path graph when pruning all leaf nodes (p2=0).

### Parameters n: int

The expected number of nodes in the backbone

**p1**: float

Probability of adding an edge to the backbone

p2: float

Probability of adding an edge one level beyond backbone

seed: int, optional

Seed for random number generator (default=None).

## 6.5.14 random shell graph

### random\_shell\_graph (constructor, seed=None)

Return a random shell graph for the constructor given.

## Parameters constructor: a list of three-tuples

(n,m,d) for each shell starting at the center shell.

n: int

The number of nodes in the shell

 $\mathbf{m}: \mathrm{int}$ 

The number or edges in the shell

d: float

The ratio of inter-shell (next) edges to intra-shell edges. d=0 means no intra shell edges, d=1 for the last shell

seed: int, optional

Seed for random number generator (default=None).

## **Examples**

```
>>> constructor=[(10,20,0.8),(20,40,0.8)]
>>> G=nx.random_shell_graph(constructor)
```

## 6.5.15 random powerlaw tree

random\_powerlaw\_tree (n, gamma=3, seed=None, tries=100)

Return a tree with a powerlaw degree distribution.

Parameters n: int,

The number of nodes

gamma: float

Exponent of the power-law

**seed**: int, optional

Seed for random number generator (default=None).

tries: int

Number of attempts to adjust sequence to make a tree

#### **Notes**

A trial powerlaw degree sequence is chosen and then elements are swapped with new elements from a powerlaw distribution until the sequence makes a tree (#edges=#nodes-1).

## 6.5.16 random\_powerlaw\_tree\_sequence

random\_powerlaw\_tree\_sequence (n, gamma=3, seed=None, tries=100)

Return a degree sequence for a tree with a powerlaw distribution.

Parameters n: int.

The number of nodes

gamma: float

Exponent of the power-law

seed: int, optional

Seed for random number generator (default=None).

tries: int

Number of attempts to adjust sequence to make a tree

#### **Notes**

A trial powerlaw degree sequence is chosen and then elements are swapped with new elements from a powerlaw distribution until the sequence makes a tree (#edges=#nodes-1).

# 6.6 Degree Sequence

Generate graphs with a given degree sequence or expected degree sequence.

<pre>configuration_model(deg_sequence[,])</pre>	Return a random graph with the given degree sequence.
directed_configuration_model([,])	Return a directed_random graph with the given degree sequences.
<pre>expected_degree_graph(w[, seed, selfloops])</pre>	Return a random graph with given expected degrees.
havel_hakimi_graph(deg_sequence[, create_using])	Return a simple graph with given degree sequence constructed using
directed_havel_hakimi_graph(in_deg_sequence,)	Return a directed graph with the given degree sequences.
degree_sequence_tree(deg_sequence[,])	Make a tree for the given degree sequence.
random_degree_sequence_graph(sequence[,])	Return a simple random graph with the given degree sequence.

## 6.6.1 configuration\_model

configuration\_model (deg\_sequence, create\_using=None, seed=None)

Return a random graph with the given degree sequence.

The configuration model generates a random pseudograph (graph with parallel edges and self loops) by randomly assigning edges to match the given degree sequence.

Parameters deg\_sequence: list of integers

Each list entry corresponds to the degree of a node.

create\_using : graph, optional (default MultiGraph)

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

Seed for random number generator.

Returns G: MultiGraph

A graph with the specified degree sequence. Nodes are labeled starting at 0 with an index corresponding to the position in deg\_sequence.

### Raises NetworkXError

If the degree sequence does not have an even sum.

### See also:

is\_valid\_degree\_sequence

### Notes

As described by Newman [R319].

A non-graphical degree sequence (not realizable by some simple graph) is allowed since this function returns graphs with self loops and parallel edges. An exception is raised if the degree sequence does not have an even sum.

This configuration model construction process can lead to duplicate edges and loops. You can remove the self-loops and parallel edges (see below) which will likely result in a graph that doesn't have the exact degree sequence specified.

The density of self-loops and parallel edges tends to decrease as the number of nodes increases. However, typically the number of self-loops will approach a Poisson distribution with a nonzero mean, and similarly for the number of parallel edges. Consider a node with k stubs. The probability of being joined to another stub of the same node is basically (k-1)/N where k is the degree and N is the number of nodes. So the probability of a self-loop scales like c/N for some constant c. As N grows, this means we expect c self-loops. Similarly for parallel edges.

#### References

[R319]

### **Examples**

```
>>> from networkx.utils import powerlaw_sequence
>>> z=nx.utils.create_degree_sequence(100, powerlaw_sequence)
>>> G=nx.configuration_model(z)

To remove parallel edges:
>>> G=nx.Graph(G)

To remove self loops:
>>> G.remove_edges_from(G.selfloop_edges())
```

## 6.6.2 directed configuration model

Return a directed\_random graph with the given degree sequences.

The configuration model generates a random directed pseudograph (graph with parallel edges and self loops) by randomly assigning edges to match the given degree sequences.

```
Parameters in_degree_sequence : list of integers
```

Each list entry corresponds to the in-degree of a node.

```
out_degree_sequence : list of integers
```

Each list entry corresponds to the out-degree of a node.

```
create_using : graph, optional (default MultiDiGraph)
```

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

Seed for random number generator.

### Returns G: MultiDiGraph

A graph with the specified degree sequences. Nodes are labeled starting at 0 with an index corresponding to the position in deg\_sequence.

#### Raises NetworkXError

If the degree sequences do not have the same sum.

### See also:

```
configuration_model
```

#### **Notes**

Algorithm as described by Newman [R320].

A non-graphical degree sequence (not realizable by some simple graph) is allowed since this function returns graphs with self loops and parallel edges. An exception is raised if the degree sequences does not have the same sum.

This configuration model construction process can lead to duplicate edges and loops. You can remove the self-loops and parallel edges (see below) which will likely result in a graph that doesn't have the exact degree sequence specified. This "finite-size effect" decreases as the size of the graph increases.

#### References

[R320]

## **Examples**

```
>>> D=nx.DiGraph([(0,1),(1,2),(2,3)]) # directed path graph
>>> din=list(D.in_degree().values())
>>> dout=list(D.out_degree().values())
>>> din.append(1)
>>> dout[0]=2
>>> D=nx.directed_configuration_model(din,dout)

To remove parallel edges:
>>> D=nx.DiGraph(D)

To remove self loops:
>>> D.remove_edges_from(D.selfloop_edges())
```

## 6.6.3 expected degree graph

```
expected_degree_graph (w, seed=None, selfloops=True)
```

Return a random graph with given expected degrees.

Given a sequence of expected degrees  $W = (w_0, w_1, \dots, w_{n-1})$  of length n this algorithm assigns an edge between node u and node v with probability

$$p_{uv} = \frac{w_u w_v}{\sum_k w_k}.$$

Parameters w: list

The list of expected degrees.

### selfloops: bool (default=True)

Set to False to remove the possibility of self-loop edges.

seed: hashable object, optional

The seed for the random number generator.

### Returns Graph

### **Notes**

The nodes have integer labels corresponding to index of expected degrees input sequence.

The complexity of this algorithm is O(n+m) where n is the number of nodes and m is the expected number of edges.

The model in [R322] includes the possibility of self-loop edges. Set selfloops=False to produce a graph without self loops.

For finite graphs this model doesn't produce exactly the given expected degree sequence. Instead the expected degrees are as follows.

For the case without self loops (selfloops=False),

$$E[deg(u)] = \sum_{v \neq u} p_{uv} = w_u \left( 1 - \frac{w_u}{\sum_k w_k} \right).$$

NetworkX uses the standard convention that a self-loop edge counts 2 in the degree of a node, so with self loops (selfloops=True),

$$E[deg(u)] = \sum_{v \neq u} p_{uv} + 2p_{uu} = w_u \left( 1 + \frac{w_u}{\sum_k w_k} \right).$$

#### References

[R322], [R323]

### **Examples**

```
>>> z=[10 for i in range(100)]
>>> G=nx.expected_degree_graph(z)
```

## 6.6.4 havel\_hakimi\_graph

havel\_hakimi\_graph (deg\_sequence, create\_using=None)

Return a simple graph with given degree sequence constructed using the Havel-Hakimi algorithm.

### Parameters deg\_sequence: list of integers

Each integer corresponds to the degree of a node (need not be sorted).

create\_using : graph, optional (default Graph)

Return graph of this type. The instance will be cleared. Directed graphs are not allowed.

### Raises NetworkXException

For a non-graphical degree sequence (i.e. one not realizable by some simple graph).

### **Notes**

The Havel-Hakimi algorithm constructs a simple graph by successively connecting the node of highest degree to other nodes of highest degree, resorting remaining nodes by degree, and repeating the process. The resulting graph has a high degree-associativity. Nodes are labeled 1,.., len(deg\_sequence), corresponding to their position in deg\_sequence.

The basic algorithm is from Hakimi [R324] and was generalized by Kleitman and Wang [R325].

#### References

[R324], [R325]

## 6.6.5 directed havel hakimi graph

**directed\_havel\_hakimi\_graph** (*in\_deg\_sequence*, *out\_deg\_sequence*, *create\_using=None*)

Return a directed graph with the given degree sequences.

Parameters in\_deg\_sequence : list of integers

Each list entry corresponds to the in-degree of a node.

out\_deg\_sequence : list of integers

Each list entry corresponds to the out-degree of a node.

create\_using : graph, optional (default DiGraph)

Return graph of this type. The instance will be cleared.

## Returns G: DiGraph

A graph with the specified degree sequences. Nodes are labeled starting at 0 with an index corresponding to the position in deg\_sequence

### Raises NetworkXError

If the degree sequences are not digraphical.

#### See also:

```
configuration_model
```

### **Notes**

Algorithm as described by Kleitman and Wang [R321].

### References

[R321]

## 6.6.6 degree sequence tree

```
{\tt degree\_sequence\_tree}~(\textit{deg\_sequence}, \textit{create\_using=None})
```

Make a tree for the given degree sequence.

A tree has #nodes-#edges=1 so the degree sequence must have len(deg\_sequence)-sum(deg\_sequence)/2=1

## 6.6.7 random degree sequence graph

```
random_degree_sequence_graph (sequence, seed=None, tries=10)
```

Return a simple random graph with the given degree sequence.

If the maximum degree  $d_m$  in the sequence is  $O(m^{1/4})$  then the algorithm produces almost uniform random graphs in  $O(md_m)$  time where m is the number of edges.

### Parameters sequence: list of integers

Sequence of degrees

seed: hashable object, optional

Seed for random number generator

tries: int, optional

Maximum number of tries to create a graph

### Returns G: Graph

A graph with the specified degree sequence. Nodes are labeled starting at 0 with an index corresponding to the position in the sequence.

### Raises NetworkXUnfeasible

If the degree sequence is not graphical.

## NetworkXError

If a graph is not produced in specified number of tries

### See also:

```
is_valid_degree_sequence, configuration_model
```

#### **Notes**

The generator algorithm [R326] is not guaranteed to produce a graph.

#### References

[R326]

### **Examples**

```
>>> sequence = [1, 2, 2, 3]
>>> G = nx.random_degree_sequence_graph(sequence)
>>> sorted(G.degree().values())
[1, 2, 2, 3]
```

## 6.7 Random Clustered

Generate graphs with given degree and triangle sequence.

random\_clustered\_graph(joint\_degree\_sequence) Generate a random graph with the given joint degree and triangle degree

## 6.7.1 random clustered graph

random\_clustered\_graph (joint\_degree\_sequence, create\_using=None, seed=None)

Generate a random graph with the given joint degree and triangle degree sequence.

This uses a configuration model-like approach to generate a random pseudograph (graph with parallel edges and self loops) by randomly assigning edges to match the given independent edge and triangle degree sequence.

Parameters joint\_degree\_sequence : list of integer pairs

Each list entry corresponds to the independent edge degree and triangle degree of a node.

create\_using : graph, optional (default MultiGraph)

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

The seed for the random number generator.

Returns G: MultiGraph

A graph with the specified degree sequence. Nodes are labeled starting at 0 with an index corresponding to the position in deg\_sequence.

### Raises NetworkXError

If the independent edge degree sequence sum is not even or the triangle degree sequence sum is not divisible by 3.

#### **Notes**

As described by Miller [R341] (see also Newman [R342] for an equivalent description).

A non-graphical degree sequence (not realizable by some simple graph) is allowed since this function returns graphs with self loops and parallel edges. An exception is raised if the independent degree sequence does not have an even sum or the triangle degree sequence sum is not divisible by 3.

This configuration model-like construction process can lead to duplicate edges and loops. You can remove the self-loops and parallel edges (see below) which will likely result in a graph that doesn't have the exact degree sequence specified. This "finite-size effect" decreases as the size of the graph increases.

#### References

[R341], [R342]

6.7. Random Clustered 383

#### **Examples**

```
>>> deg_tri=[[1,0],[1,0],[1,0],[2,0],[1,0],[2,1],[0,1],[0,1]]
>>> G = nx.random_clustered_graph(deg_tri)

To remove parallel edges:
>>> G=nx.Graph(G)

To remove self loops:
>>> G.remove_edges_from(G.selfloop_edges())
```

## 6.8 Directed

Generators for some directed graphs.

gn\_graph: growing network gnc\_graph: growing network with copying gnr\_graph: growing network with redirection scale\_free\_graph: scale free directed graph

gn_graph(n[, kernel, create_using, seed])	Return the GN digraph with n nodes.
<pre>gnr_graph(n, p[, create_using, seed])</pre>	Return the GNR digraph with n nodes and redirection probability p.
<pre>gnc_graph(n[, create_using, seed])</pre>	Return the GNC digraph with n nodes.
scale_free_graph(n[, alpha, beta, gamma,])	Return a scale free directed graph.

## 6.8.1 gn\_graph

```
gn_graph (n, kernel=None, create_using=None, seed=None)
```

Return the GN digraph with n nodes.

The GN (growing network) graph is built by adding nodes one at a time with a link to one previously added node. The target node for the link is chosen with probability based on degree. The default attachment kernel is a linear function of degree.

The graph is always a (directed) tree.

### Parameters n: int

The number of nodes for the generated graph.

kernel: function

The attachment kernel.

create using: graph, optional (default DiGraph)

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

The seed for the random number generator.

### References

[R327]

## **Examples**

```
>>> D=nx.gn_graph(10)  # the GN graph
>>> G=D.to_undirected()  # the undirected version
```

To specify an attachment kernel use the kernel keyword

```
>>> D=nx.gn_graph(10,kernel=lambda x:x**1.5) # A_k=k^1.5
```

## 6.8.2 gnr\_graph

```
gnr_graph (n, p, create_using=None, seed=None)
```

Return the GNR digraph with n nodes and redirection probability p.

The GNR (growing network with redirection) graph is built by adding nodes one at a time with a link to one previously added node. The previous target node is chosen uniformly at random. With probability p the link is instead "redirected" to the successor node of the target. The graph is always a (directed) tree.

#### Parameters n: int

The number of nodes for the generated graph.

p: float

The redirection probability.

create\_using : graph, optional (default DiGraph)

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

The seed for the random number generator.

#### References

[R329]

#### **Examples**

```
>>> D=nx.gnr_graph(10,0.5) # the GNR graph
>>> G=D.to_undirected() # the undirected version
```

## 6.8.3 gnc graph

```
gnc_graph (n, create_using=None, seed=None)
```

Return the GNC digraph with n nodes.

The GNC (growing network with copying) graph is built by adding nodes one at a time with a links to one previously added node (chosen uniformly at random) and to all of that node's successors.

Parameters n: int

The number of nodes for the generated graph.

create\_using : graph, optional (default DiGraph)

6.8. Directed 385

Return graph of this type. The instance will be cleared.

seed: hashable object, optional

The seed for the random number generator.

#### References

[R328]

## 6.8.4 scale free graph

```
scale_free_graph (n, alpha=0.41, beta=0.54, gamma=0.05, delta_in=0.2, delta_out=0, cre-
ate_using=None, seed=None)
Return a scale free directed graph.
```

arm a scale free directed graph

Parameters n : integer

Number of nodes in graph

alpha: float

Probability for adding a new node connected to an existing node chosen randomly according to the in-degree distribution.

beta: float

Probability for adding an edge between two existing nodes. One existing node is chosen randomly according the in-degree distribution and the other chosen randomly according to the out-degree distribution.

gamma: float

Probability for adding a new node conecgted to an existing node chosen randomly according to the out-degree distribution.

delta\_in: float

Bias for choosing ndoes from in-degree distribution.

delta out: float

Bias for choosing ndoes from out-degree distribution.

create\_using : graph, optional (default MultiDiGraph)

Use this graph instance to start the process (default=3-cycle).

**seed**: integer, optional

Seed for random number generator

### **Notes**

The sum of alpha, beta, and gamma must be 1.

#### References

[R330]

### **Examples**

```
>>> G=nx.scale_free_graph(100)
```

## 6.9 Geometric

Generators for geometric graphs.

<pre>random_geometric_graph(n, radius[, dim, pos])</pre>	Return the random geometric graph in the unit cube.
<pre>geographical_threshold_graph(n, theta[,])</pre>	Return a geographical threshold graph.
waxman_graph(n[, alpha, beta, L, domain])	Return a Waxman random graph.
$navigable\_small\_world\_graph(n[, p, q, r,])$	Return a navigable small-world graph.

## 6.9.1 random\_geometric\_graph

```
random_geometric_graph (n, radius, dim=2, pos=None)
```

Return the random geometric graph in the unit cube.

The random geometric graph model places n nodes uniformly at random in the unit cube Two nodes u, v are connected with an edge if d(u, v) <= r where d is the Euclidean distance and r is a radius threshold.

#### Parameters n: int

Number of nodes

#### radius: float

Distance threshold value

dim: int, optional

Dimension of graph

pos: dict, optional

A dictionary keyed by node with node positions as values.

Returns Graph

#### **Notes**

This uses an  $n^2$  algorithm to build the graph. A faster algorithm is possible using k-d trees.

The pos keyword can be used to specify node positions so you can create an arbitrary distribution and domain for positions. If you need a distance function other than Euclidean you'll have to hack the algorithm.

E.g to use a 2d Gaussian distribution of node positions with mean (0,0) and std. dev. 2

```
>>> import random
>>> n=20
>>> p=dict((i,(random.gauss(0,2),random.gauss(0,2))) for i in range(n))
>>> G = nx.random_geometric_graph(n,0.2,pos=p)
```

6.9. Geometric 387

#### References

[R335]

## **Examples**

```
>>> G = nx.random_geometric_graph(20,0.1)
```

## 6.9.2 geographical threshold graph

```
geographical_threshold_graph (n, theta, alpha=2, dim=2, pos=None, weight=None) Return a geographical threshold graph.
```

The geographical threshold graph model places n nodes uniformly at random in a rectangular domain. Each node u is assigned a weight  $w_u$ . Two nodes u, v are connected with an edge if

$$w_u + w_v \ge \theta r^{\alpha}$$

where r is the Euclidean distance between u and v, and  $\theta$ ,  $\alpha$  are parameters.

Parameters n: int

Number of nodes

theta: float

Threshold value

alpha: float, optional

Exponent of distance function

dim: int, optional

Dimension of graph

pos: dict

Node positions as a dictionary of tuples keyed by node.

weight: dict

Node weights as a dictionary of numbers keyed by node.

Returns Graph

### **Notes**

If weights are not specified they are assigned to nodes by drawing randomly from an the exponential distribution with rate parameter  $\lambda=1$ . To specify a weights from a different distribution assign them to a dictionary and pass it as the weight= keyword

```
>>> import random
>>> n = 20
>>> w=dict((i,random.expovariate(5.0)) for i in range(n))
>>> G = nx.geographical_threshold_graph(20,50,weight=w)
```

If node positions are not specified they are randomly assigned from the uniform distribution.

#### References

```
[R332], [R333]
```

### **Examples**

```
>>> G = nx.geographical_threshold_graph(20,50)
```

## 6.9.3 waxman graph

```
\mathbf{waxman\_graph} \ (n, alpha=0.4, beta=0.1, L=None, domain=(0, 0, 1, 1))
```

Return a Waxman random graph.

The Waxman random graph models place n nodes uniformly at random in a rectangular domain. Two nodes u,v are connected with an edge with probability

$$p = \alpha * exp(-d/(\beta * L)).$$

This function implements both Waxman models.

**Waxman-1:** L **not specified** The distance d is the Euclidean distance between the nodes u and v. L is the maximum distance between all nodes in the graph.

**Waxman-2:** L specified The distance d is chosen randomly in [0, L].

Parameters n: int

Number of nodes

alpha: float

Model parameter

beta: float

Model parameter

L: float, optional

Maximum distance between nodes. If not specified the actual distance is calculated.

domain: tuple of numbers, optional

Domain size (xmin, ymin, xmax, ymax)

Returns G: Graph

## References

[R336]

# 6.9.4 navigable\_small\_world\_graph

```
navigable_small_world_graph (n, p=1, q=1, r=2, dim=2, seed=None) Return a navigable small-world graph.
```

6.9. Geometric 389

A navigable small-world graph is a directed grid with additional long-range connections that are chosen randomly. From [R334]:

Begin with a set of nodes that are identified with the set of lattice points in an  $n \times n$  square,  $(i, j) : i \in 1, 2, ..., n, j \in 1, 2, ..., n$  and define the lattice distance between two nodes (i, j) and (k, l) to be the number of "lattice steps" separating them: d((i, j), (k, l)) = |k - i| + |l - j|.

For a universal constant p, the node u has a directed edge to every other node within lattice distance p (local contacts).

For universal constants  $q \ge 0$  and  $r \ge 0$  construct directed edges from u to q other nodes (long-range contacts) using independent random trials; the i'th directed edge from u has endpoint v with probability proportional to  $d(u,v)^{-r}$ .

#### Parameters n: int

The number of nodes.

p: int

The diameter of short range connections. Each node is connected to every other node within lattice distance p.

q: int

The number of long-range connections for each node.

r: float

Exponent for decaying probability of connections. The probability of connecting to a node at lattice distance d is 1/d^r.

dim: int

Dimension of grid

seed: int, optional

Seed for random number generator (default=None).

#### References

[R334]

# 6.10 Hybrid

### Hybrid

kl_connected_subgraph(G, k, l[, low_memory,])	Returns the maximum locally (k,l) connected subgraph of G.
<pre>is_kl_connected(G, k, l[, low_memory])</pre>	Returns True if G is kl connected.

## 6.10.1 kl\_connected\_subgraph

**k1\_connected\_subgraph** (*G*, *k*, *l*, *low\_memory=False*, *same\_as\_graph=False*) Returns the maximum locally (k,l) connected subgraph of G.

(k,l)-connected subgraphs are presented by Fan Chung and Li in "The Small World Phenomenon in hybrid power law graphs" to appear in "Complex Networks" (Ed. E. Ben-Naim) Lecture Notes in Physics, Springer

(2004)

low\_memory=True then use a slightly slower, but lower memory version same\_as\_graph=True then return a tuple with subgraph and pflag for if G is kl-connected

## 6.10.2 is kl connected

**is\_kl\_connected** (*G*, *k*, *l*, *low\_memory=False*)

Returns True if G is kl connected.

# 6.11 Line Graph

Line graph algorithms.

## 6.11.1 Undirected Graphs

For an undirected graph G without multiple edges, each edge can be written as a set  $\{u,v\}$ . Its line graph L has the edges of G as its nodes. If x and y are two nodes in L, then  $\{x,y\}$  is an edge in L if and only if the intersection of x and y is nonempty. Thus, the set of all edges is determined by the set of all pair-wise intersections of edges in G.

Trivially, every edge  $x=\{u,v\}$  in G would have a nonzero intersection with itself, and so every node in L should have a self-loop. This is not so interesting, and the original context of line graphs was with simple graphs, which had no self-loops or multiple edges. The line graph was also meant to be simple graph and thus, self-loops in L are not part of the standard definition of a line graph. In a pair-wise intersection matrix, this is analogous to not including the diagonal as part of the line graph definition.

Self-loops and multiple edges in G add nodes to L in a natural way, and do not require any fundamental changes to the definition. It might be argued that the self-loops we excluded before should now be included. However, the self-loops are still "trivial" in some sense and thus, are usually excluded.

## 6.11.2 Directed Graphs

For a directed graph G without multiple edges, each edge can be written as a tuple (u,v). Its line graph L has the edges of G as its nodes. If x=(a,b) and y=(c,d) are two nodes in L, then (x,y) is an edge in L if and only if the tail of x matches the head of y=e,g,b=c.

Due to the directed nature of the edges, it is no longer the case that every edge x=(u,v) should be connected to itself with a self-loop in L. Now, the only time self-loops arise is if G itself has a self-loop. So such self-loops are no longer "trivial" but instead, represent essential features of the topology of G. For this reason, the historical development of line digraphs is such that self-loops are included. When the graph G has multiple edges, once again only superficial changes are required to the definition.

### 6.11.3 References

**Harary, Frank, and Norman, Robert Z., "Some properties of line digraphs",** Rend. Circ. Mat. Palermo, II. Ser. 9 (1960), 161–168.

Hemminger, R. L.; Beineke, L. W. (1978), "Line graphs and line digraphs", in Beineke, L. W.; Wilson, R. J., Selected Topics in Graph Theory, Academic Press Inc., pp. 271–305.

line\_graph(G[, create\_using]) Return the line graph of the graph or digraph G.

6.11. Line Graph 391

## 6.11.4 line graph

```
line_graph(G, create_using=None)
```

Return the line graph of the graph or digraph G.

The line graph of a graph G has a node for each edge in G and an edge between those nodes if the two edges in G share a common node. For directed graphs, nodes are connected only if they form a directed path of length 2.

The nodes of the line graph are 2-tuples of nodes in the original graph (or 3-tuples for multigraphs, with the key of the edge as the 3rd element).

For more discussion, see the docstring in networks.generators.line.

**Parameters G**: graph

A NetworkX Graph, DiGraph, MultiGraph, or MultiDigraph.

Returns L: graph

The line graph of G.

#### **Notes**

Graph, node, and edge data are not propagated to the new graph. For undirected graphs, the nodes in G must be sortable—otherwise, the constructed line graph may not be correct.

### **Examples**

```
>>> G = nx.star_graph(3)
>>> L = nx.line_graph(G)
>>> print(sorted(map(sorted, L.edges()))) # makes a clique, K3
[[(0, 1), (0, 2)], [(0, 1), (0, 3)], [(0, 2), (0, 3)]]
```

# 6.12 Ego Graph

Ego graph.

ego\_graph(G, n[, radius, center, ...]) Returns induced subgraph of neighbors centered at node n within a given radius.

## 6.12.1 ego\_graph

```
ego\_graph(G, n, radius=1, center=True, undirected=False, distance=None)
```

Returns induced subgraph of neighbors centered at node n within a given radius.

Parameters G: graph

A NetworkX Graph or DiGraph

n: node

A single node

radius: number, optional

Include all neighbors of distance<=radius from n.

center: bool, optional

If False, do not include center node in graph

undirected: bool, optional

If True use both in- and out-neighbors of directed graphs.

distance: key, optional

Use specified edge data key as distance. For example, setting distance='weight' will use the edge weight to measure the distance from the node n.

#### **Notes**

For directed graphs D this produces the "out" neighborhood or successors. If you want the neighborhood of predecessors first reverse the graph with D.reverse(). If you want both directions use the keyword argument undirected=True.

Node, edge, and graph attributes are copied to the returned subgraph.

## 6.13 Stochastic

Stocastic graph.

stochastic\_graph(G[, copy, weight]) Return a right-stochastic representation of G.

## 6.13.1 stochastic graph

stochastic\_graph(G, copy=True, weight='weight')

Return a right-stochastic representation of G.

A right-stochastic graph is a weighted digraph in which all of the node (out) neighbors edge weights sum to 1.

Parameters G: directed graph

A NetworkX DiGraph

copy: boolean, optional

If True make a copy of the graph, otherwise modify the original graph

**weight**: edge attribute key (optional, default='weight')

Edge data key used for weight. If no attribute is found for an edge the edge weight is set to 1. Weights must be positive numbers.

## 6.14 Intersection

Generators for random intersection graphs.

$\verb"uniform_random_intersection_graph"(n, m, p[,])$	Return a uniform random intersection graph.
$k$ _random_intersection_graph $(n, m, k)$	Return a intersection graph with randomly chosen attribute sets for e
<pre>general_random_intersection_graph(n, m, p)</pre>	Return a random intersection graph with independent probabilities f

6.14. Intersection 393

## 6.14.1 uniform random intersection graph

```
uniform\_random\_intersection\_graph(n, m, p, seed=None)
     Return a uniform random intersection graph.
           Parameters n: int
                   The number of nodes in the first bipartite set (nodes)
               m: int
                   The number of nodes in the second bipartite set (attributes)
               p: float
                   Probability of connecting nodes between bipartite sets
               seed: int, optional
                   Seed for random number generator (default=None).
     See also:
     gnp_random_graph
     References
     [R339], [R340]
6.14.2 k random intersection graph
k_random_intersection_graph(n, m, k)
     Return a intersection graph with randomly chosen attribute sets for each node that are of equal size (k).
           Parameters n: int
                   The number of nodes in the first bipartite set (nodes)
               m: int
                   The number of nodes in the second bipartite set (attributes)
               k: float
                   Size of attribute set to assign to each node.
               seed: int, optional
                   Seed for random number generator (default=None).
     See also:
     gnp_random_graph, uniform_random_intersection_graph
     References
```

[R338]

## 6.14.3 general\_random\_intersection\_graph

### $general\_random\_intersection\_graph(n, m, p)$

Return a random intersection graph with independent probabilities for connections between node and attribute sets.

#### Parameters n: int

The number of nodes in the first bipartite set (nodes)

m: int

The number of nodes in the second bipartite set (attributes)

p: list of floats of length m

Probabilities for connecting nodes to each attribute

seed: int, optional

Seed for random number generator (default=None).

#### See also:

```
gnp_random_graph, uniform_random_intersection_graph
```

#### References

[R337]

## 6.15 Social Networks

Famous social networks.

karate_club_graph()	Return Zachary's Karate club graph.
davis_southern_women_graph()	Return Davis Southern women social network.
florentine_families_graph()	Return Florentine families graph.

## 6.15.1 karate\_club\_graph

#### karate\_club\_graph()

Return Zachary's Karate club graph.

### References

[R359], [R360]

## 6.15.2 davis\_southern\_women\_graph

## ${\tt davis\_southern\_women\_graph}\;(\;)$

Return Davis Southern women social network.

This is a bipartite graph.

6.15. Social Networks 395

### References

[R357]

# 6.15.3 florentine\_families\_graph

## florentine\_families\_graph()

Return Florentine families graph.

## References

[R358]

## LINEAR ALGEBRA

# 7.1 Graph Matrix

Adjacency matrix and incidence matrix of graphs.

adjacency_matrix(G[, nodelist, weight])	Return adjacency matrix of G.
<pre>incidence_matrix(G[, nodelist, edgelist,])</pre>	Return incidence matrix of G.

## 7.1.1 adjacency\_matrix

 $\verb"adjacency_matrix" (G, nodelist=None, weight='weight')$ 

Return adjacency matrix of G.

Parameters G: graph

A NetworkX graph

nodelist: list, optional

The rows and columns are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

weight : string or None, optional (default='weight')

The edge data key used to provide each value in the matrix. If None, then each edge has weight 1.

**Returns** A : SciPy sparse matrix

Adjacency matrix representation of G.

### See also:

```
to_numpy_matrix, to_scipy_sparse_matrix, to_dict_of_dicts
```

## **Notes**

For directed graphs, entry i,j corresponds to an edge from i to j.

If you want a pure Python adjacency matrix representation try networkx.convert.to\_dict\_of\_dicts which will return a dictionary-of-dictionaries format that can be addressed as a sparse matrix.

For MultiGraph/MultiDiGraph with parallel edges the weights are summed. See to\_numpy\_matrix for other options.

The convention used for self-loop edges in graphs is to assign the diagonal matrix entry value to the edge weight attribute (or the number 1 if the edge has no weight attribute). If the alternate convention of doubling the edge weight is desired the resulting Scipy sparse matrix can be modified as follows:

```
>>> import scipy as sp
>>> G = nx.Graph([(1,1)])
>>> A = nx.adjacency_matrix(G)
>>> print(A.todense())
[[1]]
>>> A.setdiag(A.diagonal()*2)
>>> print(A.todense())
[[2]]
```

## 7.1.2 incidence\_matrix

incidence\_matrix (G, nodelist=None, edgelist=None, oriented=False, weight=None)
Return incidence matrix of G.

The incidence matrix assigns each row to a node and each column to an edge. For a standard incidence matrix a 1 appears wherever a row's node is incident on the column's edge. For an oriented incidence matrix each edge is assigned an orientation (arbitrarily for undirected and aligning to direction for directed). A -1 appears for the tail of an edge and 1 for the head of the edge. The elements are zero otherwise.

### Parameters G: graph

A NetworkX graph

nodelist: list, optional (default= all nodes in G)

The rows are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

edgelist: list, optional (default= all edges in G)

The columns are ordered according to the edges in edgelist. If edgelist is None, then the ordering is produced by G.edges().

### oriented: bool, optional (default=False)

If True, matrix elements are +1 or -1 for the head or tail node respectively of each edge. If False, +1 occurs at both nodes.

```
weight : string or None, optional (default=None)
```

The edge data key used to provide each value in the matrix. If None, then each edge has weight 1. Edge weights, if used, should be positive so that the orientation can provide the sign.

### Returns A: SciPy sparse matrix

The incidence matrix of G.

#### **Notes**

For MultiGraph/MultiDiGraph, the edges in edgelist should be (u,v,key) 3-tuples.

"Networks are the best discrete model for so many problems in applied mathematics" [R361].

#### References

[R361]

# 7.2 Laplacian Matrix

Laplacian matrix of graphs.

laplacian_matrix(G[, nodelist, weight])	Return the Laplacian matrix of G.
normalized_laplacian_matrix(G[, nodelist,])	Return the normalized Laplacian matrix of G.
directed_laplacian_matrix(G[, nodelist,])	Return the directed Laplacian matrix of G.

## 7.2.1 laplacian\_matrix

laplacian\_matrix (G, nodelist=None, weight='weight')

Return the Laplacian matrix of G.

The graph Laplacian is the matrix L = D - A, where A is the adjacency matrix and D is the diagonal matrix of node degrees.

Parameters G: graph

A NetworkX graph

nodelist: list, optional

The rows and columns are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

weight : string or None, optional (default='weight')

The edge data key used to compute each value in the matrix. If None, then each edge has weight 1.

**Returns** L : SciPy sparse matrix

The Laplacian matrix of G.

### See also:

to\_numpy\_matrix, normalized\_laplacian\_matrix

### **Notes**

For MultiGraph/MultiDiGraph, the edges weights are summed.

## 7.2.2 normalized laplacian matrix

normalized\_laplacian\_matrix(G, nodelist=None, weight='weight')

Return the normalized Laplacian matrix of G.

The normalized graph Laplacian is the matrix

$$N = D^{-1/2}LD^{-1/2}$$

where L is the graph Laplacian and D is the diagonal matrix of node degrees.

Parameters G: graph

A NetworkX graph

nodelist: list, optional

The rows and columns are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

weight : string or None, optional (default='weight')

The edge data key used to compute each value in the matrix. If None, then each edge has weight 1.

**Returns N**: NumPy matrix

The normalized Laplacian matrix of G.

#### See also:

laplacian\_matrix

#### **Notes**

For MultiGraph/MultiDiGraph, the edges weights are summed. See to\_numpy\_matrix for other options.

If the Graph contains selfloops, D is defined as diag(sum(A,1)), where A is the adjacency matrix [R364].

#### References

[R363], [R364]

## 7.2.3 directed laplacian matrix

**directed\_laplacian\_matrix** (*G*, nodelist=None, weight='weight', walk\_type=None, alpha=0.95)

Return the directed Laplacian matrix of G.

The graph directed Laplacian is the matrix

$$L = I - (\Phi^{1/2}P\Phi^{-1/2} + \Phi^{-1/2}P^T\Phi^{1/2})/2$$

where I is the identity matrix, P is the transition matrix of the graph, and  $\Phi$  a matrix with the Perron vector of P in the diagonal and zeros elsewhere.

Depending on the value of walk\_type, P can be the transition matrix induced by a random walk, a lazy random walk, or a random walk with teleportation (PageRank).

**Parameters G**: DiGraph

A NetworkX graph

nodelist: list, optional

The rows and columns are ordered according to the nodes in nodelist. If nodelist is None, then the ordering is produced by G.nodes().

weight : string or None, optional (default='weight')

The edge data key used to compute each value in the matrix. If None, then each edge has weight 1.

walk\_type : string or None, optional (default=None)

If None, P is selected depending on the properties of the graph. Otherwise is one of 'random', 'lazy', or 'pagerank'

alpha: real

(1 - alpha) is the teleportation probability used with pagerank

**Returns** L : NumPy array

Normalized Laplacian of G.

Raises NetworkXError

If NumPy cannot be imported

### NetworkXNotImplemnted

If G is not a DiGraph

### See also:

laplacian\_matrix

#### **Notes**

Only implemented for DiGraphs

#### References

[R362]

# 7.3 Spectrum

Eigenvalue spectrum of graphs.

laplacian_spectrum( $G[$ , weight])	Return eigenvalues of the Laplacian of G
adjacency_spectrum(G[, weight])	Return eigenvalues of the adjacency matrix of G.

## 7.3.1 laplacian\_spectrum

laplacian\_spectrum(G, weight='weight')

Return eigenvalues of the Laplacian of G

Parameters G: graph

A NetworkX graph

weight : string or None, optional (default='weight')

The edge data key used to compute each value in the matrix. If None, then each edge has weight 1.

Returns evals: NumPy array

Eigenvalues

7.3. Spectrum 401

#### See also:

laplacian\_matrix

#### **Notes**

For MultiGraph/MultiDiGraph, the edges weights are summed. See to\_numpy\_matrix for other options.

## 7.3.2 adjacency\_spectrum

```
adjacency_spectrum(G, weight='weight')
```

Return eigenvalues of the adjacency matrix of G.

Parameters G: graph

A NetworkX graph

weight : string or None, optional (default='weight')

The edge data key used to compute each value in the matrix. If None, then each edge has weight 1.

Returns evals: NumPy array

Eigenvalues

### See also:

adjacency\_matrix

#### **Notes**

For MultiGraph/MultiDiGraph, the edges weights are summed. See to\_numpy\_matrix for other options.

# 7.4 Algebraic Connectivity

Algebraic connectivity and Fiedler vectors of undirected graphs.

$algebraic\_connectivity(G[,weight,])$	Return the algebraic connectivity of an undirected graph.
fiedler_vector(G[, weight, normalized, tol,])	Return the Fiedler vector of a connected undirected graph.
$spectral\_ordering(G[, weight, normalized,])$	Compute the spectral_ordering of a graph.

## 7.4.1 algebraic\_connectivity

**algebraic\_connectivity** (*G*, weight='weight', normalized=False, tol=1e-08, method='tracemin') Return the algebraic connectivity of an undirected graph.

The algebraic connectivity of a connected undirected graph is the second smallest eigenvalue of its Laplacian matrix.

**Parameters G**: NetworkX graph

An undirected graph.

weight: object, optional

The data key used to determine the weight of each edge. If None, then each edge has unit weight. Default value: None.

normalized: bool, optional

Whether the normalized Laplacian matrix is used. Default value: False.

tol: float, optional

Tolerance of relative residual in eigenvalue computation. Default value: 1e-8.

method: string, optional

Method of eigenvalue computation. It should be one of 'tracemin' (TraceMIN), 'lanczos' (Lanczos iteration) and 'lobpcg' (LOBPCG). Default value: 'tracemin'.

The TraceMIN algorithm uses a linear system solver. The following values allow specifying the solver to be used.

Value	Solver
'tracemin_pcg'	Preconditioned conjugate gradient method
'tracemin_chol'	Cholesky factorization
'tracemin_lu'	LU factorization

Returns algebraic\_connectivity: float

Algebraic connectivity.

### Raises NetworkXNotImplemented

If G is directed.

### NetworkXError

If G has less than two nodes.

#### See also:

laplacian\_matrix

#### Notes

Edge weights are interpreted by their absolute values. For MultiGraph's, weights of parallel edges are summed. Zero-weighted edges are ignored.

To use Cholesky factorization in the TraceMIN algorithm, the scikits.sparse package must be installed.

## 7.4.2 fiedler\_vector

**fiedler\_vector** (*G*, weight='weight', normalized=False, tol=1e-08, method='tracemin')

Return the Fiedler vector of a connected undirected graph.

The Fiedler vector of a connected undirected graph is the eigenvector corresponding to the second smallest eigenvalue of the Laplacian matrix of of the graph.

Parameters G: NetworkX graph

An undirected graph.

weight: object, optional

The data key used to determine the weight of each edge. If None, then each edge has unit weight. Default value: None.

normalized: bool, optional

Whether the normalized Laplacian matrix is used. Default value: False.

tol: float, optional

Tolerance of relative residual in eigenvalue computation. Default value: 1e-8.

method: string, optional

Method of eigenvalue computation. It should be one of 'tracemin' (TraceMIN), 'lanczos' (Lanczos iteration) and 'lobpcg' (LOBPCG). Default value: 'tracemin'.

The TraceMIN algorithm uses a linear system solver. The following values allow specifying the solver to be used.

Value	Solver
'tracemin_pcg'	Preconditioned conjugate gradient method
'tracemin_chol'	Cholesky factorization
'tracemin_lu'	LU factorization

Returns fiedler\_vector: NumPy array of floats.

Fiedler vector.

#### Raises NetworkXNotImplemented

If G is directed.

#### NetworkXError

If G has less than two nodes or is not connected.

#### See also:

laplacian matrix

### Notes

Edge weights are interpreted by their absolute values. For MultiGraph's, weights of parallel edges are summed. Zero-weighted edges are ignored.

To use Cholesky factorization in the TraceMIN algorithm, the scikits.sparse package must be installed.

## 7.4.3 spectral\_ordering

 $\begin{tabular}{ll} \textbf{spectral\_ordering} (G, weight='weight', normalized=False, tol=1e-08, method='tracemin') \\ \textbf{Compute the spectral\_ordering of a graph.} \end{tabular}$ 

The spectral ordering of a graph is an ordering of its nodes where nodes in the same weakly connected components appear contiguous and ordered by their corresponding elements in the Fiedler vector of the component.

### Parameters G: NetworkX graph

A graph.

weight: object, optional

The data key used to determine the weight of each edge. If None, then each edge has unit weight. Default value: None.

normalized: bool, optional

Whether the normalized Laplacian matrix is used. Default value: False.

tol: float, optional

Tolerance of relative residual in eigenvalue computation. Default value: 1e-8.

method: string, optional

Method of eigenvalue computation. It should be one of 'tracemin' (TraceMIN), 'lanczos' (Lanczos iteration) and 'lobpcg' (LOBPCG). Default value: 'tracemin'.

The TraceMIN algorithm uses a linear system solver. The following values allow specifying the solver to be used.

Value	Solver
'tracemin_pcg'	Preconditioned conjugate gradient method
'tracemin_chol'	Cholesky factorization
'tracemin_lu'	LU factorization

**Returns** spectral\_ordering: NumPy array of floats.

Spectral ordering of nodes.

#### Raises NetworkXError

If G is empty.

#### See also:

laplacian\_matrix

#### **Notes**

Edge weights are interpreted by their absolute values. For MultiGraph's, weights of parallel edges are summed. Zero-weighted edges are ignored.

To use Cholesky factorization in the TraceMIN algorithm, the scikits.sparse package must be installed.

## 7.5 Attribute Matrices

Functions for constructing matrix-like objects from graph attributes.

<pre>attr_matrix(G[, edge_attr, node_attr,])</pre>	Returns a NumPy matrix using attributes from G.
attr_sparse_matrix(G[, edge_attr,])	Returns a SciPy sparse matrix using attributes from G.

## 7.5.1 attr\_matrix

Returns a NumPy matrix using attributes from G.

If only G is passed in, then the adjacency matrix is constructed.

Let A be a discrete set of values for the node attribute  $node_a ttr$ . Then the elements of A represent the rows and columns of the constructed matrix. Now, iterate through every edge e=(u,v) in G and consider the value of the edge attribute  $edge_a ttr$ . If ua and va are the values of the node attribute  $node_a ttr$  for u and v, respectively, then the value of the edge attribute is added to the matrix element at (ua, va).

Parameters G: graph

7.5. Attribute Matrices 405

The NetworkX graph used to construct the NumPy matrix.

```
edge_attr: str, optional
```

Each element of the matrix represents a running total of the specified edge attribute for edges whose node attributes correspond to the rows/cols of the matrix. The attribute must be present for all edges in the graph. If no attribute is specified, then we just count the number of edges whose node attributes correspond to the matrix element.

```
node attr: str, optional
```

Each row and column in the matrix represents a particular value of the node attribute. The attribute must be present for all nodes in the graph. Note, the values of this attribute should be reliably hashable. So, float values are not recommended. If no attribute is specified, then the rows and columns will be the nodes of the graph.

#### normalized: bool, optional

If True, then each row is normalized by the summation of its values.

```
rc_order: list, optional
```

A list of the node attribute values. This list specifies the ordering of rows and columns of the array. If no ordering is provided, then the ordering will be random (and also, a return value).

### **Returns** M: NumPy matrix

The attribute matrix.

#### ordering: list

If  $rc_order$  was specified, then only the matrix is returned. However, if  $rc_order$  was None, then the ordering used to construct the matrix is returned as well.

### Other Parameters dtype: NumPy data-type, optional

A valid NumPy dtype used to initialize the array. Keep in mind certain dtypes can yield unexpected results if the array is to be normalized. The parameter is passed to numpy.zeros(). If unspecified, the NumPy default is used.

```
order: {'C', 'F'}, optional
```

Whether to store multidimensional data in C- or Fortran-contiguous (row- or columnwise) order in memory. This parameter is passed to numpy.zeros(). If unspecified, the NumPy default is used.

#### **Examples**

Construct an adjacency matrix:

Alternatively, we can obtain the matrix describing edge thickness.

We can also color the nodes and ask for the probability distribution over all edges (u,v) describing:

 $Pr(v \text{ has color } Y \mid u \text{ has color } X)$ 

For example, the above tells us that for all edges (u,v):

```
Pr(v \text{ is red} \mid u \text{ is red}) = 1/3 Pr(v \text{ is blue} \mid u \text{ is red}) = 2/3
```

 $Pr(v \text{ is red} \mid u \text{ is blue}) = 1 Pr(v \text{ is blue} \mid u \text{ is blue}) = 0$ 

Finally, we can obtain the total weights listed by the node colors.

Thus, the total weight over all edges (u,v) with u and v having colors:

(red, red) is 3 # the sole contribution is from edge (0,1) (red, blue) is 2 # contributions from edges (0,2) and (1,2) (blue, red) is 2 # same as (red, blue) since graph is undirected (blue, blue) is 0 # there are no edges with blue endpoints

### 7.5.2 attr sparse matrix

Returns a SciPy sparse matrix using attributes from G.

If only G is passed in, then the adjacency matrix is constructed.

Let A be a discrete set of values for the node attribute  $node_attr$ . Then the elements of A represent the rows and columns of the constructed matrix. Now, iterate through every edge e=(u,v) in G and consider the value of the edge attribute  $edge_attr$ . If ua and va are the values of the node attribute  $node_attr$  for u and v, respectively, then the value of the edge attribute is added to the matrix element at (ua, va).

Parameters G: graph

The NetworkX graph used to construct the NumPy matrix.

```
edge_attr : str, optional
```

Each element of the matrix represents a running total of the specified edge attribute for edges whose node attributes correspond to the rows/cols of the matrix. The attribute must be present for all edges in the graph. If no attribute is specified, then we just count the number of edges whose node attributes correspond to the matrix element.

node\_attr: str, optional

7.5. Attribute Matrices 407

Each row and column in the matrix represents a particular value of the node attribute. The attribute must be present for all nodes in the graph. Note, the values of this attribute should be reliably hashable. So, float values are not recommended. If no attribute is specified, then the rows and columns will be the nodes of the graph.

### normalized: bool, optional

If True, then each row is normalized by the summation of its values.

```
rc order: list, optional
```

A list of the node attribute values. This list specifies the ordering of rows and columns of the array. If no ordering is provided, then the ordering will be random (and also, a return value).

### **Returns** M : SciPy sparse matrix

The attribute matrix.

#### ordering: list

If  $rc_order$  was specified, then only the matrix is returned. However, if  $rc_order$  was None, then the ordering used to construct the matrix is returned as well.

### Other Parameters dtype: NumPy data-type, optional

A valid NumPy dtype used to initialize the array. Keep in mind certain dtypes can yield unexpected results if the array is to be normalized. The parameter is passed to numpy.zeros(). If unspecified, the NumPy default is used.

#### **Examples**

Construct an adjacency matrix:

Alternatively, we can obtain the matrix describing edge thickness.

We can also color the nodes and ask for the probability distribution over all edges (u,v) describing:

Pr(v has color Y | u has color X)

```
>>> G.node[0]['color'] = 'red'
>>> G.node[1]['color'] = 'red'
>>> G.node[2]['color'] = 'blue'
>>> rc = ['red', 'blue']
>>> M = nx.attr_sparse_matrix(G, node_attr='color',
```

normalized

node\_attr=

409

For example, the above tells us that for all edges (u,v):

```
Pr(v \text{ is red} \mid u \text{ is red}) = 1/3 Pr(v \text{ is blue} \mid u \text{ is red}) = 2/3
```

 $Pr(v \text{ is red} \mid u \text{ is blue}) = 1 Pr(v \text{ is blue} \mid u \text{ is blue}) = 0$ 

Finally, we can obtain the total weights listed by the node colors.

Thus, the total weight over all edges (u,v) with u and v having colors:

(red, red) is 3 # the sole contribution is from edge (0,1) (red, blue) is 2 # contributions from edges (0,2) and (1,2) (blue, red) is 2 # same as (red, blue) since graph is undirected (blue, blue) is 0 # there are no edges with blue endpoints

7.5. Attribute Matrices

**CHAPTER** 

**EIGHT** 

## CONVERTING TO AND FROM OTHER DATA FORMATS

# 8.1 To NetworkX Graph

Functions to convert NetworkX graphs to and from other formats.

The preferred way of converting data to a NetworkX graph is through the graph constuctor. The constructor calls the to\_networkx\_graph() function which attempts to guess the input type and convert it automatically.

## 8.1.1 Examples

Create a graph with a single edge from a dictionary of dictionaries

```
>>> d={0: {1: 1}} # dict-of-dicts single edge (0,1) >>> G=nx.Graph(d)
```

### 8.1.2 See Also

nx\_pygraphviz, nx\_pydot

to\_networkx\_graph(data[, create\_using, ...]) Make a NetworkX graph from a known data structure.

## 8.1.3 to networkx graph

 $\verb"to_networkx_graph" (data, create\_using=None, multigraph\_input=False)$ 

Make a NetworkX graph from a known data structure.

The preferred way to call this is automatically from the class constructor

```
>>> d={0: {1: {'weight':1}}} # dict-of-dicts single edge (0,1)
>>> G=nx.Graph(d)
instead of the equivalent
>>> G=nx.from_dict_of_dicts(d)
```

Parameters data: a object to be converted

**Current known types are:** any NetworkX graph dict-of-dicts dist-of-lists list of edges numpy matrix numpy ndarray scipy sparse matrix pygraphviz agraph

create\_using : NetworkX graph

Use specified graph for result. Otherwise a new graph is created.

```
multigraph_input : bool (default False)
```

If True and data is a dict\_of\_dicts, try to create a multigraph assuming dict\_of\_dict\_of\_lists. If data and create\_using are both multigraphs then create a multigraph from a multigraph.

## 8.2 Dictionaries

to_dict_of_dicts(G[, nodelist, edge_data])	Return adjacency representation of graph as a dictionary of dictionaries.
<pre>from_dict_of_dicts(d[, create_using,])</pre>	Return a graph from a dictionary of dictionaries.

## 8.2.1 to dict of dicts

```
to_dict_of_dicts (G, nodelist=None, edge_data=None)
```

Return adjacency representation of graph as a dictionary of dictionaries.

Parameters G: graph

A NetworkX graph

nodelist : list

Use only nodes specified in nodelist

edge\_data: list, optional

If provided, the value of the dictionary will be set to edge\_data for all edges. This is useful to make an adjacency matrix type representation with 1 as the edge data. If edgedata is None, the edgedata in G is used to fill the values. If G is a multigraph, the edgedata is a dict for each pair (u,v).

## 8.2.2 from dict of dicts

from\_dict\_of\_dicts (d, create\_using=None, multigraph\_input=False)

Return a graph from a dictionary of dictionaries.

**Parameters d**: dictionary of dictionaries

A dictionary of dictionaries adjacency representation.

create\_using : NetworkX graph

Use specified graph for result. Otherwise a new graph is created.

multigraph input: bool (default False)

When True, the values of the inner dict are assumed to be containers of edge data for multiple edges. Otherwise this routine assumes the edge data are singletons.

#### **Examples**

```
>>> dod= {0: {1:{'weight':1}}} # single edge (0,1)
>>> G=nx.from_dict_of_dicts(dod)
```

or >>> G=nx.Graph(dod) # use Graph constructor

## 8.3 Lists

<pre>to_dict_of_lists(G[, nodelist])</pre>	Return adjacency representation of graph as a dictionary of lists.
<pre>from_dict_of_lists(d[, create_using])</pre>	Return a graph from a dictionary of lists.
$to\_edgelist(G[, nodelist])$	Return a list of edges in the graph.
<pre>from_edgelist(edgelist[, create_using])</pre>	Return a graph from a list of edges.

## 8.3.1 to\_dict\_of\_lists

to\_dict\_of\_lists(G, nodelist=None)

Return adjacency representation of graph as a dictionary of lists.

Parameters G: graph

A NetworkX graph

nodelist: list

Use only nodes specified in nodelist

#### **Notes**

Completely ignores edge data for MultiGraph and MultiDiGraph.

## 8.3.2 from\_dict\_of\_lists

from\_dict\_of\_lists(d, create\_using=None)

Return a graph from a dictionary of lists.

**Parameters d**: dictionary of lists

A dictionary of lists adjacency representation.

create\_using : NetworkX graph

Use specified graph for result. Otherwise a new graph is created.

### **Examples**

```
>>> dol= {0:[1]} # single edge (0,1)
>>> G=nx.from_dict_of_lists(dol)
```

or >>> G=nx.Graph(dol) # use Graph constructor

## 8.3.3 to edgelist

to\_edgelist(G, nodelist=None)

Return a list of edges in the graph.

Parameters G: graph

8.3. Lists 413

A NetworkX graph

nodelist: list

Use only nodes specified in nodelist

## 8.3.4 from\_edgelist

```
from_edgelist (edgelist, create_using=None)
```

Return a graph from a list of edges.

Parameters edgelist: list or iterator

Edge tuples

create\_using : NetworkX graph

Use specified graph for result. Otherwise a new graph is created.

#### **Examples**

```
>>> edgelist= [(0,1)] # single edge (0,1)
>>> G=nx.from_edgelist(edgelist)
```

or >>> G=nx.Graph(edgelist) # use Graph constructor

# 8.4 Numpy

Functions to convert NetworkX graphs to and from numpy/scipy matrices.

The preferred way of converting data to a NetworkX graph is through the graph constuctor. The constructor calls the to\_networkx\_graph() function which attempts to guess the input type and convert it automatically.

## 8.4.1 Examples

Create a 10 node random graph from a numpy matrix

```
>>> import numpy
>>> a = numpy.reshape(numpy.random.random_integers(0,1,size=100),(10,10))
>>> D = nx.DiGraph(a)

or equivalently
>>> D = nx.to_networkx_graph(a,create_using=nx.DiGraph())
```

## 8.4.2 See Also

nx\_pygraphviz, nx\_pydot

to_numpy_matrix(G[, nodelist, dtype, order,])	Return the graph adjacency matrix as a NumPy matrix.
to_numpy_recarray(G[, nodelist, dtype, order])	Return the graph adjacency matrix as a NumPy recarray.
from_numpy_matrix(A[, parallel_edges,])	Return a graph from numpy matrix.

## 8.4.3 to numpy matrix

 $\label{local_constraint} \begin{tabular}{ll} \textbf{to\_numpy\_matrix} (G, nodelist=None, dtype=None, order=None, multigraph\_weight=<br/>built-in function $sum>$, weight='weight', nonedge=0.0) \end{tabular}$ 

Return the graph adjacency matrix as a NumPy matrix.

Parameters G: graph

The NetworkX graph used to construct the NumPy matrix.

nodelist: list, optional

The rows and columns are ordered according to the nodes in *nodelist*. If *nodelist* is None, then the ordering is produced by G.nodes().

dtype: NumPy data type, optional

A valid single NumPy data type used to initialize the array. This must be a simple type such as int or numpy.float64 and not a compound data type (see to\_numpy\_recarray) If None, then the NumPy default is used.

order: {'C', 'F'}, optional

Whether to store multidimensional data in C- or Fortran-contiguous (row- or columnwise) order in memory. If None, then the NumPy default is used.

multigraph\_weight : {sum, min, max}, optional

An operator that determines how weights in multigraphs are handled. The default is to sum the weights of the multiple edges.

weight: string or None optional (default='weight')

The edge attribute that holds the numerical value used for the edge weight. If an edge does not have that attribute, then the value 1 is used instead.

**nonedge**: float (default=0.0)

The matrix values corresponding to nonedges are typically set to zero. However, this could be undesirable if there are matrix values corresponding to actual edges that also have the value zero. If so, one might prefer nonedges to have some other value, such as nan.

**Returns** M: NumPy matrix

Graph adjacency matrix

## See also:

to\_numpy\_recarray, from\_numpy\_matrix

#### **Notes**

The matrix entries are assigned to the weight edge attribute. When an edge does not have a weight attribute, the value of the entry is set to the number 1. For multiple (parallel) edges, the values of the entries are determined by the 'multigraph\_weight' parameter. The default is to sum the weight attributes for each of the parallel edges.

When nodelist does not contain every node in G, the matrix is built from the subgraph of G that is induced by the nodes in nodelist.

The convention used for self-loop edges in graphs is to assign the diagonal matrix entry value to the weight attribute of the edge (or the number 1 if the edge has no weight attribute). If the alternate convention of doubling the edge weight is desired the resulting Numpy matrix can be modified as follows:

8.4. Numpy 415

```
>>> import numpy as np
>>> G = nx.Graph([(1,1)])
>>> A = nx.to_numpy_matrix(G)
>>> A
matrix([[ 1.]])
>>> A.A[np.diag_indices_from(A)] *= 2
>>> A
matrix([[ 2.]])
```

### **Examples**

## 8.4.4 to numpy recarray

to\_numpy\_recarray (*G*, nodelist=None, dtype=[('weight', <type 'float'>)], order=None)
Return the graph adjacency matrix as a NumPy recarray.

### Parameters G: graph

The NetworkX graph used to construct the NumPy matrix.

### nodelist: list, optional

The rows and columns are ordered according to the nodes in *nodelist*. If *nodelist* is None, then the ordering is produced by G.nodes().

### dtype: NumPy data-type, optional

A valid NumPy named dtype used to initialize the NumPy recarray. The data type names are assumed to be keys in the graph edge attribute dictionary.

```
order: {'C', 'F'}, optional
```

Whether to store multidimensional data in C- or Fortran-contiguous (row- or columnwise) order in memory. If None, then the NumPy default is used.

### **Returns** M : NumPy recarray

The graph with specified edge data as a Numpy recarray

### **Notes**

When nodelist does not contain every node in G, the matrix is built from the subgraph of G that is induced by the nodes in nodelist.

### **Examples**

```
>>> G = nx.Graph()
>>> G.add_edge(1,2,weight=7.0,cost=5)
>>> A=nx.to_numpy_recarray(G,dtype=[('weight',float),('cost',int)])
>>> print(A.weight)
[[ 0.    7.]
   [ 7.   0.]]
>>> print(A.cost)
[[ 0   5]
   [ 5   0]]
```

## 8.4.5 from\_numpy\_matrix

from\_numpy\_matrix (A, parallel\_edges=False, create\_using=None)

Return a graph from numpy matrix.

The numpy matrix is interpreted as an adjacency matrix for the graph.

**Parameters A**: numpy matrix

An adjacency matrix representation of a graph

parallel\_edges: Boolean

If this is True,  $create_u sing$  is a multigraph, and A is an integer matrix, then entry (i, j) in the matrix is interpreted as the number of parallel edges joining vertices i and j in the graph. If it is False, then the entries in the adjacency matrix are interpreted as the weight of a single edge joining the vertices.

create\_using : NetworkX graph

Use specified graph for result. The default is Graph()

### See also:

```
to_numpy_matrix, to_numpy_recarray
```

## **Notes**

If  $create_u sing$  is an instance of networkx. MultiGraph or networkx. MultiDiGraph,  $parallel_e dges$  is True, and the entries of A are of type int, then this function returns a multigraph (of the same type as  $create_u sing$ ) with parallel edges.

If  $create_u sing$  is an undirected multigraph, then only the edges indicated by the upper triangle of the matrix A will be added to the graph.

If the numpy matrix has a single data type for each matrix entry it will be converted to an appropriate Python data type.

If the numpy matrix has a user-specified compound data type the names of the data fields will be used as attribute keys in the resulting NetworkX graph.

#### **Examples**

Simple integer weights on edges:

8.4. Numpy 417

```
>>> import numpy
>>> A=numpy.matrix([[1,1],[2,1]])
>>> G=nx.from_numpy_matrix(A)
```

If  $create_u sing$  is a multigraph and the matrix has only integer entries, the entries will be interpreted as weighted edges joining the vertices (without creating parallel edges):

```
>>> import numpy
>>> A = numpy.matrix([[1, 1], [1, 2]])
>>> G = nx.from_numpy_matrix(A, create_using=nx.MultiGraph())
>>> G[1][1]
{0: {'weight': 2}}
```

If  $create_u sing$  is a multigraph and the matrix has only integer entries but  $parallel_e dges$  is True, then the entries will be interpreted as the number of parallel edges joining those two vertices:

```
>>> import numpy
>>> A = numpy.matrix([[1, 1], [1, 2]])
>>> temp = nx.MultiGraph()
>>> G = nx.from_numpy_matrix(A, parallel_edges=True, create_using=temp)
>>> G[1][1]
{0: {'weight': 1}, 1: {'weight': 1}}
```

User defined compound data type on edges:

```
>>> import numpy
>>> dt=[('weight',float),('cost',int)]
>>> A=numpy.matrix([[(1.0,2)]],dtype=dt)
>>> G=nx.from_numpy_matrix(A)
>>> G.edges()
[(0, 0)]
>>> G[0][0]['cost']
2
>>> G[0][0]['weight']
1.0
```

# 8.5 Scipy

```
to_scipy_sparse_matrix(G[, nodelist, dtype, ...]) Return the graph adjacency matrix as a SciPy sparse matrix.

from_scipy_sparse_matrix(A[, ...]) Creates a new graph from an adjacency matrix given as a SciPy sparse matrix.
```

## 8.5.1 to\_scipy\_sparse\_matrix

```
to_scipy_sparse_matrix (G, nodelist=None, dtype=None, weight='weight', format='csr')
Return the graph adjacency matrix as a SciPy sparse matrix.
```

Parameters G: graph

The NetworkX graph used to construct the NumPy matrix.

nodelist: list, optional

The rows and columns are ordered according to the nodes in *nodelist*. If *nodelist* is None, then the ordering is produced by G.nodes().

dtype: NumPy data-type, optional

A valid NumPy dtype used to initialize the array. If None, then the NumPy default is used.

```
weight: string or None optional (default='weight')
```

The edge attribute that holds the numerical value used for the edge weight. If None then all edge weights are 1.

```
format: str in {'bsr', 'csr', 'csc', 'coo', 'lil', 'dia', 'dok'}
```

The type of the matrix to be returned (default 'csr'). For some algorithms different implementations of sparse matrices can perform better. See [R318] for details.

### **Returns** M : SciPy sparse matrix

Graph adjacency matrix.

#### **Notes**

The matrix entries are populated using the edge attribute held in parameter weight. When an edge does not have that attribute, the value of the entry is 1.

For multiple edges the matrix values are the sums of the edge weights.

When nodelist does not contain every node in G, the matrix is built from the subgraph of G that is induced by the nodes in nodelist.

Uses coo\_matrix format. To convert to other formats specify the format= keyword.

The convention used for self-loop edges in graphs is to assign the diagonal matrix entry value to the weight attribute of the edge (or the number 1 if the edge has no weight attribute). If the alternate convention of doubling the edge weight is desired the resulting Scipy sparse matrix can be modified as follows:

```
>>> import scipy as sp
>>> G = nx.Graph([(1,1)])
>>> A = nx.to_scipy_sparse_matrix(G)
>>> print(A.todense())
[[1]]
>>> A.setdiag(A.diagonal()*2)
>>> print(A.todense())
[[2]]
```

### References

[R318]

### **Examples**

```
>>> G = nx.MultiDiGraph()
>>> G.add_edge(0,1,weight=2)
>>> G.add_edge(1,0)
>>> G.add_edge(2,2,weight=3)
>>> G.add_edge(2,2)
>>> S = nx.to_scipy_sparse_matrix(G, nodelist=[0,1,2])
>>> print(S.todense())
[[0 2 0]
  [1 0 0]
  [0 0 4]]
```

8.5. Scipy 419

## 8.5.2 from scipy sparse matrix

**from\_scipy\_sparse\_matrix** (A, parallel\_edges=False, create\_using=None, edge\_attribute='weight')

Creates a new graph from an adjacency matrix given as a SciPy sparse matrix.

### Parameters A: scipy sparse matrix

An adjacency matrix representation of a graph

```
parallel_edges: Boolean
```

If this is True,  $create_u sing$  is a multigraph, and A is an integer matrix, then entry (i, j) in the matrix is interpreted as the number of parallel edges joining vertices i and j in the graph. If it is False, then the entries in the adjacency matrix are interpreted as the weight of a single edge joining the vertices.

### create\_using: NetworkX graph

Use specified graph for result. The default is Graph()

#### edge\_attribute: string

Name of edge attribute to store matrix numeric value. The data will have the same type as the matrix entry (int, float, (real,imag)).

#### **Notes**

If  $create_u sing$  is an instance of networkx. MultiGraph or networkx. MultiDiGraph,  $parallel_e dges$  is True, and the entries of A are of type int, then this function returns a multigraph (of the same type as  $create_u sing$ ) with parallel edges. In this case,  $edge_a ttribute$  will be ignored.

If  $create_u sing$  is an undirected multigraph, then only the edges indicated by the upper triangle of the matrix A will be added to the graph.

#### **Examples**

```
>>> import scipy.sparse
>>> A = scipy.sparse.eye(2,2,1)
>>> G = nx.from_scipy_sparse_matrix(A)
```

If  $create_u sing$  is a multigraph and the matrix has only integer entries, the entries will be interpreted as weighted edges joining the vertices (without creating parallel edges):

```
>>> import scipy
>>> A = scipy.sparse.csr_matrix([[1, 1], [1, 2]])
>>> G = nx.from_scipy_sparse_matrix(A, create_using=nx.MultiGraph())
>>> G[1][1]
{0: {'weight': 2}}
```

If  $create_u sing$  is a multigraph and the matrix has only integer entries but  $parallel_e dges$  is True, then the entries will be interpreted as the number of parallel edges joining those two vertices:

```
>>> import scipy
>>> A = scipy.sparse.csr_matrix([[1, 1], [1, 2]])
>>> G = nx.from_scipy_sparse_matrix(A, parallel_edges=True,
```

```
create_using=nx.MultiGraph())
>>> G[1][1]
{0: {'weight': 1}, 1: {'weight': 1}}
```

## 8.6 Pandas

to_pandas_dataframe( $G[$ , nodelist,])	Return the graph adjacency matrix as a Pandas DataFrame.
<pre>from_pandas_dataframe(df[, create_using])</pre>	Return a graph from Pandas DataFrame.

## 8.6.1 to pandas dataframe

 $\begin{tabular}{ll} \textbf{to\_pandas\_dataframe} (G, & nodelist=None, & multigraph\_weight=<built-in & function & sum>, \\ & weight='weight', nonedge=0.0) \\ & \textbf{Return the graph adjacency matrix as a Pandas DataFrame}. \\ \end{tabular}$ 

Parameters G: graph

The NetworkX graph used to construct the Pandas DataFrame.

nodelist: list, optional

The rows and columns are ordered according to the nodes in *nodelist*. If *nodelist* is None, then the ordering is produced by G.nodes().

multigraph\_weight: {sum, min, max}, optional

An operator that determines how weights in multigraphs are handled. The default is to sum the weights of the multiple edges.

weight: string or None, optional

The edge attribute that holds the numerical value used for the edge weight. If an edge does not have that attribute, then the value 1 is used instead.

nonedge: float, optional

The matrix values corresponding to nonedges are typically set to zero. However, this could be undesirable if there are matrix values corresponding to actual edges that also have the value zero. If so, one might prefer nonedges to have some other value, such as nan.

Returns df: Pandas DataFrame

Graph adjacency matrix

#### **Notes**

The DataFrame entries are assigned to the weight edge attribute. When an edge does not have a weight attribute, the value of the entry is set to the number 1. For multiple (parallel) edges, the values of the entries are determined by the 'multigraph\_weight' parameter. The default is to sum the weight attributes for each of the parallel edges.

When nodelist does not contain every node in G, the matrix is built from the subgraph of G that is induced by the nodes in nodelist.

8.6. Pandas 421

The convention used for self-loop edges in graphs is to assign the diagonal matrix entry value to the weight attribute of the edge (or the number 1 if the edge has no weight attribute). If the alternate convention of doubling the edge weight is desired the resulting Pandas DataFrame can be modified as follows:

### **Examples**

## 8.6.2 from pandas dataframe

```
from_pandas_dataframe (df, create_using=None)
```

Return a graph from Pandas DataFrame.

The Pandas DataFrame is interpreted as an adjacency matrix for the graph.

Parameters df: Pandas DataFrame

An adjacency matrix representation of a graph

create\_using : NetworkX graph

Use specified graph for result. The default is Graph()

#### See also:

```
from_pandas_dataframe
```

### **Notes**

If the numpy matrix has a single data type for each matrix entry it will be converted to an appropriate Python data type.

If the numpy matrix has a user-specified compound data type the names of the data fields will be used as attribute keys in the resulting NetworkX graph.

## **Examples**

Simple integer weights on edges:

```
>>> import pandas as pd
>>> df=pd.DataFrame([[1,1],[2,1]])
>>> G=nx.from_pandas_dataframe(df)
```

8.6. Pandas 423

NetworkX Reference, Release 2.0.dev2015	0317000012	
424	Chapter 9	Converting to and from other data formats

**CHAPTER** 

**NINE** 

## READING AND WRITING GRAPHS

# 9.1 Adjacency List

Read and write NetworkX graphs as adjacency lists.

Adjacency list format is useful for graphs without data associated with nodes or edges and for nodes that can be meaningfully represented as strings.

## **9.1.1 Format**

The adjacency list format consists of lines with node labels. The first label in a line is the source node. Further labels in the line are considered target nodes and are added to the graph along with an edge between the source node and target node.

The graph with edges a-b, a-c, d-e can be represented as the following adjacency list (anything following the # in a line is a comment):

```
a b c # source target target
d e
```

<pre>read_adjlist(path[, comments, delimiter,])</pre>	Read graph in adjacency list format from path.
<pre>write_adjlist(G, path[, comments,])</pre>	Write graph G in single-line adjacency-list format to path.
<pre>parse_adjlist(lines[, comments, delimiter,])</pre>	Parse lines of a graph adjacency list representation.
<pre>generate_adjlist(G[, delimiter])</pre>	Generate a single line of the graph G in adjacency list format.

## 9.1.2 read adjlist

read\_adjlist (path, comments='#', delimiter=None, create\_using=None, nodetype=None, encoding='utf-8')

Read graph in adjacency list format from path.

Parameters path: string or file

Filename or file handle to read. Filenames ending in .gz or .bz2 will be uncompressed.

create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

nodetype: Python type, optionalConvert nodes to this type.comments: string, optional

Marker for comment lines

delimiter: string, optional

Separator for node labels. The default is whitespace.

## create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

Returns G: NetworkX graph

The graph corresponding to the lines in adjacency list format.

#### See also:

```
write_adjlist
```

#### **Notes**

This format does not store graph or node data.

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_adjlist(G, "test.adjlist")
>>> G=nx.read_adjlist("test.adjlist")
```

The path can be a filehandle or a string with the name of the file. If a filehandle is provided, it has to be opened in 'rb' mode.

```
>>> fh=open("test.adjlist", 'rb')
>>> G=nx.read_adjlist(fh)
```

Filenames ending in .gz or .bz2 will be compressed.

```
>>> nx.write_adjlist(G,"test.adjlist.gz")
>>> G=nx.read_adjlist("test.adjlist.gz")
```

The optional nodetype is a function to convert node strings to nodetype.

For example

```
>>> G=nx.read_adjlist("test.adjlist", nodetype=int)
```

will attempt to convert all nodes to integer type.

Since nodes must be hashable, the function nodetype must return hashable types (e.g. int, float, str, frozenset - or tuples of those, etc.)

The optional create\_using parameter is a NetworkX graph container. The default is Graph(), an undirected graph. To read the data as a directed graph use

```
>>> G=nx.read_adjlist("test.adjlist", create_using=nx.DiGraph())
```

## 9.1.3 write adjlist

```
write_adjlist (G, path, comments='#', delimiter=' ', encoding='utf-8')
Write graph G in single-line adjacency-list format to path.
```

```
Parameters G : NetworkX graph
    path : string or file
        Filename or file handle for data output. Filenames ending in .gz or .bz2 will be compressed.
    comments : string, optional
```

**delimiter**: string, optional

Separator for node labels

Marker for comment lines

encoding : string, optional

Text encoding.

#### See also:

```
read_adjlist, generate_adjlist
```

#### **Notes**

This format does not store graph, node, or edge data.

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_adjlist(G,"test.adjlist")
```

The path can be a filehandle or a string with the name of the file. If a filehandle is provided, it has to be opened in 'wb' mode.

```
>>> fh=open("test.adjlist",'wb')
>>> nx.write_adjlist(G, fh)
```

## 9.1.4 parse adjlist

```
parse_adjlist (lines, comments='#', delimiter=None, create_using=None, nodetype=None)
Parse lines of a graph adjacency list representation.
```

Parameters lines: list or iterator of strings

Input data in adjlist format

create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

**nodetype**: Python type, optional Convert nodes to this type.

comments : string, optional
 Marker for comment lines
delimiter : string, optional

Separator for node labels. The default is whitespace.

9.1. Adjacency List

### create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

## Returns G: NetworkX graph

The graph corresponding to the lines in adjacency list format.

#### See also:

```
read_adjlist
```

### **Examples**

## 9.1.5 generate\_adjlist

```
generate_adjlist(G, delimiter=' ')
```

Generate a single line of the graph G in adjacency list format.

Parameters G: NetworkX graph

delimiter: string, optional

Separator for node labels

**Returns lines**: string

Lines of data in adjlist format.

#### See also:

```
write_adjlist, read_adjlist
```

### **Examples**

# 9.2 Multiline Adjacency List

Read and write NetworkX graphs as multi-line adjacency lists.

The multi-line adjacency list format is useful for graphs with nodes that can be meaningfully represented as strings. With this format simple edge data can be stored but node or graph data is not.

### **9.2.1 Format**

The first label in a line is the source node label followed by the node degree d. The next d lines are target node labels and optional edge data. That pattern repeats for all nodes in the graph.

The graph with edges a-b, a-c, d-e can be represented as the following adjacency list (anything following the # in a line is a comment):

```
# example.multiline-adjlist
a 2
b
c
d 1
```

read_multiline_adjlist(path[, comments,])	Read graph in multi-line adjacency list format from path.
write_multiline_adjlist(G, path[,])	Write the graph G in multiline adjacency list format to path
<pre>parse_multiline_adjlist(lines[, comments,])</pre>	Parse lines of a multiline adjacency list representation of a graph.
<pre>generate_multiline_adjlist(G[, delimiter])</pre>	Generate a single line of the graph G in multiline adjacency list format.

## 9.2.2 read\_multiline\_adjlist

```
read_multiline_adjlist (path, comments='#', delimiter=None, create_using=None, nodetype=None, edgetype=None, encoding='utf-8')

Read graph in multi-line adjacency list format from path.
```

**Parameters path**: string or file

Filename or file handle to read. Filenames ending in .gz or .bz2 will be uncompressed.

## create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

**nodetype**: Python type, optional Convert nodes to this type.

**edgetype**: Python type, optional Convert edge data to this type.

comments : string, optional
 Marker for comment lines
delimiter : string, optional

Separator for node labels. The default is whitespace.

## create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

### Returns G: NetworkX graph

#### See also:

```
write_multiline_adjlist
```

#### **Notes**

This format does not store graph, node, or edge data.

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_multiline_adjlist(G,"test.adjlist")
>>> G=nx.read_multiline_adjlist("test.adjlist")
```

The path can be a file or a string with the name of the file. If a file s provided, it has to be opened in 'rb' mode.

```
>>> fh=open("test.adjlist", 'rb')
>>> G=nx.read_multiline_adjlist(fh)
```

Filenames ending in .gz or .bz2 will be compressed.

```
>>> nx.write_multiline_adjlist(G,"test.adjlist.gz")
>>> G=nx.read_multiline_adjlist("test.adjlist.gz")
```

The optional nodetype is a function to convert node strings to nodetype.

For example

```
>>> G=nx.read_multiline_adjlist("test.adjlist", nodetype=int)
```

will attempt to convert all nodes to integer type.

The optional edgetype is a function to convert edge data strings to edgetype.

```
>>> G=nx.read_multiline_adjlist("test.adjlist")
```

The optional create\_using parameter is a NetworkX graph container. The default is Graph(), an undirected graph. To read the data as a directed graph use

```
>>> G=nx.read_multiline_adjlist("test.adjlist", create_using=nx.DiGraph())
```

## 9.2.3 write\_multiline\_adjlist

```
write_multiline_adjlist (G, path, delimiter=' ', comments='#', encoding='utf-8')
Write the graph G in multiline adjacency list format to path
```

```
Parameters G: NetworkX graph
comments: string, optional
Marker for comment lines
delimiter: string, optional
Separator for node labels
encoding: string, optional
```

Text encoding.

### See also:

```
read_multiline_adjlist
```

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_multiline_adjlist(G,"test.adjlist")
```

The path can be a file handle or a string with the name of the file. If a file handle is provided, it has to be opened in 'wb' mode.

```
>>> fh=open("test.adjlist",'wb')
>>> nx.write_multiline_adjlist(G,fh)
```

Filenames ending in .gz or .bz2 will be compressed.

```
>>> nx.write_multiline_adjlist(G,"test.adjlist.gz")
```

## 9.2.4 parse multiline adjlist

```
parse_multiline_adjlist(lines, comments='#', delimiter=None, create_using=None, node-
type=None, edgetype=None)
```

Parse lines of a multiline adjacency list representation of a graph.

Parameters lines: list or iterator of strings

Input data in multiline adjlist format

### create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

nodetype: Python type, optional

Convert nodes to this type.

comments: string, optional

Marker for comment lines

delimiter: string, optional

Separator for node labels. The default is whitespace.

## create\_using: NetworkX graph container

Use given NetworkX graph for holding nodes or edges.

Returns G: NetworkX graph

The graph corresponding to the lines in multiline adjacency list format.

### **Examples**

## 9.2.5 generate\_multiline\_adjlist

```
\begin{tabular}{ll} \beg
```

Parameters G: NetworkX graph delimiter: string, optional Separator for node labels

Returns lines: string

Lines of data in multiline adjlist format.

### See also:

```
write_multiline_adjlist, read_multiline_adjlist
```

### **Examples**

```
>>> G = nx.lollipop_graph(4, 3)
>>> for line in nx.generate_multiline_adjlist(G):
        print(line)
0 3
1 {}
2 {}
3 {}
1 2
2 {}
3 {}
2 1
3 {}
3 1
4 {}
4 1
5 {}
5 1
6 {}
6 0
```

# 9.3 Edge List

Read and write NetworkX graphs as edge lists.

The multi-line adjacency list format is useful for graphs with nodes that can be meaningfully represented as strings. With the edgelist format simple edge data can be stored but node or graph data is not. There is no way of representing isolated nodes unless the node has a self-loop edge.

## **9.3.1 Format**

You can read or write three formats of edge lists with these functions.

Node pairs with no data:

1 2

Python dictionary as data:

```
1 2 {'weight':7, 'color':'green'}
```

## Arbitrary data:

1 2 7 green

read_edgelist(path[, comments, delimiter,])	Read a graph from a list of edges.
write_edgelist(G, path[, comments,])	Write graph as a list of edges.
<pre>read_weighted_edgelist(path[, comments,])</pre>	Read a graph as list of edges with numeric weights.
<pre>write_weighted_edgelist(G, path[, comments,])</pre>	Write graph G as a list of edges with numeric weights.
<pre>generate_edgelist(G[, delimiter, data])</pre>	Generate a single line of the graph G in edge list format.
<pre>parse_edgelist(lines[, comments, delimiter,])</pre>	Parse lines of an edge list representation of a graph.

## 9.3.2 read\_edgelist

 $\begin{tabular}{ll} \textbf{read\_edgelist} (path, comments='\#', delimiter=None, create\_using=None, nodetype=None, data=True, \\ edgetype=None, encoding='utf-8') \end{tabular}$ 

Read a graph from a list of edges.

Parameters path: file or string

File or filename to read. If a file is provided, it must be opened in 'rb' mode. Filenames ending in .gz or .bz2 will be uncompressed.

comments: string, optional

The character used to indicate the start of a comment.

delimiter: string, optional

The string used to separate values. The default is whitespace.

create\_using : Graph container, optional,

Use specified container to build graph. The default is networkx. Graph, an undirected graph.

nodetype: int, float, str, Python type, optional

Convert node data from strings to specified type

data: bool or list of (label,type) tuples

Tuples specifying dictionary key names and types for edge data

edgetype: int, float, str, Python type, optional OBSOLETE

9.3. Edge List 433

Convert edge data from strings to specified type and use as 'weight'

### encoding: string, optional

Specify which encoding to use when reading file.

## Returns G: graph

A networkx Graph or other type specified with create using

#### See also:

```
parse_edgelist
```

#### **Notes**

Since nodes must be hashable, the function nodetype must return hashable types (e.g. int, float, str, frozenset - or tuples of those, etc.)

#### **Examples**

```
>>> nx.write_edgelist(nx.path_graph(4), "test.edgelist")
>>> G=nx.read_edgelist("test.edgelist")
>>> fh=open("test.edgelist", 'rb')
>>> G=nx.read_edgelist(fh)
>>> fh.close()
>>> G=nx.read_edgelist("test.edgelist", nodetype=int)
>>> G=nx.read_edgelist("test.edgelist",create_using=nx.DiGraph())
Edgelist with data in a list:
>>> textline = '1 2 3'
>>> fh = open('test.edgelist','w')
>>> d = fh.write(textline)
>>> fh.close()
>>> G = nx.read_edgelist('test.edgelist', nodetype=int, data=(('weight',float),))
>>> G.nodes()
[1, 2]
>>> G.edges(data = True)
[(1, 2, {'weight': 3.0})]
```

See parse\_edgelist() for more examples of formatting.

## 9.3.3 write\_edgelist

```
write_edgelist (G, path, comments='\#', delimiter='', data=True, encoding='utf-8') Write graph as a list of edges.
```

```
Parameters G: graph
```

A NetworkX graph

path: file or string

File or filename to write. If a file is provided, it must be opened in 'wb' mode. Filenames ending in .gz or .bz2 will be compressed.

```
comments: string, optional
```

The character used to indicate the start of a comment

### delimiter: string, optional

The string used to separate values. The default is whitespace.

```
data: bool or list, optional
```

If False write no edge data. If True write a string representation of the edge data dictionary.. If a list (or other iterable) is provided, write the keys specified in the list.

## encoding: string, optional

Specify which encoding to use when writing file.

#### See also:

```
write_edgelist, write_weighted_edgelist
```

## **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_edgelist(G, "test.edgelist")
>>> G=nx.path_graph(4)
>>> fh=open("test.edgelist",'wb')
>>> nx.write_edgelist(G, fh)
>>> nx.write_edgelist(G, "test.edgelist.gz")
>>> nx.write_edgelist(G, "test.edgelist.gz", data=False)
>>> G=nx.Graph()
>>> G.add_edge(1,2,weight=7,color='red')
>>> nx.write_edgelist(G,'test.edgelist',data=False)
>>> nx.write_edgelist(G,'test.edgelist',data=['color'])
>>> nx.write_edgelist(G,'test.edgelist',data=['color','weight'])
```

## 9.3.4 read weighted edgelist

Read a graph as list of edges with numeric weights.

## Parameters path: file or string

File or filename to read. If a file is provided, it must be opened in 'rb' mode. Filenames ending in .gz or .bz2 will be uncompressed.

### comments: string, optional

The character used to indicate the start of a comment.

### **delimiter**: string, optional

The string used to separate values. The default is whitespace.

create\_using : Graph container, optional,

Use specified container to build graph. The default is networkx. Graph, an undirected graph.

nodetype: int, float, str, Python type, optional

9.3. Edge List 435

Convert node data from strings to specified type

```
encoding: string, optional
```

Specify which encoding to use when reading file.

Returns G: graph

A networkx Graph or other type specified with create using

#### **Notes**

Since nodes must be hashable, the function nodetype must return hashable types (e.g. int, float, str, frozenset - or tuples of those, etc.)

Example edgelist file format.

With numeric edge data:

```
# read with
# >>> G=nx.read_weighted_edgelist(fh)
# source target data
a b 1
a c 3.14159
d e 42
```

## 9.3.5 write\_weighted\_edgelist

```
write_weighted_edgelist (G, path, comments='#', delimiter=' ', encoding='utf-8')
Write graph G as a list of edges with numeric weights.
```

```
Parameters G: graph
```

A NetworkX graph

path: file or string

File or filename to write. If a file is provided, it must be opened in 'wb' mode. Filenames ending in .gz or .bz2 will be compressed.

comments: string, optional

The character used to indicate the start of a comment

delimiter: string, optional

The string used to separate values. The default is whitespace.

## encoding: string, optional

Specify which encoding to use when writing file.

#### See also:

```
read_edgelist, write_edgelist, write_weighted_edgelist
```

### **Examples**

```
>>> G=nx.Graph()
>>> G.add_edge(1,2,weight=7)
>>> nx.write_weighted_edgelist(G, 'test.weighted.edgelist')
```

## 9.3.6 generate edgelist

generate\_edgelist(G, delimiter=' ', data=True)

```
Generate a single line of the graph G in edge list format.
     Parameters G: NetworkX graph
         delimiter: string, optional
             Separator for node labels
         data: bool or list of keys
             If False generate no edge data. If True use a dictionary representation of edge data. If a
             list of keys use a list of data values corresponding to the keys.
     Returns lines: string
            Lines of data in adjlist format.
See also:
write_adjlist, read_adjlist
Examples
\rightarrow \rightarrow G = nx.lollipop_graph(4, 3)
>>> G[1][2]['weight'] = 3
>>> G[3][4]['capacity'] = 12
>>> for line in nx.generate_edgelist(G, data=False):
         print(line)
0 1
0 2
0 3
1 2
1 3
2 3
3 4
4 5
5 6
>>> for line in nx.generate_edgelist(G):
         print(line)
. . .
0 1 {}
0 2 {}
0 3 {}
1 2 {'weight': 3}
1 3 {}
2 3 {}
3 4 {'capacity': 12}
4 5 {}
5 6 {}
>>> for line in nx.generate_edgelist(G, data=['weight']):
         print(line)
0 1
0 2
```

9.3. Edge List 437

```
2 3
3 4
4 5
5 6
```

## 9.3.7 parse edgelist

```
parse_edgelist (lines, comments='#', delimiter=None, create_using=None, nodetype=None, data=True)
Parse lines of an edge list representation of a graph.

Parameters lines: list or iterator of strings
Input data in edgelist format
comments: string, optional
```

Marker for comment lines

delimiter: string, optional

Separator for node labels

## create\_using: NetworkX graph container, optional

Use given NetworkX graph for holding nodes or edges.

**nodetype**: Python type, optional Convert nodes to this type.

data: bool or list of (label,type) tuples

If False generate no edge data or if True use a dictionary representation of edge data or a list tuples specifying dictionary key names and types for edge data.

### Returns G: NetworkX Graph

The graph corresponding to lines

#### See also:

```
read weighted edgelist
```

## **Examples**

Edgelist with no data:

Edgelist with data in Python dictionary representation:

```
>>> lines = ["1 2 {'weight':3}",
           "2 3 {'weight':27}",
            "3 4 {'weight':3.0}"]
>>> G = nx.parse_edgelist(lines, nodetype = int)
>>> G.nodes()
[1, 2, 3, 4]
>>> G.edges(data = True)
[(1, 2, {'weight': 3}), (2, 3, {'weight': 27}), (3, 4, {'weight': 3.0})]
Edgelist with data in a list:
>>> lines = ["1 2 3",
             "2 3 27",
             "3 4 3.0"1
>>> G = nx.parse_edgelist(lines, nodetype = int, data=(('weight',float),))
>>> G.nodes()
[1, 2, 3, 4]
>>> G.edges(data = True)
[(1, 2, {'weight': 3.0}), (2, 3, {'weight': 27.0}), (3, 4, {'weight': 3.0})]
```

## **9.4 GEXF**

Read and write graphs in GEXF format.

GEXF (Graph Exchange XML Format) is a language for describing complex network structures, their associated data and dynamics.

This implementation does not support mixed graphs (directed and undirected edges together).

## **9.4.1 Format**

GEXF is an XML format. See <a href="http://gexf.net/format/schema.html">http://gexf.net/format/schema.html</a> for the specification and <a href="http://gexf.net/format/basic.html">http://gexf.net/format/basic.html</a> for examples.

<pre>read_gexf(path[, node_type, relabel, version])</pre>	Read graph in GEXF format from path.
<pre>write_gexf(G, path[, encoding, prettyprint,])</pre>	Write G in GEXF format to path.
relabel_gexf_graph(G)	Relabel graph using "label" node keyword for node label.

## 9.4.2 read gexf

```
read_gexf (path, node_type=None, relabel=False, version='1.1draft')
Read graph in GEXF format from path.
```

"GEXF (Graph Exchange XML Format) is a language for describing complex networks structures, their associated data and dynamics" [R365].

**Parameters path**: file or string

File or file name to write. File names ending in .gz or .bz2 will be compressed.

node\_type: Python type (default: None)

Convert node ids to this type if not None.

relabel: bool (default: False)

9.4. GEXF 439

If True relabel the nodes to use the GEXF node "label" attribute instead of the node "id" attribute as the NetworkX node label.

Returns graph: NetworkX graph

If no parallel edges are found a Graph or DiGraph is returned. Otherwise a MultiGraph or MultiDiGraph is returned.

#### **Notes**

This implementation does not support mixed graphs (directed and undirected edges together).

#### References

[R365]

## 9.4.3 write\_gexf

```
write_gexf (G, path, encoding='utf-8', prettyprint=True, version='1.1draft')
Write G in GEXF format to path.
```

"GEXF (Graph Exchange XML Format) is a language for describing complex networks structures, their associated data and dynamics" [R366].

```
Parameters G: graph

A NetworkX graph

path: file or string

File or file name to write. File names ending in .gz or .bz2 will be compressed.

encoding: string (optional)

Encoding for text data.

prettyprint: bool (optional)

If True use line breaks and indenting in output XML.
```

#### **Notes**

This implementation does not support mixed graphs (directed and undirected edges together).

The node id attribute is set to be the string of the node label. If you want to specify an id use set it as node data, e.g. node['a']['id']=1 to set the id of node 'a' to 1.

#### References

[R366]

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_gexf(G, "test.gexf")
```

## 9.4.4 relabel\_gexf\_graph

 $relabel_gexf_graph(G)$ 

Relabel graph using "label" node keyword for node label.

Parameters G: graph

A NetworkX graph read from GEXF data

Returns H: graph

A NetworkX graph with relabed nodes

#### **Notes**

This function relabels the nodes in a NetworkX graph with the "label" attribute. It also handles relabeling the specific GEXF node attributes "parents", and "pid".

## 9.5 GML

Read graphs in GML format.

"GML, the G>raph Modelling Language, is our proposal for a portable file format for graphs. GML's key features are portability, simple syntax, extensibility and flexibility. A GML file consists of a hierarchical key-value lists. Graphs can be annotated with arbitrary data structures. The idea for a common file format was born at the GD'95; this proposal is the outcome of many discussions. GML is the standard file format in the Graphlet graph editor system. It has been overtaken and adapted by several other systems for drawing graphs."

See http://www.infosun.fim.uni-passau.de/Graphlet/GML/gml-tr.html

## **9.5.1 Format**

See http://www.infosun.fim.uni-passau.de/Graphlet/GML/gml-tr.html for format specification.

Example graphs in GML format: http://www-personal.umich.edu/~mejn/netdata/

<pre>read_gml(path[, label, destringizer])</pre>	Read graph in GML format from path.
<pre>write_gml(G, path[, stringizer])</pre>	Write a graph G in GML format to the file or file handle path.
<pre>parse_gml(lines[, label, destringizer])</pre>	Parse GML graph from a string or iterable.
<pre>generate_gml(G[, stringizer])</pre>	Generate a single entry of the graph G in GML format.
literal_destringizer(rep)	Convert a Python literal to the value it represents.
literal_stringizer(value)	Convert a value to a Python literal in GML representation.

9.5. GML 441

## 9.5.2 read gml

```
read_gml (path, label='label', destringizer=None)
Read graph in GML format from path.
```

Parameters path: filename or filehandle

The filename or filehandle to read from.

label: string, optional

If not None, the parsed nodes will be renamed according to node attributes indicated by label. Default value: 'label'.

destringizer: callable, optional

A destringizer that recovers values stored as strings in GML. If it cannot convert a string to a value, a ValueError is raised. Default value: None.

Returns G: NetworkX graph

The parsed graph.

Raises NetworkXError

If the input cannot be parsed.

#### See also:

```
write_gml, parse_gml
```

#### **Notes**

The GML specification says that files should be ASCII encoded, with any extended ASCII characters (iso8859-1) appearing as HTML character entities.

#### References

GML specification: http://www.infosun.fim.uni-passau.de/Graphlet/GML/gml-tr.html

## **Examples**

```
>>> G = nx.path_graph(4)
>>> nx.write_gml(G, 'test.gml')
>>> H = nx.read_gml('test.gml')
```

## 9.5.3 write\_gml

write\_gml (G, path, stringizer=None)

Write a graph G in GML format to the file or file handle path.

Parameters G: NetworkX graph

The graph to be converted to GML.

path: filename or filehandle

The filename or filehandle to write. Files whose names end with .gz or .bz2 will be compressed.

stringizer: callable, optional

A stringizer which converts non-int/float/dict values into strings. If it cannot convert a value into a string, it should raise a ValueError raised to indicate that. Default value: None.

#### Raises NetworkXError

If stringizer cannot convert a value into a string, or the value to convert is not a string while stringizer is None.

#### See also:

```
read_gml, generate_gml
```

#### **Notes**

Graph attributes named 'directed', 'multigraph', 'node' or 'edge', node attributes named 'id' or 'label', edge attributes named 'source' or 'target' (or 'key' if G is a multigraph) are ignored because these attribute names are used to encode the graph structure.

### **Examples**

```
>>> G = nx.path_graph(4)
>>> nx.write_gml(G, "test.gml")
```

Filenames ending in .gz or .bz2 will be compressed.

```
>>> nx.write_gml(G, "test.gml.gz")
```

## 9.5.4 parse\_gml

```
parse_gml (lines, label='label', destringizer=None)
Parse GML graph from a string or iterable.
```

**Parameters lines**: string or iterable of strings

Data in GML format.

label: string, optional

If not None, the parsed nodes will be renamed according to node attributes indicated by label. Default value: 'label'.

destringizer: callable, optional

A destringizer that recovers values stored as strings in GML. If it cannot convert a string to a value, a ValueError is raised. Default value: None.

Returns G: NetworkX graph

The parsed graph.

### Raises NetworkXError

If the input cannot be parsed.

9.5. GML 443

#### See also:

```
write_gml, read_gml
```

#### **Notes**

This stores nested GML attributes as dictionaries in the NetworkX graph, node, and edge attribute structures.

#### References

GML specification: http://www.infosun.fim.uni-passau.de/Graphlet/GML/gml-tr.html

## 9.5.5 generate\_gml

### generate\_gml (G, stringizer=None)

Generate a single entry of the graph G in GML format.

Parameters G: NetworkX graph

The graph to be converted to GML.

stringizer: callable, optional

A stringizer which converts non-int/float/dict values into strings. If it cannot convert a value into a string, it should raise a ValueError raised to indicate that. Default value: None.

**Returns** lines: generator of strings

Lines of GML data. Newlines are not appended.

#### Raises NetworkXError

If stringizer cannot convert a value into a string, or the value to convert is not a string while stringizer is None.

## **Notes**

Graph attributes named 'directed', 'multigraph', 'node' or 'edge', node attributes named 'id' or 'label', edge attributes named 'source' or 'target' (or 'key' if G is a multigraph) are ignored because these attribute names are used to encode the graph structure.

# 9.5.6 literal\_destringizer

## ${\tt literal\_destringizer}\ (\textit{rep})$

Convert a Python literal to the value it represents.

**Parameters rep**: string A Python literal.

Returns value: object

The value of the Python literal.

Raises ValueError

If rep is not a Python literal.

## 9.5.7 literal\_stringizer

#### literal stringizer(value)

Convert a value to a Python literal in GML representation.

Parameters value: object

The value to be converted to GML representation.

**Returns** rep: string

A double-quoted Python literal representing value. Unprintable characters are replaced by XML character references.

#### Raises ValueError

If value cannot be converted to GML.

#### **Notes**

literal\_stringizer is largely the same as repr in terms of functionality but attempts prefix unicode and bytes literals with u and b to provide better interoperability of data generated by Python 2 and Python 3.

The original value can be recovered using the networkx.readwrite.gml.literal\_destringizer function.

## 9.6 Pickle

Read and write NetworkX graphs as Python pickles.

"The pickle module implements a fundamental, but powerful algorithm for serializing and de-serializing a Python object structure. "Pickling" is the process whereby a Python object hierarchy is converted into a byte stream, and "unpickling" is the inverse operation, whereby a byte stream is converted back into an object hierarchy."

Note that NetworkX graphs can contain any hashable Python object as node (not just integers and strings). For arbitrary data types it may be difficult to represent the data as text. In that case using Python pickles to store the graph data can be used.

## 9.6.1 Format

See http://docs.python.org/library/pickle.html

read_gpickle(path)	Read graph object in Python pickle format.
<pre>write_gpickle(G, path[, protocol])</pre>	Write graph in Python pickle format.

## 9.6.2 read gpickle

## read\_gpickle(path)

Read graph object in Python pickle format.

Pickles are a serialized byte stream of a Python object [R367]. This format will preserve Python objects used as

9.6. Pickle 445

nodes or edges.

Parameters path: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be uncompressed.

Returns G: graph

A NetworkX graph

### References

[R367]

### **Examples**

```
>>> G = nx.path_graph(4)
>>> nx.write_gpickle(G, "test.gpickle")
>>> G = nx.read_gpickle("test.gpickle")
```

## 9.6.3 write\_gpickle

```
write_gpickle (G, path, protocol=2)
```

Write graph in Python pickle format.

Pickles are a serialized byte stream of a Python object [R368]. This format will preserve Python objects used as nodes or edges.

```
Parameters G: graph
```

A NetworkX graph

path: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be compressed.

protocol: integer

Pickling protocol to use. Default value: pickle.HIGHEST\_PROTOCOL.

#### References

[R368]

## **Examples**

```
>>> G = nx.path_graph(4)
>>> nx.write_gpickle(G, "test.gpickle")
```

# 9.7 GraphML

Read and write graphs in GraphML format.

This implementation does not support mixed graphs (directed and unidirected edges together), hyperedges, nested graphs, or ports.

"GraphML is a comprehensive and easy-to-use file format for graphs. It consists of a language core to describe the structural properties of a graph and a flexible extension mechanism to add application-specific data. Its main features include support of

- · directed, undirected, and mixed graphs,
- · hypergraphs,
- · hierarchical graphs,
- · graphical representations,
- · references to external data,
- · application-specific attribute data, and
- light-weight parsers.

Unlike many other file formats for graphs, GraphML does not use a custom syntax. Instead, it is based on XML and hence ideally suited as a common denominator for all kinds of services generating, archiving, or processing graphs."

http://graphml.graphdrawing.org/

### **9.7.1 Format**

GraphML is an XML format. See <a href="http://graphml.graphdrawing.org/specification.html">http://graphml.graphdrawing.org/specification.html</a> for the specification and <a href="http://graphml.graphdrawing.org/primer/graphml-primer.html">http://graphml.graphdrawing.org/primer/graphml-primer.html</a> for examples.

read_graphml(path[, node_type])	Read graph in GraphML format from path.
<pre>write_graphml(G, path[, encoding, prettyprint])</pre>	Write G in GraphML XML format to path

## 9.7.2 read\_graphml

**read\_graphml** (*path*, *node\_type*=<*type* '*str*'>)
Read graph in GraphML format from path.

**Parameters path**: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be compressed.

node\_type: Python type (default: str)

Convert node ids to this type

**Returns** graph: NetworkX graph

If no parallel edges are found a Graph or DiGraph is returned. Otherwise a MultiGraph or MultiDiGraph is returned.

9.7. GraphML 447

#### **Notes**

This implementation does not support mixed graphs (directed and unidirected edges together), hypergraphs, nested graphs, or ports.

For multigraphs the GraphML edge "id" will be used as the edge key. If not specified then they "key" attribute will be used. If there is no "key" attribute a default NetworkX multigraph edge key will be provided.

Files with the yEd "yfiles" extension will can be read but the graphics information is discarded.

yEd compressed files ("file.graphmlz" extension) can be read by renaming the file to "file.graphml.gz".

## 9.7.3 write graphml

```
write_graphml (G, path, encoding='utf-8', prettyprint=True)
    Write G in GraphML XML format to path
    Parameters G: graph
        A networkx graph
    path: file or string
        File or filename to write. Filenames ending in .gz or .bz2 will be compressed.
    encoding: string (optional)
        Encoding for text data.
    prettyprint: bool (optional)
        If True use line breaks and indenting in output XML.
```

## **Notes**

This implementation does not support mixed graphs (directed and unidirected edges together) hyperedges, nested graphs, or ports.

## **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_graphml(G, "test.graphml")
```

## **9.8 JSON**

Generate and parse JSON serializable data for NetworkX graphs.

These formats are suitable for use with the d3.js examples http://d3js.org/

The three formats that you can generate with NetworkX are:

- node-link like in the d3.js example http://bl.ocks.org/mbostock/4062045
- tree like in the d3.js example http://bl.ocks.org/mbostock/4063550
- adjacency like in the d3.js example http://bost.ocks.org/mike/miserables/

$node_link_data(G[, attrs])$	Return data in node-link format that is suitable for JSON serialization and use in Javascr
node_link_graph(data[, directed,])	Return graph from node-link data format.
adjacency_data(G[, attrs])	Return data in adjacency format that is suitable for JSON serialization and use in Javasca
adjacency_graph(data[, directed,])	Return graph from adjacency data format.
tree_data(G, root[, attrs])	Return data in tree format that is suitable for JSON serialization and use in Javascript do
tree_graph(data[, attrs])	Return graph from tree data format.

## 9.8.1 node\_link\_data

node\_link\_data (*G*, attrs={'source': 'source', 'target': 'target', 'key': 'key', 'id': 'id'})

Return data in node-link format that is suitable for JSON serialization and use in Javascript documents.

## Parameters G: NetworkX graph

attrs: dict

A dictionary that contains four keys 'id', 'source', 'target' and 'key'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. The values should be unique. Default value: dict(id='id', source='source', target='target', key='key').

If some user-defined graph data use these attribute names as data keys, they may be silently dropped.

#### Returns data: dict

A dictionary with node-link formatted data.

### Raises NetworkXError

If values in attrs are not unique.

### See also:

```
node_link_graph, adjacency_data, tree_data
```

#### **Notes**

Graph, node, and link attributes are stored in this format. Note that attribute keys will be converted to strings in order to comply with JSON.

The default value of attrs will be changed in a future release of NetworkX.

#### **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.Graph([(1,2)])
>>> data = json_graph.node_link_data(G)

To serialize with json
>>> import json
>>> s = json.dumps(data)
```

9.8. JSON 449

## 9.8.2 node link graph

Parameters data: dict

node-link formatted graph data

directed: bool

If True, and direction not specified in data, return a directed graph.

multigraph: bool

If True, and multigraph not specified in data, return a multigraph.

attrs: dict

A dictionary that contains four keys 'id', 'source', 'target' and 'key'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. Default value: dict(id='id', source='source', target='target', key='key').

**Returns** G: NetworkX graph

A NetworkX graph object

#### See also:

```
node_link_data, adjacency_data, tree_data
```

#### **Notes**

The default value of attrs will be changed in a future release of NetworkX.

### **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.Graph([(1,2)])
>>> data = json_graph.node_link_data(G)
>>> H = json_graph.node_link_graph(data)
```

## 9.8.3 adjacency\_data

```
adjacency_data(G, attrs={'id': 'id', 'key': 'key'})
```

Return data in adjacency format that is suitable for JSON serialization and use in Javascript documents.

### **Parameters G**: NetworkX graph

attrs: dict

A dictionary that contains two keys 'id' and 'key'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. The values should be unique. Default value: dict(id='id', key='key').

If some user-defined graph data use these attribute names as data keys, they may be silently dropped.

#### Returns data: dict

A dictionary with adjacency formatted data.

### Raises NetworkXError

If values in attrs are not unique.

#### See also:

```
adjacency_graph, node_link_data, tree_data
```

#### **Notes**

Graph, node, and link attributes will be written when using this format but attribute keys must be strings if you want to serialize the resulting data with JSON.

The default value of attrs will be changed in a future release of NetworkX.

#### **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.Graph([(1,2)])
>>> data = json_graph.adjacency_data(G)
To serialize with json
>>> import json
>>> s = json.dumps(data)
```

## 9.8.4 adjacency graph

adjacency\_graph (data, directed=False, multigraph=True, attrs={'id': 'id', 'key': 'key'})
Return graph from adjacency data format.

Parameters data: dict

Adjacency list formatted graph data

Returns G: NetworkX graph

A NetworkX graph object

directed: bool

If True, and direction not specified in data, return a directed graph.

multigraph: bool

If True, and multigraph not specified in data, return a multigraph.

attrs: dict

A dictionary that contains two keys 'id' and 'key'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. The values should be unique. Default value: dict(id='id', key='key').

### See also:

```
adjacency_graph, node_link_data, tree_data
```

9.8. JSON 451

#### **Notes**

The default value of attrs will be changed in a future release of NetworkX.

## **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.Graph([(1,2)])
>>> data = json_graph.adjacency_data(G)
>>> H = json_graph.adjacency_graph(data)
```

## 9.8.5 tree\_data

```
tree_data(G, root, attrs={'children': 'children', 'id': 'id'})
```

Return data in tree format that is suitable for JSON serialization and use in Javascript documents.

## **Parameters G**: NetworkX graph

G must be an oriented tree

root: node

The root of the tree

attrs: dict

A dictionary that contains two keys 'id' and 'children'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. The values should be unique. Default value: dict(id='id', children='children').

If some user-defined graph data use these attribute names as data keys, they may be silently dropped.

#### Returns data: dict

A dictionary with node-link formatted data.

### Raises NetworkXError

If values in attrs are not unique.

### See also:

```
tree_graph, node_link_data, node_link_data
```

#### **Notes**

Node attributes are stored in this format but keys for attributes must be strings if you want to serialize with JSON.

Graph and edge attributes are not stored.

The default value of attrs will be changed in a future release of NetworkX.

### **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.DiGraph([(1,2)])
>>> data = json_graph.tree_data(G,root=1)

To serialize with json
>>> import json
>>> s = json.dumps(data)
```

## 9.8.6 tree\_graph

```
tree_graph (data, attrs={'children': 'children', 'id': 'id'})

Return graph from tree data format.
```

Parameters data: dict

Tree formatted graph data

**Returns** G: NetworkX DiGraph

attrs: dict

A dictionary that contains two keys 'id' and 'children'. The corresponding values provide the attribute names for storing NetworkX-internal graph data. The values should be unique. Default value: dict(id='id', children='children').

### See also:

```
tree_graph, node_link_data, adjacency_data
```

### **Notes**

The default value of attrs will be changed in a future release of NetworkX.

## **Examples**

```
>>> from networkx.readwrite import json_graph
>>> G = nx.DiGraph([(1,2)])
>>> data = json_graph.tree_data(G,root=1)
>>> H = json_graph.tree_graph(data)
```

## **9.9 LEDA**

Read graphs in LEDA format.

LEDA is a C++ class library for efficient data types and algorithms.

## 9.9.1 Format

See http://www.algorithmic-solutions.info/leda\_guide/graphs/leda\_native\_graph\_fileformat.html

9.9. LEDA 453

read_leda(path[, encoding])	Read graph in LEDA format from path.
parse_leda(lines)	Read graph in LEDA format from string or iterable.

## 9.9.2 read\_leda

read\_leda (path, encoding='UTF-8')

Read graph in LEDA format from path.

Parameters path: file or string

File or filename to read. Filenames ending in .gz or .bz2 will be uncompressed.

Returns G: NetworkX graph

### References

[R370]

### **Examples**

G=nx.read\_leda('file.leda')

## 9.9.3 parse\_leda

parse\_leda(lines)

Read graph in LEDA format from string or iterable.

Parameters lines: string or iterable

Data in LEDA format.

Returns G: NetworkX graph

### References

[R369]

## **Examples**

G=nx.parse\_leda(string)

## 9.10 YAML

Read and write NetworkX graphs in YAML format.

"YAML is a data serialization format designed for human readability and interaction with scripting languages." See http://www.yaml.org for documentation.

## 9.10.1 Format

http://pyyaml.org/wiki/PyYAML

read_yaml(path)	Read graph in YAML format from path.
<pre>write_yaml(G, path[, encoding])</pre>	Write graph G in YAML format to path.

## 9.10.2 read\_yaml

### read\_yaml (path)

Read graph in YAML format from path.

YAML is a data serialization format designed for human readability and interaction with scripting languages [R373].

Parameters path: file or string

File or filename to read. Filenames ending in .gz or .bz2 will be uncompressed.

Returns G: NetworkX graph

#### References

[R373]

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_yaml(G,'test.yaml')
>>> G=nx.read_yaml('test.yaml')
```

## 9.10.3 write yaml

```
\label{eq:write_yaml} \textbf{(}G, path, encoding = \text{'}UTF\text{-}8\text{'}, \text{ **}kwds\text{'})
```

Write graph G in YAML format to path.

YAML is a data serialization format designed for human readability and interaction with scripting languages [R374].

Parameters G: graph

A NetworkX graph

path: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be compressed.

encoding: string, optional

Specify which encoding to use when writing file.

References

[R374]

9.10. YAML 455

### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_yaml(G,'test.yaml')
```

# 9.11 SparseGraph6

## 9.11.1 Graph6

Graph6

Read and write graphs in graph6 format.

#### **Format**

"graph6 and sparse6 are formats for storing undirected graphs in a compact manner, using only printable ASCII characters. Files in these formats have text type and contain one line per graph."

See http://cs.anu.edu.au/~bdm/data/formats.txt for details.

parse_graph6(string)	Read a simple undirected graph in graph6 format from string.
read_graph6(path)	Read simple undirected graphs in graph6 format from path.
$generate\_graph6(G[, nodes, header])$	Generate graph6 format string from a simple undirected graph.
<pre>write_graph6(G, path[, nodes, header])</pre>	Write a simple undirected graph to path in graph6 format.

## parse\_graph6

parse\_graph6 (string)

Read a simple undirected graph in graph6 format from string.

Parameters string: string

Data in graph6 format

Returns G: Graph

Raises NetworkXError

If the string is unable to be parsed in graph6 format

### See also:

```
generate_graph6, read_graph6, write_graph6
```

#### References

Graph6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

## **Examples**

```
>>> G = nx.parse_graph6('A_')
>>> sorted(G.edges())
[(0, 1)]
```

## read graph6

## read\_graph6 (path)

Read simple undirected graphs in graph6 format from path.

Parameters path: file or string

File or filename to write.

**Returns** G: Graph or list of Graphs

If the file contains multiple lines then a list of graphs is returned

Raises NetworkXError

If the string is unable to be parsed in graph6 format

#### See also:

```
generate_graph6, parse_graph6, write_graph6
```

#### References

Graph6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

## **Examples**

```
>>> nx.write_graph6(nx.Graph([(0,1)]), 'test.g6')
>>> G = nx.read_graph6('test.g6')
>>> sorted(G.edges())
[(0, 1)]
```

## generate\_graph6

```
generate_graph6 (G, nodes=None, header=True)
```

Generate graph6 format string from a simple undirected graph.

**Parameters G**: Graph (undirected)

nodes: list or iterable

Nodes are labeled 0...n-1 in the order provided. If None the ordering given by G.nodes() is used.

header: bool

If True add '>>graph6<<' string to head of data

**Returns** s : string

String in graph6 format

## Raises NetworkXError

If the graph is directed or has parallel edges

#### See also:

```
read_graph6, parse_graph6, write_graph6
```

#### **Notes**

The format does not support edge or node labels, parallel edges or self loops. If self loops are present they are silently ignored.

#### References

Graph6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

### **Examples**

```
>>> G = nx.Graph([(0, 1)])
>>> nx.generate_graph6(G)
'>>graph6<<A_'</pre>
```

### write\_graph6

```
write_graph6 (G, path, nodes=None, header=True)
```

Write a simple undirected graph to path in graph6 format.

```
Parameters G: Graph (undirected)
```

path: file or string

File or filename to write.

### nodes: list or iterable

Nodes are labeled 0...n-1 in the order provided. If None the ordering given by G.nodes() is used.

### header: bool

If True add '>>graph6<<' string to head of data

## Raises NetworkXError

If the graph is directed or has parallel edges

#### See also:

```
generate_graph6, parse_graph6, read_graph6
```

## **Notes**

The format does not support edge or node labels, parallel edges or self loops. If self loops are present they are silently ignored.

## References

Graph6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

### **Examples**

```
>>> G = nx.Graph([(0, 1)])
>>> nx.write_graph6(G, 'test.g6')
```

## 9.11.2 Sparse6

Sparse6

Read and write graphs in sparse6 format.

#### **Format**

"graph6 and sparse6 are formats for storing undirected graphs in a compact manner, using only printable ASCII characters. Files in these formats have text type and contain one line per graph."

See http://cs.anu.edu.au/~bdm/data/formats.txt for details.

parse_sparse6(string)	Read an undirected graph in sparse6 format from string.
read_sparse6(path)	Read an undirected graph in sparse6 format from path.
$generate\_sparse6(G[, nodes, header])$	Generate sparse6 format string from an undirected graph.
<pre>write_sparse6(G, path[, nodes, header])</pre>	Write graph G to given path in sparse6 format.

## parse\_sparse6

## parse\_sparse6 (string)

Read an undirected graph in sparse6 format from string.

Parameters string: string

Data in sparse6 format

Returns G: Graph

Raises NetworkXError

If the string is unable to be parsed in sparse6 format

### See also:

```
generate_sparse6, read_sparse6, write_sparse6
```

#### References

Sparse6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt

## **Examples**

```
>>> G = nx.parse_sparse6(':A_')
>>> sorted(G.edges())
[(0, 1), (0, 1), (0, 1)]
```

### read sparse6

### read\_sparse6 (path)

Read an undirected graph in sparse6 format from path.

Parameters path: file or string

File or filename to write.

**Returns** G: Graph/Multigraph or list of Graphs/MultiGraphs

If the file contains multple lines then a list of graphs is returned

Raises NetworkXError

If the string is unable to be parsed in sparse6 format

#### See also:

```
generate_sparse6, read_sparse6, parse_sparse6
```

#### References

Sparse6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt

### **Examples**

```
>>> nx.write_sparse6(nx.Graph([(0,1),(0,1),(0,1)]), 'test.s6')
>>> G = nx.read_sparse6('test.s6')
>>> sorted(G.edges())
[(0, 1)]
```

### generate\_sparse6

```
generate_sparse6 (G, nodes=None, header=True)
```

Generate sparse6 format string from an undirected graph.

Parameters G: Graph (undirected)

nodes: list or iterable

Nodes are labeled 0...n-1 in the order provided. If None the ordering given by G.nodes() is used.

header: bool

If True add '>>sparse6<<' string to head of data

**Returns** s: string

String in sparse6 format

Raises NetworkXError

If the graph is directed

### See also:

```
read_sparse6, parse_sparse6, write_sparse6
```

#### **Notes**

The format does not support edge or node labels. References ———— Sparse6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

### **Examples**

```
>>> G = nx.MultiGraph([(0, 1), (0, 1), (0, 1)])
>>> nx.generate_sparse6(G)
'>>sparse6<<:A_'</pre>
```

## write\_sparse6

```
write_sparse6 (G, path, nodes=None, header=True)
```

Write graph G to given path in sparse6 format. Parameters ———— G : Graph (undirected)

path [file or string] File or filename to write

**nodes:** list or iterable Nodes are labeled 0...n-1 in the order provided. If None the ordering given by G.nodes() is used.

header: bool If True add '>>sparse6<<' string to head of data

#### Raises NetworkXError

If the graph is directed

#### See also:

```
read_sparse6, parse_sparse6, generate_sparse6
```

#### **Notes**

The format does not support edge or node labels.

#### References

Sparse6 specification: http://cs.anu.edu.au/~bdm/data/formats.txt for details.

### **Examples**

```
>>> G = nx.Graph([(0, 1), (0, 1), (0, 1)])
>>> nx.write_sparse6(G, 'test.s6')
```

# 9.12 Pajek

Read graphs in Pajek format.

This implementation handles directed and undirected graphs including those with self loops and parallel edges.

9.12. Pajek 461

## 9.12.1 Format

See http://vlado.fmf.uni-lj.si/pub/networks/pajek/doc/draweps.htm for format information.

read_pajek(path[, encoding])	Read graph in Pajek format from path.
<pre>write_pajek(G, path[, encoding])</pre>	Write graph in Pajek format to path.
parse_pajek(lines)	Parse Pajek format graph from string or iterable.

## 9.12.2 read pajek

read\_pajek (path, encoding='UTF-8')

Read graph in Pajek format from path.

Parameters path: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be uncompressed.

**Returns G**: NetworkX MultiGraph or MultiDiGraph.

#### References

See http://vlado.fmf.uni-lj.si/pub/networks/pajek/doc/draweps.htm for format information.

## **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_pajek(G, "test.net")
>>> G=nx.read_pajek("test.net")
```

To create a Graph instead of a MultiGraph use

```
>>> G1=nx.Graph(G)
```

## 9.12.3 write\_pajek

```
write_pajek (G, path, encoding='UTF-8')
```

Write graph in Pajek format to path.

Parameters G: graph

A Networkx graph

path: file or string

File or filename to write. Filenames ending in .gz or .bz2 will be compressed.

### References

See http://vlado.fmf.uni-lj.si/pub/networks/pajek/doc/draweps.htm for format information.

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> nx.write_pajek(G, "test.net")
```

## 9.12.4 parse pajek

```
parse_pajek (lines)
```

Parse Pajek format graph from string or iterable.

**Parameters lines**: string or iterable

Data in Pajek format.

Returns G: NetworkX graph

See also:

read\_pajek

# 9.13 GIS Shapefile

Generates a networkx. DiGraph from point and line shapefiles.

"The Esri Shapefile or simply a shapefile is a popular geospatial vector data format for geographic information systems software. It is developed and regulated by Esri as a (mostly) open specification for data interoperability among Esri and other software products." See <a href="http://en.wikipedia.org/wiki/Shapefile">http://en.wikipedia.org/wiki/Shapefile</a> for additional information.

read_shp(path)	Generates a networkx.DiGraph from shapefiles.
write_shp(G, outdir)	Writes a networkx.DiGraph to two shapefiles, edges and nodes.

## 9.13.1 read\_shp

#### read\_shp (path)

Generates a networkx.DiGraph from shapefiles. Point geometries are translated into nodes, lines into edges. Coordinate tuples are used as keys. Attributes are preserved, line geometries are simplified into start and end coordinates. Accepts a single shapefile or directory of many shapefiles.

"The Esri Shapefile or simply a shapefile is a popular geospatial vector data format for geographic information systems software [R371]."

Parameters path: file or string

File, directory, or filename to read.

**Returns G** : NetworkX graph

#### References

[R371]

9.13. GIS Shapefile 463

#### **Examples**

```
>>> G=nx.read_shp('test.shp')
```

## 9.13.2 write shp

### $write\_shp(G, outdir)$

Writes a networkx.DiGraph to two shapefiles, edges and nodes. Nodes and edges are expected to have a Well Known Binary (Wkb) or Well Known Text (Wkt) key in order to generate geometries. Also acceptable are nodes with a numeric tuple key (x,y).

"The Esri Shapefile or simply a shapefile is a popular geospatial vector data format for geographic information systems software [R372]."

Parameters outdir: directory path

Output directory for the two shapefiles.

Returns None

#### References

[R372]

## **Examples**

nx.write\_shp(digraph, '/shapefiles') # doctest +SKIP

**TEN** 

## **DRAWING**

# 10.1 Matplotlib

Draw networks with matplotlib.

### 10.1.1 See Also

matplotlib: http://matplotlib.org/

pygraphviz: http://pygraphviz.github.io/

draw(G[, pos, ax, hold])	Draw the graph G with Matplotlib.
$draw_networkx(G[, pos, with_labels])$	Draw the graph G using Matplotlib.
draw_networkx_nodes(G, pos[, nodelist,])	Draw the nodes of the graph G.
draw_networkx_edges(G, pos[, edgelist,])	Draw the edges of the graph G.
draw_networkx_labels(G, pos[, labels,])	Draw node labels on the graph G.
draw_networkx_edge_labels(G, pos[,])	Draw edge labels.
draw_circular(G, **kwargs)	Draw the graph G with a circular layout.
draw_random(G, **kwargs)	Draw the graph G with a random layout.
draw_spectral(G, **kwargs)	Draw the graph G with a spectral layout.
draw_spring(G, **kwargs)	Draw the graph G with a spring layout.
draw_shell(G, **kwargs)	Draw networkx graph with shell layout.
$draw\_graphviz(G[,prog])$	Draw networkx graph with graphviz layout.

## 10.1.2 draw

draw(G, pos=None, ax=None, hold=None, \*\*kwds)

Draw the graph G with Matplotlib.

Draw the graph as a simple representation with no node labels or edge labels and using the full Matplotlib figure area and no axis labels by default. See draw\_networkx() for more full-featured drawing that allows title, axis labels etc.

 $Parameters \ G: {\sf graph}$ 

A networkx graph

pos: dictionary, optional

A dictionary with nodes as keys and positions as values. If not specified a spring layout positioning will be computed. See networkx.layout for functions that compute node positions.

ax : Matplotlib Axes object, optional

Draw the graph in specified Matplotlib axes.

hold: bool, optional

Set the Matplotlib hold state. If True subsequent draw commands will be added to the current axes.

\*\*kwds: optional keywords

See networkx.draw\_networkx() for a description of optional keywords.

#### See also:

```
draw_networkx, draw_networkx_nodes, draw_networkx_edges, draw_networkx_labels,
draw_networkx_edge_labels
```

#### **Notes**

This function has the same name as pylab.draw and pyplot.draw so beware when using

```
>>> from networkx import *
```

since you might overwrite the pylab.draw function.

With pyplot use

```
>>> import matplotlib.pyplot as plt
>>> import networkx as nx
>>> G=nx.dodecahedral_graph()
>>> nx.draw(G) # networkx draw()
>>> plt.draw() # pyplot draw()
```

Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html

#### **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> nx.draw(G)
>>> nx.draw(G,pos=nx.spring_layout(G)) # use spring layout
```

# 10.1.3 draw\_networkx

```
draw_networkx (G, pos=None, with_labels=True, **kwds)
```

Draw the graph G using Matplotlib.

Draw the graph with Matplotlib with options for node positions, labeling, titles, and many other drawing features. See draw() for simple drawing without labels or axes.

```
Parameters G: graph
```

A networkx graph

pos: dictionary, optional

A dictionary with nodes as keys and positions as values. If not specified a spring layout positioning will be computed. See networkx.layout for functions that compute node positions.

with\_labels : bool, optional (default=True)

Set to True to draw labels on the nodes.

ax : Matplotlib Axes object, optional

Draw the graph in the specified Matplotlib axes.

**nodelist**: list, optional (default G.nodes())

Draw only specified nodes

edgelist : list, optional (default=G.edges())

Draw only specified edges

node\_size : scalar or array, optional (default=300)

Size of nodes. If an array is specified it must be the same length as nodelist.

**node\_color**: color string, or array of floats, (default='r')

Node color. Can be a single color format string, or a sequence of colors with the same length as nodelist. If numeric values are specified they will be mapped to colors using the cmap and vmin,vmax parameters. See matplotlib.scatter for more details.

node\_shape : string, optional (default='o')

The shape of the node. Specification is as matplotlib.scatter marker, one of 'so^>v<dph8'.

alpha: float, optional (default=1.0)

The node and edge transparency

cmap : Matplotlib colormap, optional (default=None)

Colormap for mapping intensities of nodes

vmin,vmax : float, optional (default=None)

Minimum and maximum for node colormap scaling

**linewidths** : [None | scalar | sequence]

Line width of symbol border (default =1.0)

width: float, optional (default=1.0)

Line width of edges

edge\_color : color string, or array of floats (default='r')

Edge color. Can be a single color format string, or a sequence of colors with the same length as edgelist. If numeric values are specified they will be mapped to colors using the edge\_cmap and edge\_vmin,edge\_vmax parameters.

edge\_cmap: Matplotlib colormap, optional (default=None)

Colormap for mapping intensities of edges

edge\_vmin,edge\_vmax : floats, optional (default=None)

Minimum and maximum for edge colormap scaling

**style**: string, optional (default='solid')

Edge line style (solidldashedldotted,dashdot)

**labels**: dictionary, optional (default=None)

10.1. Matplotlib 467

```
Node labels in a dictionary keyed by node of text labels
              font size: int, optional (default=12)
                  Font size for text labels
              font_color : string, optional (default='k' black)
                  Font color string
              font_weight : string, optional (default='normal')
                  Font weight
              font_family : string, optional (default='sans-serif')
                  Font family
              label: string, optional
                  Label for graph legend
     See also:
     draw.
                draw_networkx_nodes,
                                              draw_networkx_edges,
                                                                             draw_networkx_labels,
     draw networkx edge labels
     Examples
     >>> G=nx.dodecahedral_graph()
     >>> nx.draw(G)
     >>> nx.draw(G,pos=nx.spring_layout(G)) # use spring layout
     >>> import matplotlib.pyplot as plt
     >>> limits=plt.axis('off') # turn of axis
     Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html
10.1.4 draw_networkx_nodes
draw_networkx_nodes (G, pos, nodelist=None, node_size=300, node_color='r', node_shape='o', al-
                          pha=1.0, cmap=None, vmin=None, vmax=None, ax=None, linewidths=None, la-
                          bel=None, **kwds)
     Draw the nodes of the graph G.
     This draws only the nodes of the graph G.
          Parameters G: graph
                  A networkx graph
              pos: dictionary
                  A dictionary with nodes as keys and positions as values. Positions should be sequences
                  of length 2.
              ax : Matplotlib Axes object, optional
                  Draw the graph in the specified Matplotlib axes.
              nodelist: list, optional
```

Draw only specified nodes (default G.nodes())

```
node_size : scalar or array
```

Size of nodes (default=300). If an array is specified it must be the same length as nodelist.

node\_color : color string, or array of floats

Node color. Can be a single color format string (default='r'), or a sequence of colors with the same length as nodelist. If numeric values are specified they will be mapped to colors using the cmap and vmin,vmax parameters. See matplotlib.scatter for more details.

#### node\_shape: string

The shape of the node. Specification is as matplotlib.scatter marker, one of 'so^>v<dph8' (default='o').

alpha: float

The node transparency (default=1.0)

cmap: Matplotlib colormap

Colormap for mapping intensities of nodes (default=None)

vmin,vmax: floats

Minimum and maximum for node colormap scaling (default=None)

**linewidths**: [None | scalar | sequence]

Line width of symbol border (default =1.0)

label : [Nonel string]

Label for legend

Returns matplotlib.collections.PathCollection

PathCollection of the nodes.

#### See also:

```
draw, draw_networkx, draw_networkx_edges, draw_networkx_labels,
```

### **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> nodes=nx.draw_networkx_nodes(G,pos=nx.spring_layout(G))
```

Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html

## 10.1.5 draw\_networkx\_edges

Draw the edges of the graph G.

This draws only the edges of the graph G.

**Parameters G**: graph

10.1. Matplotlib 469

```
A networkx graph
```

**pos**: dictionary

A dictionary with nodes as keys and positions as values. Positions should be sequences of length 2.

edgelist: collection of edge tuples

Draw only specified edges(default=G.edges())

width: float

Line width of edges (default =1.0)

edge\_color : color string, or array of floats

Edge color. Can be a single color format string (default='r'), or a sequence of colors with the same length as edgelist. If numeric values are specified they will be mapped to colors using the edge\_cmap and edge\_vmin,edge\_vmax parameters.

style: string

Edge line style (default='solid') (solidldashedldotted,dashdot)

alpha: float

The edge transparency (default=1.0)

edge\_cmap: Matplotlib colormap

Colormap for mapping intensities of edges (default=None)

edge\_vmin,edge\_vmax: floats

Minimum and maximum for edge colormap scaling (default=None)

ax : Matplotlib Axes object, optional

Draw the graph in the specified Matplotlib axes.

arrows : bool, optional (default=True)

For directed graphs, if True draw arrowheads.

label: [Nonel string]

Label for legend

Returns matplotlib.collection.LineCollection

LineCollection of the edges

#### See also:

```
draw, draw_networkx, draw_networkx_nodes, draw_networkx_labels,
```

#### Notes

For directed graphs, "arrows" (actually just thicker stubs) are drawn at the head end. Arrows can be turned off with keyword arrows=False. Yes, it is ugly but drawing proper arrows with Matplotlib this way is tricky.

#### **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> edges=nx.draw_networkx_edges(G,pos=nx.spring_layout(G))
```

Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html

```
10.1.6 draw networkx labels
draw_networkx_labels (G, pos, labels=None, font_size=12, font_color='k', font_family='sans-serif',
                              font weight='normal', alpha=1.0, bbox=None, ax=None, **kwds)
     Draw node labels on the graph G.
           Parameters G: graph
                   A networkx graph
               pos: dictionary
                   A dictionary with nodes as keys and positions as values. Positions should be sequences
                   of length 2.
               labels: dictionary, optional (default=None)
                   Node labels in a dictionary keyed by node of text labels
               font_size : int
                   Font size for text labels (default=12)
               font_color : string
                   Font color string (default='k' black)
               font_family: string
                   Font family (default='sans-serif')
               font_weight : string
                   Font weight (default='normal')
               alpha: float
                   The text transparency (default=1.0)
               ax : Matplotlib Axes object, optional
                   Draw the graph in the specified Matplotlib axes.
           Returns dict
                   dict of labels keyed on the nodes
     See also:
```

```
draw, draw_networkx, draw_networkx_nodes, draw_networkx_edges, draw_networkx_edge_labels
```

### **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> labels=nx.draw_networkx_labels(G,pos=nx.spring_layout(G))
```

10.1. Matplotlib 471

Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html

## 10.1.7 draw\_networkx\_edge\_labels

```
draw_networkx_edge_labels(G,
                                           pos,
                                                    edge labels=None,
                                                                           label pos=0.5,
                                                                                              font size=10,
                                                                                     font_weight='normal',
                                     font_color='k',
                                                       font_family='sans-serif',
                                     alpha=1.0, bbox=None, ax=None, rotate=True, **kwds)
      Draw edge labels.
           Parameters G: graph
                    A networkx graph
               pos: dictionary
                    A dictionary with nodes as keys and positions as values. Positions should be sequences
                    of length 2.
               ax: Matplotlib Axes object, optional
                    Draw the graph in the specified Matplotlib axes.
               alpha: float
                   The text transparency (default=1.0)
               edge_labels: dictionary
                   Edge labels in a dictionary keyed by edge two-tuple of text labels (default=None). Only
                    labels for the keys in the dictionary are drawn.
               label_pos: float
                    Position of edge label along edge (0=head, 0.5=center, 1=tail)
               font size: int
                    Font size for text labels (default=12)
               font_color : string
                   Font color string (default='k' black)
               font_weight : string
                    Font weight (default='normal')
               font family: string
                   Font family (default='sans-serif')
               bbox : Matplotlib bbox
                    Specify text box shape and colors.
               clip_on: bool
                   Turn on clipping at axis boundaries (default=True)
           Returns dict
                    dict of labels keyed on the edges
      See also:
```

draw networkx nodes,

draw networkx edges,

draw networkx,

draw\_networkx\_labels

#### **Examples**

```
>>> G=nx.dodecahedral_graph()
>>> edge_labels=nx.draw_networkx_edge_labels(G,pos=nx.spring_layout(G))
```

Also see the NetworkX drawing examples at http://networkx.github.io/documentation/latest/gallery.html

## 10.1.8 draw\_circular

```
draw_circular(G, **kwargs)
```

Draw the graph G with a circular layout.

**Parameters G**: graph

A networkx graph

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords, with the exception of the pos parameter which is not used by this function.

## 10.1.9 draw\_random

```
draw_random(G, **kwargs)
```

Draw the graph G with a random layout.

Parameters G: graph

A networkx graph

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords, with the exception of the pos parameter which is not used by this function.

## 10.1.10 draw\_spectral

```
draw_spectral(G, **kwargs)
```

Draw the graph G with a spectral layout.

Parameters G: graph

A networkx graph

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords, with the exception of the pos parameter which is not used by this function.

## 10.1.11 draw\_spring

```
draw_spring(G, **kwargs)
```

Draw the graph G with a spring layout.

Parameters G: graph

A networkx graph

10.1. Matplotlib 473

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords, with the exception of the pos parameter which is not used by this function.

## 10.1.12 draw\_shell

```
draw_shell(G, **kwargs)
```

Draw networkx graph with shell layout.

Parameters G: graph

A networkx graph

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords, with the exception of the pos parameter which is not used by this function.

## 10.1.13 draw\_graphviz

draw\_graphviz (G, prog='neato', \*\*kwargs)

Draw networkx graph with graphviz layout.

Parameters G: graph

A networkx graph

prog: string, optional

Name of Graphviz layout program

\*\*kwargs: optional keywords

See networkx.draw\_networkx() for a description of optional keywords.

# 10.2 Graphviz AGraph (dot)

Interface to pygraphviz AGraph class.

## 10.2.1 Examples

```
>>> G=nx.complete_graph(5)
>>> A=nx.to_agraph(G)
>>> H=nx.from_agraph(A)
```

## 10.2.2 See Also

Pygraphviz: http://pygraphviz.github.io/

from_agraph(A[, create_using])	Return a NetworkX Graph or DiGraph from a PyGraphviz graph.
to_agraph $(N)$	Return a pygraphviz graph from a NetworkX graph N.
write_dot(G, path)	Write NetworkX graph G to Graphviz dot format on path.
	Continued on next page

Table 10.2 – continued from previous page

read_dot(path)	Return a NetworkX graph from a dot file on path.
<pre>graphviz_layout(G[, prog, root, args])</pre>	Create node positions for G using Graphviz.
<pre>pygraphviz_layout(G[, prog, root, args])</pre>	Create node positions for G using Graphviz.

## 10.2.3 from agraph

from\_agraph (A, create\_using=None)

Return a NetworkX Graph or DiGraph from a PyGraphviz graph.

Parameters A: PyGraphviz AGraph

A graph created with PyGraphviz

create\_using : NetworkX graph class instance

The output is created using the given graph class instance

#### **Notes**

The Graph G will have a dictionary G.graph\_attr containing the default graphviz attributes for graphs, nodes and edges.

Default node attributes will be in the dictionary G.node\_attr which is keyed by node.

Edge attributes will be returned as edge data in G. With edge\_attr=False the edge data will be the Graphviz edge weight attribute or the value 1 if no edge weight attribute is found.

#### **Examples**

```
>>> K5=nx.complete_graph(5)
>>> A=nx.to_agraph(K5)
>>> G=nx.from_agraph(A)
>>> G=nx.from_agraph(A)
```

## 10.2.4 to agraph

## $\verb"to_agraph"\,(N)$

Return a pygraphviz graph from a NetworkX graph N.

Parameters N: NetworkX graph

A graph created with NetworkX

#### **Notes**

If N has an dict N.graph\_attr an attempt will be made first to copy properties attached to the graph (see from\_agraph) and then updated with the calling arguments if any.

#### **Examples**

```
>>> K5=nx.complete_graph(5)
>>> A=nx.to_agraph(K5)
```

## 10.2.5 write\_dot

write\_dot(G, path)

Write NetworkX graph G to Graphviz dot format on path.

Parameters G: graph

A networkx graph

path: filename

Filename or file handle to write

## 10.2.6 read dot

read\_dot (path)

Return a NetworkX graph from a dot file on path.

Parameters path: file or string

File name or file handle to read.

# 10.2.7 graphviz\_layout

```
graphviz_layout (G, prog='neato', root=None, args='') Create node positions for G using Graphviz.
```

Parameters G: NetworkX graph

A graph created with NetworkX

prog : string

Name of Graphviz layout program

root: string, optional

Root node for twopi layout

args: string, optional

Extra arguments to Graphviz layout program

Returns: dictionary

Dictionary of x,y, positions keyed by node.

#### **Notes**

This is a wrapper for pygraphviz\_layout.

#### **Examples**

```
>>> G=nx.petersen_graph()
>>> pos=nx.graphviz_layout(G)
>>> pos=nx.graphviz_layout(G,prog='dot')
```

## 10.2.8 pygraphviz\_layout

```
pygraphviz_layout (G, prog='neato', root=None, args='') Create node positions for G using Graphviz.
```

Parameters G: NetworkX graph

A graph created with NetworkX

prog : string

Name of Graphviz layout program

root: string, optional

Root node for twopi layout

args: string, optional

Extra arguments to Graphviz layout program

Returns: dictionary

Dictionary of x,y, positions keyed by node.

#### **Examples**

```
>>> G=nx.petersen_graph()
>>> pos=nx.graphviz_layout(G)
>>> pos=nx.graphviz_layout(G,prog='dot')
```

# 10.3 Graphviz with pydot

Import and export NetworkX graphs in Graphviz dot format using pydot.

Either this module or nx\_pygraphviz can be used to interface with graphviz.

## 10.3.1 See Also

Pydot: http://code.google.com/p/pydot/ Graphviz: http://www.research.att.com/sw/tools/graphviz/ DOT Language: http://www.graphviz.org/doc/info/lang.html

from_pydot(P)	Return a NetworkX graph from a Pydot graph.	
to_pydot(N[, strict])	Return a pydot graph from a NetworkX graph N.	
write_dot(G, path)	Write NetworkX graph G to Graphviz dot format on path.	
read_dot(path)	Return a NetworkX MultiGraph or MultiDiGraph from a dot file on path.	
$graphviz_layout(G[, prog, root])$	Create node positions using Pydot and Graphviz.	
<pre>pydot_layout(G[, prog, root])</pre>	Create node positions using Pydot and Graphviz.	

## 10.3.2 from pydot

```
from\_pydot(P)
```

Return a NetworkX graph from a Pydot graph.

Parameters P: Pydot graph

A graph created with Pydot

**Returns** G: NetworkX multigraph

A MultiGraph or MultiDiGraph.

### **Examples**

```
>>> K5=nx.complete_graph(5)
>>> A=nx.to_pydot(K5)
>>> G=nx.from_pydot(A) # return MultiGraph
>>> G=nx.Graph(nx.from_pydot(A)) # make a Graph instead of MultiGraph
```

## 10.3.3 to\_pydot

to\_pydot (N, strict=True)

Return a pydot graph from a NetworkX graph N.

Parameters N: NetworkX graph

A graph created with NetworkX

#### **Examples**

```
>>> K5=nx.complete_graph(5)
>>> P=nx.to_pydot(K5)
```

## 10.3.4 write\_dot

```
write_dot(G, path)
```

Write NetworkX graph G to Graphviz dot format on path.

Path can be a string or a file handle.

## 10.3.5 read dot

```
read_dot (path)
```

Return a NetworkX MultiGraph or MultiDiGraph from a dot file on path.

Parameters path: filename or file handle

**Returns G** : NetworkX multigraph

A MultiGraph or MultiDiGraph.

#### **Notes**

Use G=nx.Graph(nx.read\_dot(path)) to return a Graph instead of a MultiGraph.

# 10.3.6 graphviz\_layout

```
graphviz_layout (G, prog='neato', root=None, **kwds)

Create node positions using Pydot and Graphviz.
```

Returns a dictionary of positions keyed by node.

#### Notes

This is a wrapper for pydot\_layout.

#### **Examples**

```
>>> G=nx.complete_graph(4)
>>> pos=nx.graphviz_layout(G)
>>> pos=nx.graphviz_layout(G,prog='dot')
```

## 10.3.7 pydot\_layout

```
pydot_layout (G, prog='neato', root=None, **kwds)
Create node positions using Pydot and Graphviz.
```

Returns a dictionary of positions keyed by node.

#### **Examples**

```
>>> G=nx.complete_graph(4)
>>> pos=nx.pydot_layout(G)
>>> pos=nx.pydot_layout(G,prog='dot')
```

# 10.4 Graph Layout

Node positioning algorithms for graph drawing.

circular_layout(G[, dim, scale])	Position nodes on a circle.
$random\_layout(G[,dim])$	Position nodes uniformly at random in the unit square.
<pre>shell_layout(G[, nlist, dim, scale])</pre>	Position nodes in concentric circles.
spring_layout(G[, dim, k, pos, fixed,])	Position nodes using Fruchterman-Reingold force-directed algorithm.
<pre>spectral_layout(G[, dim, weight, scale])</pre>	Position nodes using the eigenvectors of the graph Laplacian.

10.4. Graph Layout 479

## 10.4.1 circular layout

```
circular_layout (G, dim=2, scale=1)
   Position nodes on a circle.

Parameters G: NetworkX graph

dim: int

   Dimension of layout, currently only dim=2 is supported
   scale: float
        Scale factor for positions

Returns dict:

        A dictionary of positions keyed by node
```

#### **Notes**

This algorithm currently only works in two dimensions and does not try to minimize edge crossings.

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> pos=nx.circular_layout(G)
```

# 10.4.2 random\_layout

```
random_layout (G, dim=2)
```

Position nodes uniformly at random in the unit square.

For every node, a position is generated by choosing each of dim coordinates uniformly at random on the interval [0.0, 1.0).

NumPy (http://scipy.org) is required for this function.

```
Parameters G: NetworkX graph
```

A position will be assigned to every node in G.

dim: int

Dimension of layout.

Returns dict:

A dictionary of positions keyed by node

#### **Examples**

```
>>> G = nx.lollipop_graph(4, 3)
>>> pos = nx.random_layout(G)
```

## 10.4.3 shell layout

A dictionary of positions keyed by node

#### **Notes**

This algorithm currently only works in two dimensions and does not try to minimize edge crossings.

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> shells=[[0],[1,2,3]]
>>> pos=nx.shell_layout(G,shells)
```

## 10.4.4 spring\_layout

**spring\_layout** (*G*, *dim=2*, *k=None*, *pos=None*, *fixed=None*, *iterations=50*, *weight='weight'*, *scale=1.0*) Position nodes using Fruchterman-Reingold force-directed algorithm.

```
Parameters G: NetworkX graph
```

dim: int

Dimension of layout

**k**: float (default=None)

Optimal distance between nodes. If None the distance is set to 1/sqrt(n) where n is the number of nodes. Increase this value to move nodes farther apart.

pos : dict or None optional (default=None)

Initial positions for nodes as a dictionary with node as keys and values as a list or tuple. If None, then use random initial positions.

fixed: list or None optional (default=None)

Nodes to keep fixed at initial position.

iterations: int optional (default=50)

Number of iterations of spring-force relaxation

10.4. Graph Layout 481

```
weight: string or None optional (default='weight')
```

The edge attribute that holds the numerical value used for the edge weight. If None, then all edge weights are 1.

```
scale : float (default=1.0)
```

Scale factor for positions. The nodes are positioned in a box of size [0,scale] x [0,scale].

#### Returns dict:

A dictionary of positions keyed by node

#### **Examples**

```
>>> G=nx.path_graph(4)
>>> pos=nx.spring_layout(G)
```

# The same using longer function name >>> pos=nx.fruchterman\_reingold\_layout(G)

## 10.4.5 spectral\_layout

```
spectral_layout (G, dim=2, weight='weight', scale=1)
```

Position nodes using the eigenvectors of the graph Laplacian.

#### **Parameters G**: NetworkX graph

dim: int

Dimension of layout

weight : string or None optional (default='weight')

The edge attribute that holds the numerical value used for the edge weight. If None, then all edge weights are 1.

scale: float

Scale factor for positions

#### Returns dict:

A dictionary of positions keyed by node

### **Notes**

Directed graphs will be considered as undirected graphs when positioning the nodes.

For larger graphs (>500 nodes) this will use the SciPy sparse eigenvalue solver (ARPACK).

### **Examples**

```
>>> G=nx.path_graph(4)
>>> pos=nx.spectral_layout(G)
```

**CHAPTER** 

## **ELEVEN**

## **EXCEPTIONS**

Base exceptions and errors for NetworkX.

### class NetworkXException

Base class for exceptions in NetworkX.

#### class NetworkXError

Exception for a serious error in NetworkX

#### class NetworkXPointlessConcept

Harary, F. and Read, R. "Is the Null Graph a Pointless Concept?" In Graphs and Combinatorics Conference, George Washington University. New York: Springer-Verlag, 1973.

#### class NetworkXAlgorithmError

Exception for unexpected termination of algorithms.

#### class NetworkXUnfeasible

Exception raised by algorithms trying to solve a problem instance that has no feasible solution.

### class NetworkXNoPath

Exception for algorithms that should return a path when running on graphs where such a path does not exist.

#### class NetworkXUnbounded

Exception raised by algorithms trying to solve a maximization or a minimization problem instance that is unbounded.

# **TWELVE**

# **UTILITIES**

# 12.1 Helper Functions

Miscellaneous Helpers for NetworkX.

These are not imported into the base networkx namespace but can be accessed, for example, as

```
>>> import networkx
>>> networkx.utils.is_string_like('spam')
True
```

is_string_like(obj)	Check if obj is string.	
flatten(obj[, result])	Return flattened version of (possibly nested) iterable object.	
iterable(obj)	Return True if obj is iterable with a well-defined len().	
is_list_of_ints(intlist)	Return True if list is a list of ints.	
make_str(x)	Return the string representation of t.	
cumulative_sum(numbers)	Yield cumulative sum of numbers.	
<pre>generate_unique_node()</pre>	Generate a unique node label.	
default_opener(filename)	Opens filename using system's default program.	

# 12.1.1 is\_string\_like

```
is_string_like (obj)

Check if obj is string.
```

### 12.1.2 flatten

flatten (obj, result=None)

Return flattened version of (possibly nested) iterable object.

## 12.1.3 iterable

### iterable(obj)

Return True if obj is iterable with a well-defined len().

## 12.1.4 is list of ints

```
is_list_of_ints(intlist)
```

Return True if list is a list of ints.

## 12.1.5 make str

```
make_str(x)
```

Return the string representation of t.

## 12.1.6 cumulative\_sum

```
cumulative_sum (numbers)
```

Yield cumulative sum of numbers.

```
>>> import networkx.utils as utils
>>> list(utils.cumulative_sum([1,2,3,4]))
[1, 3, 6, 10]
```

## 12.1.7 generate\_unique\_node

```
generate_unique_node()
```

Generate a unique node label.

## 12.1.8 default opener

```
default_opener (filename)
```

Opens filename using system's default program.

Parameters filename: str

The path of the file to be opened.

# 12.2 Data Structures and Algorithms

Union-find data structure.

UnionFind.union(\*objects) Find the sets containing the objects and merge them all.

### 12.2.1 union

```
UnionFind.union(*objects)
```

Find the sets containing the objects and merge them all.

# 12.3 Random Sequence Generators

Utilities for generating random numbers, random sequences, and random selections.

<pre>create_degree_sequence(n[, sfunction, max_tries])</pre>	
<pre>pareto_sequence(n[, exponent])</pre>	Return sample sequence of length n from a Pareto distribution.
<pre>powerlaw_sequence(n[, exponent])</pre>	Return sample sequence of length n from a power law distribution.
uniform_sequence(n)	Return sample sequence of length n from a uniform distribution.
<pre>cumulative_distribution(distribution)</pre>	Return normalized cumulative distribution from discrete distribution.
discrete_sequence(n[, distribution,])	Return sample sequence of length n from a given discrete distribution or
<pre>zipf_sequence(n[, alpha, xmin])</pre>	Return a sample sequence of length n from a Zipf distribution with expo
<pre>zipf_rv(alpha[, xmin, seed])</pre>	Return a random value chosen from the Zipf distribution.
random_weighted_sample(mapping, k)	Return k items without replacement from a weighted sample.
weighted_choice(mapping)	Return a single element from a weighted sample.

## 12.3.1 create\_degree\_sequence

create\_degree\_sequence (n, sfunction=None, max\_tries=50, \*\*kwds)

## 12.3.2 pareto\_sequence

#### pareto\_sequence (n, exponent=1.0)

Return sample sequence of length n from a Pareto distribution.

## 12.3.3 powerlaw sequence

#### powerlaw\_sequence (n, exponent=2.0)

Return sample sequence of length n from a power law distribution.

## 12.3.4 uniform\_sequence

#### $uniform_sequence(n)$

Return sample sequence of length n from a uniform distribution.

## 12.3.5 cumulative\_distribution

### cumulative\_distribution (distribution)

Return normalized cumulative distribution from discrete distribution.

## 12.3.6 discrete sequence

### discrete\_sequence (n, distribution=None, cdistribution=None)

Return sample sequence of length n from a given discrete distribution or discrete cumulative distribution.

One of the following must be specified.

distribution = histogram of values, will be normalized

cdistribution = normalized discrete cumulative distribution

## 12.3.7 zipf sequence

```
zipf_sequence(n, alpha=2.0, xmin=1)
```

Return a sample sequence of length n from a Zipf distribution with exponent parameter alpha and minimum value xmin.

#### See also:

```
zipf_rv
```

## 12.3.8 zipf\_rv

```
zipf_rv (alpha, xmin=1, seed=None)
```

Return a random value chosen from the Zipf distribution.

The return value is an integer drawn from the probability distribution ::math:

```
p(x) = \frac{x^{-\alpha}}{x^{-\alpha}} \{ (\alpha, x_{\min}) \},
```

where  $\zeta(\alpha, x_{min})$  is the Hurwitz zeta function.

Parameters alpha: float

Exponent value of the distribution

xmin: int

Minimum value

seed: int

Seed value for random number generator

Returns x: int

Random value from Zipf distribution

#### Raises ValueError:

If xmin < 1 or If alpha <= 1

#### **Notes**

The rejection algorithm generates random values for a the power-law distribution in uniformly bounded expected time dependent on parameters. See [1] for details on its operation.

#### References

..[1] Luc Devroye, Non-Uniform Random Variate Generation, Springer-Verlag, New York, 1986.

### **Examples**

```
>>> nx.zipf_rv(alpha=2, xmin=3, seed=42)
```

## 12.3.9 random weighted sample

## $random\_weighted\_sample(mapping, k)$

Return k items without replacement from a weighted sample.

The input is a dictionary of items with weights as values.

## 12.3.10 weighted choice

```
weighted_choice (mapping)
```

Return a single element from a weighted sample.

The input is a dictionary of items with weights as values.

## 12.4 Decorators

open\_file(path\_arg[, mode]) Decorator to ensure clean opening and closing of files.

## 12.4.1 open\_file

```
open_file (path_arg, mode='r')
```

Decorator to ensure clean opening and closing of files.

#### Parameters path\_arg: int

Location of the path argument in args. Even if the argument is a named positional argument (with a default value), you must specify its index as a positional argument.

mode: str

String for opening mode.

#### Returns \_open\_file : function

Function which cleanly executes the io.

### **Examples**

#### Decorate functions like this:

```
@open_file(0,'r')
def read_function(pathname):
    pass

@open_file(1,'w')
def write_function(G,pathname):
    pass

@open_file(1,'w')
def write_function(G, pathname='graph.dot')
    pass

@open_file('path', 'w+')
def another_function(arg, **kwargs):
```

12.4. Decorators 489

```
path = kwargs['path']
pass
```

# 12.5 Cuthill-Mckee Ordering

Cuthill-McKee ordering of graph nodes to produce sparse matrices

$cuthill\_mckee\_ordering(G[, heuristic])$	Generate an ordering (permutation) of the graph nodes to make a sparse
$\verb"reverse_cuthill_mckee_ordering" (G[, heuristic])$	Generate an ordering (permutation) of the graph nodes to make a sparse

## 12.5.1 cuthill\_mckee\_ordering

```
cuthill_mckee_ordering(G, heuristic=None)
```

Generate an ordering (permutation) of the graph nodes to make a sparse matrix.

Uses the Cuthill-McKee heuristic (based on breadth-first search) [R375].

Parameters G: graph

A NetworkX graph

heuristic: function, optional

Function to choose starting node for RCM algorithm. If None a node from a psuedoperipheral pair is used. A user-defined function can be supplied that takes a graph object and returns a single node.

Returns nodes: generator

Generator of nodes in Cuthill-McKee ordering.

#### See also:

```
reverse_cuthill_mckee_ordering
```

#### **Notes**

The optimal solution the the bandwidth reduction is NP-complete [R376].

### References

[R375], [R376]

#### **Examples**

```
>>> from networkx.utils import cuthill_mckee_ordering
>>> G = nx.path_graph(4)
>>> rcm = list(cuthill_mckee_ordering(G))
>>> A = nx.adjacency_matrix(G, nodelist=rcm)
```

Smallest degree node as heuristic function:

```
>>> def smallest_degree(G):
... node, deg = min(G.degree().items(), key=lambda x: x[1])
... return node
>>> rcm = list(cuthill_mckee_ordering(G, heuristic=smallest_degree))
```

## 12.5.2 reverse\_cuthill\_mckee\_ordering

```
reverse_cuthill_mckee_ordering(G, heuristic=None)
```

Generate an ordering (permutation) of the graph nodes to make a sparse matrix.

Uses the reverse Cuthill-McKee heuristic (based on breadth-first search) [R377].

Parameters G: graph

A NetworkX graph

heuristic: function, optional

Function to choose starting node for RCM algorithm. If None a node from a psuedoperipheral pair is used. A user-defined function can be supplied that takes a graph object and returns a single node.

Returns nodes: generator

Generator of nodes in reverse Cuthill-McKee ordering.

#### See also:

```
cuthill_mckee_ordering
```

#### **Notes**

The optimal solution the the bandwidth reduction is NP-complete [R378].

#### References

```
[R377], [R378]
```

### **Examples**

```
>>> from networkx.utils import reverse_cuthill_mckee_ordering
>>> G = nx.path_graph(4)
>>> rcm = list(reverse_cuthill_mckee_ordering(G))
>>> A = nx.adjacency_matrix(G, nodelist=rcm)
```

Smallest degree node as heuristic function:

```
>>> def smallest_degree(G):
... node, deg = min(G.degree().items(), key=lambda x: x[1])
... return node
>>> rcm = list(reverse_cuthill_mckee_ordering(G, heuristic=smallest_degree))
```

# **12.6 Context Managers**

492 Chapter 12. Utilities

reversed(\*args, \*\*kwds) A context manager for temporarily reversing a directed graph in place.

## 12.6.1 reversed

reversed(\*args, \*\*kwds)

A context manager for temporarily reversing a directed graph in place.

This is a no-op for undirected graphs.

Parameters G: graph

A NetworkX graph.

494 Chapter 12. Utilities

#### **CHAPTER**

## **THIRTEEN**

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496 Chapter 13. License

**CHAPTER** 

# **FOURTEEN**

# **CITING**

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498 Chapter 14. Citing

### FIFTEEN

# **CREDITS**

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Thanks to Guido van Rossum for the idea of using Python for implementing a graph data structure https://www.python.org/doc/essays/graphs

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- Katy Bold contributed the Karate Club graph.
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**CHAPTER** 

## SIXTEEN

## **GLOSSARY**

**dictionary** A Python dictionary maps keys to values. Also known as "hashes", or "associative arrays". See http://docs.python.org/tutorial/datastructures.html#dictionaries

**ebunch** An iteratable container of edge tuples like a list, iterator, or file.

edge Edges are either two-tuples of nodes (u,v) or three tuples of nodes with an edge attribute dictionary (u,v,dict).

edge attribute Edges can have arbitrary Python objects assigned as attributes by using keyword/value pairs when adding an edge assigning to the G.edge[u][v] attribute dictionary for the specified edge u-v.

hashable An object is hashable if it has a hash value which never changes during its lifetime (it needs a \_\_hash\_\_() method), and can be compared to other objects (it needs an \_\_eq\_\_() or \_\_cmp\_\_() method). Hashable objects which compare equal must have the same hash value.

Hashability makes an object usable as a dictionary key and a set member, because these data structures use the hash value internally.

All of Python's immutable built-in objects are hashable, while no mutable containers (such as lists or dictionaries) are. Objects which are instances of user-defined classes are hashable by default; they all compare unequal, and their hash value is their id().

Definition from http://docs.python.org/glossary.html

**nbunch** An nbunch is any iterable container of nodes that is not itself a node in the graph. It can be an iterable or an iterator, e.g. a list, set, graph, file, etc..

**node** A node can be any hashable Python object except None.

**node attribute** Nodes can have arbitrary Python objects assigned as attributes by using keyword/value pairs when adding a node or assigning to the G.node[n] attribute dictionary for the specified node n.

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#### PYTHON MODULE INDEX

```
а
                                          networkx.algorithms.community.kclique,
networkx.algorithms.approximation, 123
networkx.algorithms.approximation.clique,networkx.algorithms.components, 199
                                          networkx.algorithms.components.attracting,
networkx.algorithms.approximation.clustering_coefficient,
                                          networkx.algorithms.components.biconnected,
networkx.algorithms.approximation.dominating_se20,7
                                          networkx.algorithms.components.connected,
networkx.algorithms.approximation.independent_set,
                                          networkx.algorithms.components.semiconnected,
                                                 212
networkx.algorithms.approximation.matching,
                                          networkx.algorithms.components.strongly_connected,
networkx.algorithms.approximation.ramsey,
                                          networkx.algorithms.components.weakly_connected,
       127
networkx.algorithms.approximation.vertex cover. 205
                                          networkx.algorithms.connectivity, 213
       127
                                          networkx.algorithms.connectivity.connectivity,
networkx.algorithms.assortativity, 128
                                                 213
networkx.algorithms.bipartite, 137
                                          networkx.algorithms.connectivity.cuts,
networkx.algorithms.bipartite.basic, 139
                                                 221
networkx.algorithms.bipartite.centrality,
                                          networkx.algorithms.connectivity.stoerwagner,
networkx.algorithms.bipartite.cluster,
                                          networkx.algorithms.connectivity.utils,
                                                 229
networkx.algorithms.bipartite.generators,
                                          networkx.algorithms.core, 230
                                          networkx.algorithms.cycles, 234
networkx.algorithms.bipartite.matching,
                                          networkx.algorithms.dag, 236
\verb"networkx.algorithms.bipartite.projection", \verb"networkx.algorithms.distance\_measures",
networkx.algorithms.bipartite.redundancy, networkx.algorithms.distance_regular,
                                                 241
       154
                                          networkx.algorithms.dominance, 243
networkx.algorithms.bipartite.spectral,
                                          networkx.algorithms.dominating, 244
                                          networkx.algorithms.euler, 245
networkx.algorithms.block, 163
                                          networkx.algorithms.flow, 247
networkx.algorithms.boundary, 164
                                          networkx.algorithms.graphical, 269
networkx.algorithms.centrality, 166
                                          networkx.algorithms.hierarchy, 272
networkx.algorithms.chordal.chordal_alg,
                                          networkx.algorithms.isolate, 273
       187
                                          networkx.algorithms.isomorphism, 274
networkx.algorithms.clique, 190
                                          networkx.algorithms.isomorphism.isomorphvf2,
networkx.algorithms.cluster, 193
networkx.algorithms.coloring, 197
                                          networkx.algorithms.link_analysis.hits_alg,
networkx.algorithms.community, 198
```

```
292
                                          networkx.generators.degree seg, 377
networkx.algorithms.link_analysis.pageranktwogkx.generators.directed, 384
                                          networkx.generators.ego, 392
networkx.algorithms.link_prediction, 295
                                          networkx.generators.expanders, 363
networkx.algorithms.matching, 301
                                          networkx.generators.geometric, 387
networkx.algorithms.mis, 302
                                          networkx.generators.hybrid,390
networkx.algorithms.mst, 303
                                          networkx.generators.intersection, 393
networkx.algorithms.operators.all, 308
                                          networkx.generators.line, 391
networkx.algorithms.operators.binary,
                                          networkx.generators.random clustered,
                                                 383
networkx.algorithms.operators.product,
                                          networkx.generators.random_graphs, 367
      310
                                          networkx.generators.small, 363
networkx.algorithms.operators.unary, 305
                                          networkx.generators.social, 395
networkx.algorithms.richclub, 312
                                          networkx.generators.stochastic, 393
networkx.algorithms.shortest_paths.astar,
       330
networkx.algorithms.shortest_paths.densenetworkx.linalg.algebraicconnectivity,
networkx.algorithms.shortest_paths.genermetworkx.linalg.attrmatrix,405
                                          networkx.linalg.graphmatrix,397
networkx.algorithms.shortest_paths.unweight@drkx.linalg.laplacianmatrix,399
                                          networkx.linalg.spectrum, 401
networkx.algorithms.shortest_paths.weighted,
networkx.algorithms.simple_paths, 331
                                          networkx.readwrite.adjlist, 425
networkx.algorithms.swap, 332
                                          networkx.readwrite.edgelist,432
networkx.algorithms.traversal.breadth_finetwserkcheadwrite.gexf,439
                                          networkx.readwrite.gml, 441
networkx.algorithms.traversal.depth_firsheswsfkk.readwrite.gpickle,445
                                          networkx.readwrite.graph6,456
networkx.algorithms.traversal.edgedfs,
                                          networkx.readwrite.graphml, 447
                                          networkx.readwrite.json_graph,448
networkx.algorithms.tree.branchings, 344
                                          networkx.readwrite.leda, 453
networkx.algorithms.tree.recognition,
                                          networkx.readwrite.multiline adilist,
       341
                                                 429
networkx.algorithms.vitality, 346
                                          networkx.readwrite.nx_shp, 463
                                          networkx.readwrite.nx_yaml, 454
C
                                          networkx.readwrite.pajek, 461
networkx.classes.function, 349
                                          networkx.readwrite.sparse6,459
networkx.convert,411
                                          u
networkx.convert matrix, 414
                                          networkx.utils,485
                                          networkx.utils.contextmanagers, 492
networkx.drawing.layout, 479
                                          networkx.utils.decorators, 489
networkx.drawing.nx agraph, 474
                                          networkx.utils.misc, 485
networkx.drawing.nx_pydot,477
                                          networkx.utils.random_sequence, 486
networkx.drawing.nx_pylab, 465
                                          networkx.utils.rcm, 490
                                          networkx.utils.union_find, 486
networkx.exception, 483
networkx.generators.atlas, 359
networkx.generators.classic, 359
```

514 Python Module Index

Symbols	add_node() (MultiDiGraph method), 92
contains() (DiGraph method), 54	add_node() (MultiGraph method), 66
contains() (Graph method), 27	add_nodes_from() (DiGraph method), 38
contains() (MultiDiGraph method), 111	add_nodes_from() (Graph method), 12
contains() (MultiGraph method), 82	add_nodes_from() (MultiDiGraph method), 93
getitem() (DiGraph method), 51	add_nodes_from() (MultiGraph method), 67
getitem() (Graph method), 24	add_path() (DiGraph method), 43
getitem() (MultiDiGraph method), 108	add_path() (Graph method), 18
getitem() (MultiGraph method), 80	add_path() (MultiDiGraph method), 99
init() (DiGraph method), 36	add_path() (MultiGraph method), 73
init() (DiGraph Matcher method), 281	add_star() (DiGraph method), 43
init() (Edmonds method), 346	add_star() (Graph method), 17
init() (Cannonds method), 340 init() (Graph method), 11	add_star() (MultiDiGraph method), 99
init() (Graph Matcher method), 279	add_star() (MultiGraph method), 73
The state of the s	add_weighted_edges_from() (DiGraph method), 41
init() (MultiDiGraph method), 92	add_weighted_edges_from() (Graph method), 16
init() (MultiGraph method), 66	add_weighted_edges_from() (MultiDiGraph method), 97
iter() (DiGraph method), 46	add_weighted_edges_from() (MultiGraph method), 71
iter() (Graph method), 20	adjacency_data() (in module net-
iter() (MultiDiGraph method), 102	workx.readwrite.json_graph), 450
iter() (MultiGraph method), 76	adjacency_graph() (in module net-
len() (DiGraph method), 56	workx.readwrite.json_graph), 451
len() (Graph method), 28	adjacency_iter() (DiGraph method), 53
len() (MultiDiGraph method), 112	adjacency_iter() (Graph method), 25
len() (MultiGraph method), 84	adjacency_iter() (MultiDiGraph method), 109
A	adjacency_iter() (MultiGraph method), 81
	adjacency_list() (DiGraph method), 52
adamic_adar_index() (in module net-	adjacency_list() (Graph method), 25
workx.algorithms.link_prediction), 297	adjacency_list() (MultiDiGraph method), 109
add_cycle() (DiGraph method), 44	adjacency_list() (MultiGraph method), 80
add_cycle() (Graph method), 18	adjacency_matrix() (in module net-
add_cycle() (MultiDiGraph method), 100	workx.linalg.graphmatrix), 397
add_cycle() (MultiGraph method), 74	adjacency_spectrum() (in module net-
add_edge() (DiGraph method), 40	workx.linalg.spectrum), 402
add_edge() (Graph method), 14	algebraic_connectivity() (in module net-
add_edge() (MultiDiGraph method), 95	workx.linalg.algebraicconnectivity), 402
add_edge() (MultiGraph method), 69	all_neighbors() (in module networkx.classes.function),
add_edges_from() (DiGraph method), 40	352
add_edges_from() (Graph method), 15	
add_edges_from() (MultiDiGraph method), 96	-1 - 3 -1 · · · ·
add_edges_from() (MultiGraph method), 70	workx.algorithms.shortest_paths.weighted), 323
add_node() (DiGraph method), 37	
add node() (Graph method), 11	all_pairs_dijkstra_path_length() (in module net-

workx.algorithms.shortest_paths.weighted),	workx.algorithms.assortativity), 133
324	average_neighbor_degree() (in module net-
all_pairs_node_connectivity() (in module net-	workx.algorithms.assortativity), 132
workx.algorithms.connectivity.connectivity),	average_node_connectivity() (in module net-
214	workx.algorithms.connectivity.connectivity),
all_pairs_shortest_path() (in module net-	213
workx.algorithms.shortest_paths.unweighted),	average_shortest_path_length() (in module net-
319	workx.algorithms.shortest_paths.generic),
all_pairs_shortest_path_length() (in module net-	316
workx.algorithms.shortest_paths.unweighted),	310
319	В
all_shortest_paths() (in module net-	halamand trace() (in madula naturally concentrate alassis)
workx.algorithms.shortest_paths.generic),	balanced_tree() (in module networkx.generators.classic), 360
315	
	barabasi_albert_graph() (in module net-
all_simple_paths() (in module net-	workx.generators.random_graphs), 374
workx.algorithms.simple_paths), 331	barbell_graph() (in module networkx.generators.classic),
alternating_havel_hakimi_graph() (in module net-	360
workx.algorithms.bipartite.generators), 161	bellman_ford() (in module net-
ancestors() (in module networkx.algorithms.dag), 236	workx.algorithms.shortest_paths.weighted),
approximate_current_flow_betweenness_centrality() (in	327
module networkx.algorithms.centrality), 174	betweenness_centrality() (in module net-
articulation_points() (in module net-	workx.algorithms.bipartite.centrality), 157
workx.algorithms.components.biconnected),	betweenness_centrality() (in module net-
211	workx.algorithms.centrality), 169
astar_path() (in module net-	bfs_edges() (in module net-
workx.algorithms.shortest_paths.astar), 330	workx.algorithms.traversal.breadth_first_search),
astar_path_length() (in module net-	338
workx.algorithms.shortest_paths.astar), 331	bfs_predecessors() (in module net-
attr_matrix() (in module networkx.linalg.attrmatrix), 405	workx.algorithms.traversal.breadth_first_search),
attr_sparse_matrix() (in module net-	339
workx.linalg.attrmatrix), 407	bfs_successors() (in module net-
attracting_component_subgraphs() (in module net-	workx.algorithms.traversal.breadth_first_search),
workx.algorithms.components.attracting),	workx.aigoritimis.traversai.oreautii_mst_searcii),
207	
	bfs_tree() (in module net-
attracting_components() (in module net-	workx.algorithms.traversal.breadth_first_search),
workx.algorithms.components.attracting),	338
206	biconnected_component_edges() (in module net-
attribute_assortativity_coefficient() (in module net-	workx.algorithms.components.biconnected),
workx.algorithms.assortativity), 129	209
attribute_mixing_dict() (in module net-	biconnected_component_subgraphs() (in module net-
workx.algorithms.assortativity), 137	workx.algorithms.components.biconnected),
attribute_mixing_matrix() (in module net-	210
workx.algorithms.assortativity), 135	biconnected_components() (in module net-
authority_matrix() (in module net-	workx.algorithms.components.biconnected),
workx.algorithms.link_analysis.hits_alg),	208
295	bidirectional_dijkstra() (in module net-
average_clustering() (in module net-	workx.algorithms.shortest_paths.weighted),
workx.algorithms.approximation.clustering_coef	
124	binomial_graph() (in module net-
average_clustering() (in module net-	workx.generators.random_graphs), 371
workx.algorithms.bipartite.cluster), 152	blockmodel() (in module networkx.algorithms.block),
average_clustering() (in module net-	164
workx.algorithms.cluster), 195	
average_degree_connectivity() (in module net-	branching_weight() (in module net- workx.algorithms.tree.branchings), 344
average degree connectivity) till illudule liet-	workx.argoriumis.tree.branchings), 544

build_auxiliary_edge_connectivity() (in module networkx.algorithms.connectivity.utils), 229	collaboration_weighted_projected_graph() (in module networkx.algorithms.bipartite.projection), 146
build_auxiliary_node_connectivity() (in module net- workx.algorithms.connectivity.utils), 229	color() (in module networkx.algorithms.bipartite.basic),
build_residual_network() (in module net-	common_neighbors() (in module net-
workx.algorithms.flow), 260	workx.classes.function), 353
bull_graph() (in module networkx.generators.small), 365	communicability() (in module net-
buil_graph() (in module networks.generators.smail), 505	workx.algorithms.centrality), 180
C	communicability_betweenness_centrality() (in module
	networkx.algorithms.centrality), 184
candidate_pairs_iter() (DiGraphMatcher method), 282	communicability_centrality() (in module net-
candidate_pairs_iter() (GraphMatcher method), 280	workx.algorithms.centrality), 182
capacity_scaling() (in module networkx.algorithms.flow),	communicability_centrality_exp() (in module net-
267	workx.algorithms.centrality), 183
cartesian_product() (in module net-	communicability_exp() (in module net-
workx.algorithms.operators.product), 310	workx.algorithms.centrality), 181
categorical_edge_match() (in module net- workx.algorithms.isomorphism), 283	complement() (in module net-
	workx.algorithms.operators.unary), 305
categorical_multiedge_match() (in module net- workx.algorithms.isomorphism), 283	complete_bipartite_graph() (in module net-
categorical_node_match() (in module net-	workx.algorithms.bipartite.generators), 159
workx.algorithms.isomorphism), 282	complete_graph() (in module net-
	workx.generators.classic), 361
center() (in module net- workx.algorithms.distance_measures), 239	compose() (in module net-
chordal_graph_cliques() (in module net-	workx.algorithms.operators.binary), 306
workx.algorithms.chordal.chordal_alg), 188	compose_all() (in module net-
chordal_graph_treewidth() (in module net-	workx.algorithms.operators.all), 308
workx.algorithms.chordal.chordal_alg), 189	condensation() (in module net-
chvatal_graph() (in module networkx.generators.small),	workx.algorithms.components.strongly_connected)
365	204
circular_ladder_graph() (in module net-	configuration_model() (in module net-
workx.generators.classic), 361	workx.algorithms.bipartite.generators), 159 configuration_model() (in module net-
circular_layout() (in module networkx.drawing.layout),	workx.generators.degree_seq), 377
480	connected_component_subgraphs() (in module net-
clear() (DiGraph method), 44	workx.algorithms.components.connected),
clear() (Graph method), 19	200
clear() (MultiDiGraph method), 100	connected_components() (in module net-
clear() (MultiGraph method), 74	workx.algorithms.components.connected),
clique_removal() (in module net-	200
workx.algorithms.approximation.clique), 124	connected_double_edge_swap() (in module net-
	workx.algorithms.swap), 333
cliques_containing_node() (in module net- workx.algorithms.clique), 193	connected_watts_strogatz_graph() (in module net-
closeness_centrality() (in module net-	workx.generators.random_graphs), 373
workx.algorithms.bipartite.centrality), 156	copy() (DiGraph method), 62
closeness_centrality() (in module net-	copy() (Graph method), 32
workx.algorithms.centrality), 168	copy() (MultiDiGraph method), 119
•	copy() (MultiGraph method), 88
closeness_vitality() (in module networkx.algorithms.vitality), 347	core_number() (in module networkx.algorithms.core),
• **	230
clustering() (in module net- workx.algorithms.bipartite.cluster), 150	cost_of_flow() (in module networkx.algorithms.flow),
	265
clustering() (in module networkx.algorithms.cluster), 194 cn_soundarajan_hopcroft() (in module net-	could_be_isomorphic() (in module net-
workx.algorithms.link_prediction), 298	workx.algorithms.isomorphism), 276
works.argoriumis.imk_prediction), 276	create degree sequence() (in module net-

workx.utils.random_sequence), 487	density() (in module networkx.classes.function), 349
create_empty_copy() (in module net-	desargues_graph() (in module net-
workx.classes.function), 350	workx.generators.small), 365
cubical_graph() (in module networkx.generators.small),	descendants() (in module networkx.algorithms.dag), 237
365	dfs_edges() (in module net-
cumulative_distribution() (in module net-	workx.algorithms.traversal.depth_first_search),
workx.utils.random_sequence), 487	334
cumulative_sum() (in module networkx.utils.misc), 486	dfs_labeled_edges() (in module net-
current_flow_betweenness_centrality() (in module net- workx.algorithms.centrality), 172	workx.algorithms.traversal.depth_first_search), 337
current_flow_closeness_centrality() (in module net-	dfs_postorder_nodes() (in module net-
workx.algorithms.centrality), 171	workx.algorithms.traversal.depth_first_search),
cuthill_mckee_ordering() (in module net-	336
workx.utils.rcm), 490	dfs_predecessors() (in module net-
cycle_basis() (in module networkx.algorithms.cycles),	workx.algorithms.traversal.depth_first_search),
cycle_graph() (in module networkx.generators.classic),	dfs_preorder_nodes() (in module net-
361	workx.algorithms.traversal.depth_first_search),
D	dfs_successors() (in module net-
davis_southern_women_graph() (in module net-	workx.algorithms.traversal.depth_first_search),
workx.generators.social), 395	335
default_opener() (in module networkx.utils.misc), 486	dfs_tree() (in module net-
degree() (DiGraph method), 56	workx.algorithms.traversal.depth_first_search),
degree() (Graph method), 28	334
degree() (in module networkx.classes.function), 349	diameter() (in module net-
degree() (MultiDiGraph method), 113	workx.algorithms.distance_measures), 239
degree() (MultiGraph method), 84	diamond_graph() (in module networkx.generators.small), 365
degree_assortativity_coefficient() (in module net-	
workx.algorithms.assortativity), 128	dictionary, 501 difference() (in module net-
degree_centrality() (in module net-	difference() (in module net- workx.algorithms.operators.binary), 308
workx.algorithms.bipartite.centrality), 157 degree_centrality() (in module net-	DiGraph() (in module networkx), 35
degree_centrality() (in module net- workx.algorithms.centrality), 167	dijkstra_path() (in module net-
degree_histogram() (in module net-	workx.algorithms.shortest_paths.weighted),
workx.classes.function), 349	321
degree_iter() (DiGraph method), 57	dijkstra_path_length() (in module net-
degree_iter() (Graph method), 29	workx.algorithms.shortest_paths.weighted),
degree_iter() (MultiDiGraph method), 113	321
degree_iter() (MultiGraph method), 85	dijkstra_predecessor_and_distance() (in module net-
degree_mixing_dict() (in module net-	workx.algorithms.shortest_paths.weighted),
workx.algorithms.assortativity), 136	326
degree_mixing_matrix() (in module net-	directed_configuration_model() (in module net-
workx.algorithms.assortativity), 136	workx.generators.degree_seq), 378
degree_pearson_correlation_coefficient() (in module net-	directed_havel_hakimi_graph() (in module net-
workx.algorithms.assortativity), 131	workx.generators.degree_seq), 381
degree_sequence_tree() (in module net-	directed_laplacian_matrix() (in module net-
workx.generators.degree_seq), 382	workx.linalg.laplacianmatrix), 400
degrees() (in module net-	discrete_sequence() (in module net-
workx.algorithms.bipartite.basic), 141	workx.utils.random_sequence), 487
dense_gnm_random_graph() (in module net-	disjoint_union() (in module net-
workx.generators.random_graphs), 369	workx.algorithms.operators.binary), 307 disjoint_union_all() (in module net-
density() (in module net- workx.algorithms.bipartite.basic), 141	workx.algorithms.operators.all), 309
WOLKA, algorithms, other the Justice, 141	monkangonuming.opciatorg.ang.

dispersion() (in module networkx.algorithms.centrality), 187	edge_load() (in module networkx.algorithms.centrality), 186
dodecahedral_graph() (in module net-	edges() (DiGraph method), 46
workx.generators.small), 366	edges() (Graph method), 21
dominance_frontiers() (in module net-	edges() (in module networkx.classes.function), 353
workx.algorithms.dominance), 244	edges() (MultiDiGraph method), 102
dominating_set() (in module net-	edges() (MultiGraph method), 76
workx.algorithms.dominating), 244	edges_iter() (DiGraph method), 47
dorogovtsev_goltsev_mendes_graph() (in module net-	edges_iter() (Graph method), 21
workx.generators.classic), 361	edges_iter() (in module networkx.classes.function), 354
double_edge_swap() (in module net-	edges_iter() (MultiDiGraph method), 103
workx.algorithms.swap), 332	edges_iter() (MultiGraph method), 77
draw() (in module networkx.drawing.nx_pylab), 465	Edmonds (class in networkx.algorithms.tree.branchings),
draw_circular() (in module networkx.drawing.nx_pylab),	346
473	edmonds_karp() (in module networkx.algorithms.flow),
draw_graphviz() (in module net-	254
workx.drawing.nx_pylab), 474	ego_graph() (in module networkx.generators.ego), 392
draw_networkx() (in module net-	eigenvector_centrality() (in module net-
workx.drawing.nx_pylab), 466	workx.algorithms.centrality), 175
draw_networkx_edge_labels() (in module net-	eigenvector_centrality_numpy() (in module net-
workx.drawing.nx_pylab), 472	workx.algorithms.centrality), 176
draw_networkx_edges() (in module net-	empty_graph() (in module networkx.generators.classic),
workx.drawing.nx_pylab), 469	361
draw_networkx_labels() (in module net-	enumerate_all_cliques() (in module net-
workx.drawing.nx_pylab), 471	workx.algorithms.clique), 191
draw_networkx_nodes() (in module net-	eppstein_matching() (in module net-
workx_drawing.nx_pylab), 468	workx.algorithms.bipartite.matching), 142
draw_random() (in module networkx.drawing.nx_pylab),	erdos_renyi_graph() (in module net-
473	workx.generators.random_graphs), 370
draw_shell() (in module networkx.drawing.nx_pylab),	estrada_index() (in module net-
474	workx.algorithms.centrality), 185
draw_spectral() (in module networkx.drawing.nx_pylab),	eulerian_circuit() (in module networkx.algorithms.euler),
473	246
draw_spring() (in module networkx.drawing.nx_pylab),	expected_degree_graph() (in module net-
473	workx.generators.degree_seq), 379
7/3	workx.generators.degree_seq), 377
E	F
ebunch, 501	
	fast_could_be_isomorphic() (in module net-
eccentricity() (in module net-	workx.algorithms.isomorphism), 276
workx.algorithms.distance_measures), 240	fast_gnp_random_graph() (in module net-
edge, 501	workx.generators.random_graphs), 368
edge_betweenness_centrality() (in module net-	faster_could_be_isomorphic() (in module net-
<u> </u>	workx.algorithms.isomorphism), 277
workx.algorithms.centrality), 170	fiedler_vector() (in module net-
edge_boundary() (in module net-	workx.linalg.algebraicconnectivity), 403
workx.algorithms.boundary), 164	find_cliques() (in module networkx.algorithms.clique),
edge_connectivity() (in module net-	191
workx.algorithms.connectivity.connectivity),	find_cycle() (in module networkx.algorithms.cycles), 235
214	find_induced_nodes() (in module net-
edge_current_flow_betweenness_centrality() (in module	workx.algorithms.chordal.chordal_alg), 189
networkx.algorithms.centrality), 173	flatten() (in module networkx.utils.misc), 485
edge_dfs() (in module net-	florentine_families_graph() (in module net-
workx.algorithms.traversal.edgedfs), 340	workx.generators.social), 396

flow_hierarchy()	(in	module	net-	E - E -	et-
_	rithms.hierarcl			workx.classes.function), 355	
floyd_warshall()	(in	module	net-	get_edge_data() (DiGraph method), 50	
	rithms.shortes	t_paths.dense),		get_edge_data() (Graph method), 22	
328		1.1		get_edge_data() (MultiDiGraph method), 106	
floyd_warshall_nump		module	net-	get_edge_data() (MultiGraph method), 78	
•	rithms.shortes	t_paths.dense),			et-
329				workx.classes.function), 355	
floyd_warshall_prede networkx.al		istance() (in n test_paths.dens		global_parameters() (in module n workx.algorithms.distance_regular), 242	et-
329				gn_graph() (in module networkx.generators.directe	d),
freeze() (in module no	etworkx.classe	es.function), 356	5	384	
from_agraph() (in mo 475	dule networkx	x.drawing.nx_ag	graph),	gnc_graph() (in module networkx.generators.directe 385	d),
from_dict_of_dicts()	(in module ne	tworkx.convert)	, 412	gnm_random_graph() (in module n	et-
from_dict_of_lists() (	in module net	workx.convert),	413	workx.generators.random_graphs), 370	
from_edgelist() (in m	odule network	x.convert), 414		gnmk_random_graph() (in module n	et-
from_numpy_matrix(	(in	module	net-	workx.algorithms.bipartite.generators), 163	
workx.conv	ert_matrix), 4	17		gnp_random_graph() (in module n	et-
from_pandas_datafra	me() (in	module	net-	workx.generators.random_graphs), 369	
workx.conv	ert_matrix), 4	22		gnr_graph() (in module networkx.generators.directe	d),
from_pydot() (in mo	odule network	x.drawing.nx_p	oydot),	385	
478				8 8 =	et-
from_scipy_sparse_m workx.conv	natrix() (ir ert_matrix), 4		net-	workx.algorithms.link_analysis.pagerank_alg 291	g),
frucht_graph() (in n	nodule netwo	rkx.generators.s	small),	Graph() (in module networkx), 9	
366				graph_atlas_g() (in module networkx.generators.atla 359	ıs),
G				graph_clique_number() (in module n	et-
general_random_inte	rsection_graph	n() (in module	e net-	workx.algorithms.clique), 192	
workx.gene	rators.intersec	etion), 395		graph_number_of_cliques() (in module n	et-
generate_adjlist() (in	module netwo	rkx.readwrite.a	djlist),	workx.algorithms.clique), 193	
428				graphviz_layout() (in module n	et-
generate_edgelist()	(in	module	net-	workx.drawing.nx_agraph), 476	
	write.edgelist)				et-
generate_gml() (in me	odule network	x.readwrite.gml	l), 444	workx.drawing.nx_pydot), 479	
generate_graph6()	(in	module	net-		et-
workx.read		457		workx.algorithms.tree.branchings), 344	
generate_multiline_ac	•		net-	<i>E</i>	et-
	write.multiline	•		workx.algorithms.coloring), 197	
generate_sparse6()	(in	module	net-	grid_2d_graph() (in module networkx.generators.classi	c),
	write.sparse6)			362	
generate_unique_nod	e() (in module	networkx.utils.	.misc),	grid_graph() (in module networkx.generators.classi	c),
486				362	
generic_edge_match(		module	net-	Н	
	rithms.isomorj		4		
generic_multiedge_m			net-	has_edge() (DiGraph method), 55	
	rithms.isomorj	=	nat	has_edge() (Graph method), 27	
generic_node_match(	) (in rithms.isomorj	module	net-	has_edge() (MultiGraph method), 111	
workx.argo generic_weighted_pro			e net-	has_edge() (MultiGraph method), 83 has_node() (DiGraph method), 54	
		e.projection), 14		has_node() (Graph method), 26	
geographical_thresho		(in module	net-	has_node() (MultiDiGraph method), 110	
	rators.geomet	`	1101	has node() (MultiGraph method), 82	

has_path		in	module	net-	intersection_all()		module	net-
	workx.algori	thms.shorte	est_paths.generic	),	workx.a	lgorithms.ope	erators.all), 310	
	317				intersection_array	r() (in	module	net-
hashable	, 501				workx.a	lgorithms.dist	ance_regular), 2	242
havel ha	kimi_graph()	(in	module	net-	is_aperiodic() (in			
			ite.generators), 1	60	is_arborescence()		module	net-
havel ha	kimi_graph()		module	net-		,	e.recognition), 34	
	workx.gener				is_attracting_com	_	•	
heawood			orkx.generators.	small)	_	•	nponents.attracti	
neuw oou	366	iroddio irotw	orka.generators.	oman,	206	go:::::::::::::::::::::::::::::::::::	пропенияминиен	6),
hits()	(in		nodule	net-	is biconnected()	(in	module	net-
ints()	,		malysis.hits_alg)				nponents.biconn	
	292		marysis.mis_aig)	',	207	ngoriumis.com	iipoliciits.bicoliii	iccicu),
hits_num		(in	module	nat	is_bipartite()	(in	module	nat
iiits_iiuii		*		net-	•	*		net-
	_	umis.mik_a	nnalysis.hits_alg)	),			artite.basic), 139	
1	293	··	1 1		is_bipartite_node_			net-
hits_scip		(in	module	net-			artite.basic), 139	
	_	thms.link_a	analysis.hits_alg)	),	is_branching()	(in	module	net-
	294					•	e.recognition), 34	
hopcroft_	_karp_matchi	•		net-	is_chordal()	(in	module	net-
			ite.matching), 14			•	rdal.chordal_alg	-
house_gr	• " '	odule netw	orkx.generators.	small),	is_connected()	(in	module	net-
	366					llgorithms.con	nponents.connec	cted),
house_x_	_graph() (in n	nodule netw	orkx.generators.	small),	199			
	366				is_digraphical()	(in	module	net-
hub_mat		(in	module	net-	workx.a	lgorithms.gra	phical), 270	
	workx.algori	thms.link_a	nalysis.hits_alg)	),	is_directed() (in n	nodule networ	kx.classes.funct	ion), 350
	295				is_directed_acycli	ic_graph()	(in module	e net-
hypercub	e_graph()	(in	module	net-	workx.a	lgorithms.dag	3), 238	
	workx.gener	ators.classic	2), 362		is_distance_regula	ar() (in	module	net-
_					workx.a	lgorithms.dist	ance_regular), 2	241
					is_dominating_se	-	module	net-
icosahed	ral_graph()	(in	module	net-	workx.a	lgorithms.don	ninating), 245	
reosurrea	workx.gener			1101	is_eulerian() (in n	-	_	uler), 245
immedia	te_dominators			net-	is_forest()	(in	module	net-
mmeara	workx.algori			net		lgorithms.tree	e.recognition), 34	
in degree	e() (DiGraph:				is_frozen() (in mo			
	e() (MultiDiG				is_graphical()			
	e_centrality()	(in	module	net-		lgorithms.gra		
iii_ucgici	workx.algori	*		net-	is_isolate() (in mo			late). 273
in dagra	_		• .		is_isomorphic() (l		•	
_	e_iter() (DiGr	_			is_isomorphic() (	_		
_	e_iter() (Mult	-	ietnoa), 114		is_isomorphic() (	(in	module	net-
	() (DiGraph n		106			`	morphism), 274	net-
_	() (MultiDiGr	_			is_kl_connected()	-	module	nat
_	_iter() (DiGra	-				generators.hyb		net-
_	_iter() (Multil							196
ıncıdence	e_matrix()	(in	module	net-	is_list_of_ints() (i			
	workx.linalg				is_multigraphical		module	net-
			s.function), 350			lgorithms.gra	=	
	() (DiGraphM				is_pseudographic		module	net-
initialize	() (GraphMate		l), 279			lgorithms.gra	•	
intersecti		(in	module	net-	is_semiconnected		module	net-
	workx.algori	thms.operat	tors.binary), 307			llgorithms.con	nponents.semico	onnected),
					2.1.2			

is_string_like() (in module networkx.utils.misc), 485	laplacian_spectrum() (in module net-
is_strongly_connected() (in module net-	workx.linalg.spectrum), 401
workx.algorithms.components.strongly_connecte 202	edatapy_clustering() (in module net- workx.algorithms.bipartite.cluster), 153
is_tree() (in module net-	LCF_graph() (in module networkx.generators.small), 364
workx.algorithms.tree.recognition), 342	lexicographic_product() (in module net-
	• · · · · · · · · · · · · · · · · · · ·
is_valid_degree_sequence_erdos_gallai() (in module net-	workx.algorithms.operators.product), 311
workx.algorithms.graphical), 272	line_graph() (in module networkx.generators.line), 392
is_valid_degree_sequence_havel_hakimi() (in module	literal_destringizer() (in module net-
networkx.algorithms.graphical), 271	workx.readwrite.gml), 444
is_weakly_connected() (in module net-	literal_stringizer() (in module networkx.readwrite.gml),
workx.algorithms.components.weakly_connected	d), 445
205	load_centrality() (in module net-
isolates() (in module networkx.algorithms.isolate), 274	workx.algorithms.centrality), 186
	· · · · · · · · · · · · · · · · · · ·
isomorphisms_iter() (DiGraphMatcher method), 281	, ,
isomorphisms_iter() (GraphMatcher method), 280	workx.algorithms.connectivity.connectivity),
iterable() (in module networkx.utils.misc), 485	216
	local_node_connectivity() (in module net-
J	workx.algorithms.connectivity.connectivity),
jaccard_coefficient() (in module net-	218
workx.algorithms.link_prediction), 296	lollipop_graph() (in module networkx.generators.classic),
workx.aigoriums.mik_prediction), 290	362
K	302
T\	M
k_clique_communities() (in module net-	
workx.algorithms.community.kclique), 198	make_clique_bipartite() (in module net-
k_core() (in module networkx.algorithms.core), 231	workx.algorithms.clique), 192
k_corona() (in module networkx.algorithms.core), 233	make_max_clique_graph() (in module net-
	workx.algorithms.clique), 192
k_crust() (in module networkx.algorithms.core), 232	make_small_graph() (in module net-
k_nearest_neighbors() (in module net-	,
workx.algorithms.assortativity), 134	workx.generators.small), 364
k_random_intersection_graph() (in module net-	make_str() (in module networkx.utils.misc), 486
workx.generators.intersection), 394	match() (DiGraphMatcher method), 282
k_shell() (in module networkx.algorithms.core), 231	match() (GraphMatcher method), 280
karate_club_graph() (in module net-	max_clique() (in module net-
workx.generators.social), 395	workx.algorithms.approximation.clique),
	123
katz_centrality() (in module net-	max_flow_min_cost() (in module net-
workx.algorithms.centrality), 177	workx.algorithms.flow), 266
katz_centrality_numpy() (in module net-	
workx.algorithms.centrality), 179	max_weight_matching() (in module net-
kl_connected_subgraph() (in module net-	workx.algorithms.matching), 302
workx.generators.hybrid), 390	maximal_independent_set() (in module net-
kosaraju_strongly_connected_components()	workx.algorithms.mis), 303
(in module net-	maximal_matching() (in module net-
	1 1 11 201
workx.algorithms.components.strongly_connecte	maximum_branching() (in module net-
204	workx.algorithms.tree.branchings), 345
krackhardt_kite_graph() (in module net-	
workx.generators.small), 366	maximum_flow() (in module networkx.algorithms.flow),
	247
L	maximum_flow_value() (in module net-
	workx.algorithms.flow), 249
ladder_graph() (in module networkx.generators.classic),	maximum_independent_set() (in module net-
362	workx.algorithms.approximation.independent_set)
laplacian_matrix() (in module net- workx.linalg.laplacianmatrix), 399	126

maximum_spanning_arborescence() (in module net-	neighbors() (DiGraph method), 51
workx.algorithms.tree.branchings), 345	neighbors() (Graph method), 23
min_cost_flow() (in module networkx.algorithms.flow),	neighbors() (MultiDiGraph method), 107
264	neighbors() (MultiGraph method), 79
min_cost_flow_cost() (in module net-	neighbors_iter() (DiGraph method), 51
workx.algorithms.flow), 263	neighbors_iter() (Graph method), 24
min_edge_dominating_set() (in module net-	neighbors_iter() (MultiDiGraph method), 107
workx.algorithms.approximation.dominating_se	- · · · · · · · · · · · · · · · · · · ·
125	network_simplex() (in module net-
min_maximal_matching() (in module net-	workx.algorithms.flow), 260
workx.algorithms.approximation.matching),	networkx.algorithms.approximation (module), 123
127	networkx.algorithms.approximation.clique (module), 123
min_weighted_dominating_set() (in module net-	networkx.algorithms.approximation.clustering_coefficient
workx.algorithms.approximation.dominating_se	
125	
	networkx.algorithms.approximation.dominating_set
min_weighted_vertex_cover() (in module net-	(module), 124
workx.algorithms.approximation.vertex_cover),	networkx.algorithms.approximation.independent_set
128	(module), 126
minimum_branching() (in module net-	networkx.algorithms.approximation.matching (module),
workx.algorithms.tree.branchings), 345	126
$minimum\_cut() \ \ (in \ \ module \ \ network x. algorithms. flow),$	networkx.algorithms.approximation.ramsey (module),
250	127
minimum_cut_value() (in module net-	networkx.algorithms.approximation.vertex_cover (mod-
workx.algorithms.flow), 252	ule), 127
minimum_edge_cut() (in module net-	networkx.algorithms.assortativity (module), 128
workx.algorithms.connectivity.cuts), 221	networkx.algorithms.bipartite (module), 137
minimum_node_cut() (in module net-	networkx.algorithms.bipartite.basic (module), 139
workx.algorithms.connectivity.cuts), 223	networkx.algorithms.bipartite.centrality (module), 155
minimum_spanning_arborescence() (in module net-	networkx.algorithms.bipartite.cluster (module), 150
workx.algorithms.tree.branchings), 346	networkx.algorithms.bipartite.generators (module), 158
minimum_spanning_edges() (in module net-	networkx.algorithms.bipartite.matching (module), 142
workx.algorithms.mst), 304	networkx.algorithms.bipartite.projection (module), 144
minimum_spanning_tree() (in module net-	networkx.algorithms.bipartite.redundancy (module), 154
workx.algorithms.mst), 303	networkx.algorithms.bipartite.spectral (module), 149
minimum_st_edge_cut() (in module net-	networkx.algorithms.block (module), 163
workx.algorithms.connectivity.cuts), 224	networkx.algorithms.boundary (module), 164
minimum_st_node_cut() (in module net-	networkx.algorithms.contrality (module), 164
workx.algorithms.connectivity.cuts), 226	· · · · · · · · · · · · · · · · · · ·
• • • • • • • • • • • • • • • • • • • •	networkx.algorithms.chordal.chordal_alg (module), 187
moebius_kantor_graph() (in module net-	networkx.algorithms.clique (module), 190
workx.generators.small), 366	networkx.algorithms.cluster (module), 193
MultiDiGraph() (in module networkx), 91	networkx.algorithms.coloring (module), 197
MultiGraph() (in module networkx), 65	networkx.algorithms.community (module), 198
N	networkx.algorithms.community.kclique (module), 198
IN	networkx.algorithms.components (module), 199
$navigable\_small\_world\_graph() \hspace{0.5cm} (in \hspace{0.5cm} module \hspace{0.5cm} net-$	networkx.algorithms.components.attracting (module),
workx.generators.geometric), 389	206
nbunch, 501	networkx.algorithms.components.biconnected (module),
nbunch_iter() (DiGraph method), 53	207
nbunch_iter() (Graph method), 25	networkx.algorithms.components.connected (module),
nbunch_iter() (MultiDiGraph method), 109	199
nbunch_iter() (MultiGraph method), 81	networkx.algorithms.components.semiconnected (mod-
negative_edge_cycle() (in module net-	ule), 212
workx.algorithms.shortest_paths.weighted),	networkx.algorithms.components.strongly_connected
328	(module), 201

networkx.algorithms.components.weakly_connected	networkx.algorithms.vitality (module), 346
(module), 205	networkx.classes.function (module), 349
networkx.algorithms.connectivity (module), 213	networkx.convert (module), 411
networkx.algorithms.connectivity.connectivity (module),	networkx.convert_matrix (module), 414
213	networkx.drawing.layout (module), 479
networkx.algorithms.connectivity.cuts (module), 221	networkx.drawing.nx_agraph (module), 474
networkx.algorithms.connectivity.stoerwagner (module),	networkx.drawing.nx_pydot (module), 477
228	networkx.drawing.nx_pylab (module), 465
networkx.algorithms.connectivity.utils (module), 229	networkx.exception (module), 483
networkx.algorithms.core (module), 230	networkx.generators.atlas (module), 359
networkx.algorithms.cycles (module), 234	networkx.generators.classic (module), 359
networkx.algorithms.dag (module), 236	networkx.generators.degree_seq (module), 377
networkx.algorithms.distance_measures (module), 239	networkx.generators.directed (module), 384
networkx.algorithms.distance_regular (module), 241	networkx.generators.ego (module), 392
networkx.algorithms.dominance (module), 243	networkx.generators.expanders (module), 363
networkx.algorithms.dominating (module), 244	networkx.generators.geometric (module), 387
networkx.algorithms.euler (module), 245	networkx.generators.hybrid (module), 390
networkx.algorithms.flow (module), 247	networkx.generators.intersection (module), 393
networkx.algorithms.graphical (module), 269	networkx.generators.line (module), 391
networkx.algorithms.hierarchy (module), 272	networkx.generators.random_clustered (module), 383
networkx.algorithms.isolate (module), 272	networkx.generators.random_graphs (module), 367
networkx.algorithms.isomorphism (module), 274	networkx.generators.small (module), 363
networkx.algorithms.isomorphism.isomorphyf2 (mod-	networkx.generators.social (module), 395
ule), 277	networkx.generators.stochastic (module), 393
networkx.algorithms.link_analysis.hits_alg (module),	networkx.linalg.algebraicconnectivity (module), 402
292 (module),	networkx.linalg.attrmatrix (module), 405
networkx.algorithms.link_analysis.pagerank_alg (mod-	networkx.linalg.graphmatrix (module), 403
ule), 287	networkx.linalg.laplacianmatrix (module), 397
networkx.algorithms.link_prediction (module), 295	networks.linalg.spectrum (module), 401
networkx.algorithms.matching (module), 301	networks.readwrite.adjlist (module), 425
networkx.algorithms.mis (module), 302	networkx.readwrite.edgelist (module), 432
networkx.algorithms.mst (module), 303	networks.readwrite.gexf (module), 439
networkx.algorithms.operators.all (module), 308	networkx.readwrite.gml (module), 441
networkx.algorithms.operators.binary (module), 305	networkx.readwrite.gpickle (module), 445
networkx.algorithms.operators.product (module), 310	networkx.readwrite.graph6 (module), 456
networkx.algorithms.operators.unary (module), 305	networkx.readwrite.graphml (module), 447
networkx.algorithms.richclub (module), 312	networkx.readwrite.json_graph (module), 448
networkx.algorithms.shortest_paths.astar (module), 330	networkx.readwrite.leda (module), 453
networkx.algorithms.shortest_paths.dense (module), 328	networkx.readwrite.multiline_adjlist (module), 429
networkx.algorithms.shortest_paths.generic (module),	networkx.readwrite.nx_shp (module), 463
313	networkx.readwrite.nx_yaml (module), 454
networkx.algorithms.shortest_paths.unweighted (mod-	networkx.readwrite.pajek (module), 461
ule), 317	networkx.readwrite.sparse6 (module), 459
networkx.algorithms.shortest_paths.weighted (module),	networkx.utils (module), 485
320	networkx.utils.contextmanagers (module), 492
networkx.algorithms.simple_paths (module), 331	networkx.utils.decorators (module), 489
networkx.algorithms.swap (module), 332	networkx.utils.misc (module), 485
networkx.algorithms.traversal.breadth_first_search (mod-	networkx.utils.random_sequence (module), 486
ule), 338	networkx.utils.rcm (module), 490
networkx.algorithms.traversal.depth_first_search (mod-	networkx.utils.union_find (module), 486
ule), 334	NetworkXAlgorithmError (class in networkx), 483
networkx.algorithms.traversal.edgedfs (module), 340	NetworkXError (class in networkx), 483
networkx.algorithms.tree.branchings (module), 344	NetworkXException (class in networkx), 483
networks algorithms tree recognition (module) 341	NetworkXNoPath (class in networkx) 483

NetworkXPointlessConcept (class in networkx), 483	number_of_edges() (Graph method), 30
NetworkXUnbounded (class in networkx), 483	number_of_edges() (in module net-
NetworkXUnfeasible (class in networkx), 483	workx.classes.function), 353
newman_watts_strogatz_graph() (in module net-	number_of_edges() (MultiDiGraph method), 117
workx.generators.random_graphs), 371	number_of_edges() (MultiGraph method), 86
node, <b>501</b>	number_of_nodes() (DiGraph method), 56
node attribute, 501	number_of_nodes() (Graph method), 28
node_boundary() (in module net-	number_of_nodes() (in module net-
workx.algorithms.boundary), 165	workx.classes.function), 352
node_clique_number() (in module net-	number_of_nodes() (MultiDiGraph method), 112
workx.algorithms.clique), 193	number_of_nodes() (MultiGraph method), 84
node_connected_component() (in module net-	number_of_selfloops() (DiGraph method), 62
workx.algorithms.components.connected),	number_of_selfloops() (Graph method), 32
201	number_of_selfloops() (MultiDiGraph method), 118
node_connectivity() (in module net-	number_of_selfloops() (MultiGraph method), 88
workx.algorithms.connectivity.connectivity),	number_strongly_connected_components()
220	(in module net-
node_link_data() (in module net-	workx.algorithms.components.strongly_connected)
workx.readwrite.json_graph), 449	202
C 1 V	number_weakly_connected_components() (in module net-
workx.readwrite.json_graph), 450	· ·
node_redundancy() (in module net-	workx.algorithms.components.weakly_connected),
workx.algorithms.bipartite.redundancy),	205
155	numeric_assortativity_coefficient() (in module net-
nodes() (DiGraph method), 45	workx.algorithms.assortativity), 130
nodes() (Graph method), 19	numerical_edge_match() (in module net-
nodes() (in module networkx.classes.function), 352	workx.algorithms.isomorphism), 284
nodes() (MultiDiGraph method), 101	numerical_multiedge_match() (in module net-
nodes() (MultiGraph method), 75	workx.algorithms.isomorphism), 285
nodes_iter() (DiGraph method), 45	numerical_node_match() (in module net-
nodes_iter() (Graph method), 20	workx.algorithms.isomorphism), 284
nodes_iter() (in module networkx.classes.function), 352	
nodes_iter() (MultiDiGraph method), 101	0
nodes_iter() (MultiGraph method), 75	octahedral_graph() (in module net-
nodes_with_selfloops() (DiGraph method), 61	workx.generators.small), 367
nodes_with_selfloops() (Graph method), 31	open_file() (in module networkx.utils.decorators), 489
nodes_with_selfloops() (MultiDiGraph method), 117	order() (DiGraph method), 55
nodes_with_selfloops() (MultiGraph method), 87	order() (Graph method), 28
non_edges() (in module networkx.classes.function), 354	order() (MultiDiGraph method), 112
non_neighbors() (in module networkx.classes.function),	order() (MultiGraph method), 83
352	out_degree() (DiGraph method), 59
normalized_laplacian_matrix() (in module net-	out_degree() (MultiDiGraph method), 115
workx.linalg.laplacianmatrix), 399	out_degree_centrality() (in module net-
<pre>null_graph() (in module networkx.generators.classic),</pre>	workx.algorithms.centrality), 168
363	out_degree_iter() (DiGraph method), 59
number_attracting_components() (in module net-	out_degree_iter() (MultiDiGraph method), 115
workx.algorithms.components.attracting),	out_edges() (DiGraph method), 48
206	out_edges() (MultiDiGraph method), 104
number_connected_components() (in module net-	out_edges_iter() (DiGraph method), 49
workx.algorithms.components.connected),	out_edges_iter() (MultiDiGraph method), 105
200	overlap_weighted_projected_graph() (in module net-
number_of_cliques() (in module net-	workx.algorithms.bipartite.projection), 147
workx.algorithms.clique), 193	works.argorumis.orparute.projection), 14/
number_of_edges() (DiGraph method), 60	
— — U V N I "// " "	

P	pygraphviz_layout() (in module net-
pagerank() (in module net-	workx.drawing.nx_agraph), 477
workx.algorithms.link_analysis.pagerank_alg), 288	R
pagerank_numpy() (in module net- workx.algorithms.link_analysis.pagerank_alg),	ra_index_soundarajan_hopcroft() (in module net- workx.algorithms.link_prediction), 299
289 pagerank_scipy() (in module net-	radius() (in module net- workx.algorithms.distance_measures), 240
workx.algorithms.link_analysis.pagerank_alg),	ramsey_R2() (in module net-
290	workx.algorithms.approximation.ramsey),
pappus_graph() (in module networkx.generators.small),	127
367 pareto_sequence() (in module net-	random_clustered_graph() (in module net- workx.generators.random_clustered), 383
workx.utils.random_sequence), 487	random_degree_sequence_graph() (in module net-
parse_adjlist() (in module networkx.readwrite.adjlist),	workx.generators.degree_seq), 382
427	random_geometric_graph() (in module net-
parse_edgelist() (in module networkx.readwrite.edgelist), 438	workx.generators.geometric), 387 random_graph() (in module net-
parse_gml() (in module networkx.readwrite.gml), 443	workx.algorithms.bipartite.generators), 162
parse_graph6() (in module networkx.readwrite.graph6), 456	random_layout() (in module networkx.drawing.layout), 480
parse_leda() (in module networkx.readwrite.leda), 454	random_lobster() (in module net-
parse_multiline_adjlist() (in module net-	workx.generators.random_graphs), 375
workx.readwrite.multiline_adjlist), 431 parse_pajek() (in module networkx.readwrite.pajek), 463	random_powerlaw_tree() (in module net- workx.generators.random_graphs), 376
parse_sparse6() (in module networkx.readwrite.sparse6),	random_powerlaw_tree_sequence() (in module net-
459	workx.generators.random_graphs), 376
path_graph() (in module networkx.generators.classic), 363	random_regular_graph() (in module net- workx.generators.random_graphs), 373
periphery() (in module net-	random_shell_graph() (in module net-
workx.algorithms.distance_measures), 240	workx.generators.random_graphs), 375
petersen_graph() (in module networkx.generators.small), 367	random_weighted_sample() (in module net- workx.utils.random_sequence), 489
powerlaw_cluster_graph() (in module net-	read_adjlist() (in module networkx.readwrite.adjlist), 425
workx.generators.random_graphs), 374	read_dot() (in module networkx.drawing.nx_agraph), 476
powerlaw_sequence() (in module net-	read_dot() (in module networkx.drawing.nx_pydot), 478
workx.utils.random_sequence), 487	read_edgelist() (in module networkx.readwrite.edgelist),
predecessor() (in module net- workx.algorithms.shortest_paths.unweighted),	read_gexf() (in module networkx.readwrite.gexf), 439
320	read_gml() (in module networkx.readwrite.gml), 442
predecessors() (DiGraph method), 52	read_gpickle() (in module networkx.readwrite.gpickle),
predecessors() (MultiDiGraph method), 108	445
predecessors_iter() (DiGraph method), 52	read_graph6() (in module networkx.readwrite.graph6),
predecessors_iter() (MultiDiGraph method), 108	457 read_graphml() (in module networkx.readwrite.graphml),
preferential_attachment() (in module net- workx.algorithms.link_prediction), 297	447
preferential_attachment_graph() (in module net-	read_leda() (in module networkx.readwrite.leda), 454
workx.algorithms.bipartite.generators), 161	read_multiline_adjlist() (in module net-
preflow_push() (in module networkx.algorithms.flow),	workx.readwrite.multiline_adjlist), 429
258	read_pajek() (in module networkx readwrite pajek), 462
projected_graph() (in module net- workx.algorithms.bipartite.projection), 144	read_shp() (in module networkx.readwrite.nx_shp), 463 read_sparse6() (in module networkx.readwrite.sparse6),
pydot_layout() (in module networkx.drawing.nx_pydot), 479	460

read_weighted_edgelist() (in module ne workx.readwrite.edgelist), 435	et-	sets() (in module networkx.algorithms.bipartite.basic),
read_yaml() (in module networkx.readwrite.nx_yaml	1),	shell_layout() (in module networkx.drawing.layout), 481
455		shortest_augmenting_path() (in module net-
relabel_gexf_graph() (in module ne		workx.algorithms.flow), 256
workx.readwrite.gexf), 441		shortest_path() (in module net-
remove_edge() (DiGraph method), 42		workx.algorithms.shortest_paths.generic),
remove_edge() (Graph method), 16		314
remove_edge() (MultiDiGraph method), 97		shortest_path_length() (in module net-
remove_edge() (MultiGraph method), 71		workx.algorithms.shortest_paths.generic),
remove_edges_from() (DiGraph method), 42		315
remove_edges_from() (Graph method), 17		simple_cycles() (in module networkx.algorithms.cycles),
remove_edges_from() (MultiDiGraph method), 98		234
remove_edges_from() (MultiGraph method), 72		single_source_dijkstra() (in module net-
remove_node() (DiGraph method), 39		workx.algorithms.shortest_paths.weighted),
remove_node() (Graph method), 13		324
remove_node() (MultiDiGraph method), 94		single_source_dijkstra_path() (in module net-
remove_node() (MultiGraph method), 68		workx.algorithms.shortest_paths.weighted),
remove_nodes_from() (DiGraph method), 39		322
remove_nodes_from() (Graph method), 13		single_source_dijkstra_path_length() (in module net-
remove_nodes_from() (MultiDiGraph method), 94		workx.algorithms.shortest_paths.weighted),
remove_nodes_from() (MultiGraph method), 68		323
resource_allocation_index() (in module ne	et-	single_source_shortest_path() (in module net-
workx.algorithms.link_prediction), 295		workx.algorithms.shortest_paths.unweighted),
reverse() (DiGraph method), 65		318
reverse() (in module ne	et-	single_source_shortest_path_length() (in module net-
workx.algorithms.operators.unary), 305		workx.algorithms.shortest_paths.unweighted),
reverse() (MultiDiGraph method), 121		318
reverse_cuthill_mckee_ordering() (in module ne		size() (DiGraph method), 60
workx.utils.rcm), 491		size() (Graph method), 30
reverse_havel_hakimi_graph() (in module ne		size() (MultiDiGraph method), 116
workx.algorithms.bipartite.generators), 160		size() (MultiGraph method), 85
reversed() (in module networkx.utils.contextmanagers	s),	spectral_bipartivity() (in module net-
493		workx.algorithms.bipartite.spectral), 150
rich_club_coefficient() (in module ne workx.algorithms.richclub), 312		spectral_layout() (in module networkx.drawing.layout), 482
robins_alexander_clustering() (in module ne	et-	spectral_ordering() (in module net-
workx.algorithms.bipartite.cluster), 154		workx.linalg.algebraicconnectivity), 404
S		spring_layout() (in module networkx.drawing.layout),
		481
scale_free_graph() (in module ne	et-	square_clustering() (in module net-
workx.generators.directed), 386		workx.algorithms.cluster), 196
sedgewick_maze_graph() (in module ne		star_graph() (in module networkx.generators.classic), 363
workx.generators.small), 367		stochastic_graph() (in module net-
selfloop_edges() (DiGraph method), 61		workx.generators.stochastic), 393
selfloop_edges() (Graph method), 31		stoer_wagner() (in module net-
selfloop_edges() (MultiDiGraph method), 118		workx.algorithms.connectivity.stoerwagner),
selfloop_edges() (MultiGraph method), 87		228
semantic_feasibility() (DiGraphMatcher method), 282		strong_product() (in module net-
semantic_feasibility() (GraphMatcher method), 280		workx.algorithms.operators.product), 311 strongly_connected_component_subgraphs()
set_edge_attributes() (in module ne	et-	(in module net-
workx.classes.function), 355	n+	workx.algorithms.components.strongly_connected)
set_node_attributes() (in module ne workx.classes.function), 354	t-	203
WOLKA CLASSES THREHOLD 1. 1.14		

strongly_connected_components() (in module net-	
workx.algorithms.components.strongly_connecte	edn_vertex_cover() (in module net-
202	workx.algorithms.bipartite.matching), 143
strongly_connected_components_recursive()	topological_sort() (in module networkx.algorithms.dag),
(in module net-	237
workx.algorithms.components.strongly_connecte	edopological_sort_recursive() (in module net-
203	workx.algorithms.dag), 238
subgraph() (DiGraph method), 64	transitivity() (in module networkx.algorithms.cluster),
subgraph() (Graph method), 34	194
subgraph() (MultiDiGraph method), 121	tree_data() (in module networkx.readwrite.json_graph),
subgraph() (MultiGraph method), 90	452
subgraph_is_isomorphic() (DiGraphMatcher method),	tree_graph() (in module networkx.readwrite.json_graph),
281	453
subgraph_is_isomorphic() (GraphMatcher method), 280	triangles() (in module networkx.algorithms.cluster), 193
subgraph_isomorphisms_iter() (DiGraphMatcher	trivial_graph() (in module networkx.generators.classic),
method), 282	363
subgraph_isomorphisms_iter() (GraphMatcher method),	truncated_cube_graph() (in module net-
subgraph_isomorphisms_iter() (Graphiviatener method),	workx.generators.small), 367
successors() (DiGraph method), 52	= = = = = :
successors() (MultiDiGraph method), 108	workx.generators.small), 367
successors_iter() (DiGraph method), 52	tutte_graph() (in module networkx.generators.small), 367
successors_iter() (MultiDiGraph method), 108	U
symmetric_difference() (in module net-	
workx.algorithms.operators.binary), 308	uniform_random_intersection_graph() (in module net-
syntactic_feasibility() (DiGraphMatcher method), 282	workx.generators.intersection), 394
syntactic_feasibility() (GraphMatcher method), 280	uniform_sequence() (in module net-
т	workx.utils.random_sequence), 487
Т	union() (in module net-
tensor_product() (in module net-	workx.algorithms.operators.binary), 306
workx.algorithms.operators.product), 312	union() (UnionFind method), 486
tetrahedral_graph() (in module net-	union_all() (in module net-
workx.generators.small), 367	workx.algorithms.operators.all), 309
to_agraph() (in module networkx.drawing.nx_agraph),	
475	W
to_dict_of_dicts() (in module networkx.convert), 412	watts_strogatz_graph() (in module net-
to_dict_of_lists() (in module networkx.convert), 413	workx.generators.random_graphs), 372
to_directed() (DiGraph method), 63	waxman_graph() (in module net-
to_directed() (Graph method), 33	workx.generators.geometric), 389
to_directed() (MultiDiGraph method), 120	weakly_connected_component_subgraphs()
to_directed() (MultiGraph method), 89	(in module net-
to_edgelist() (in module networkx.convert), 413	workx.algorithms.components.weakly_connected)
to_networkx_graph() (in module networkx.convert), 411	205
to_numpy_matrix() (in module net-	
workx.convert_matrix), 415	weakly_connected_components() (in module net-
to_numpy_recarray() (in module net-	workx.algorithms.components.weakly_connected)
workx.convert_matrix), 416	205
	weighted_choice() (in module net-
to_pandas_dataframe() (in module net- workx.convert_matrix), 421	workx.utils.random_sequence), 489
<del>-</del> , , ,	weighted_projected_graph() (in module net-
to_pydot() (in module networkx.drawing.nx_pydot), 478	workx.algorithms.bipartite.projection), 145
to_scipy_sparse_matrix() (in module net-	wheel_graph() (in module networkx.generators.classic),
workx.convert_matrix), 418	363
to_undirected() (DiGraph method), 63	within_inter_cluster() (in module net-
to_undirected() (Graph method), 33 to_undirected() (MultiDiGraph method), 119	workx.algorithms.link_prediction), 300

```
write_adjlist() (in module networkx.readwrite.adjlist),
         426
write_dot() (in module networkx.drawing.nx_agraph),
         476
write_dot() (in module networkx.drawing.nx_pydot), 478
write edgelist() (in module networkx.readwrite.edgelist),
write_gexf() (in module networkx.readwrite.gexf), 440
write gml() (in module networkx.readwrite.gml), 442
write_gpickle() (in module networkx.readwrite.gpickle),
write_graph6() (in module networkx.readwrite.graph6),
         458
write_graphml()
                                   module
                        (in
                                                   net-
         workx.readwrite.graphml), 448
write_multiline_adjlist()
                             (in
                                      module
                                                   net-
         workx.readwrite.multiline_adjlist), 430
write pajek() (in module networkx.readwrite.pajek), 462
write_shp() (in module networkx.readwrite.nx_shp), 464
write sparse6() (in module networkx.readwrite.sparse6),
         461
write_weighted_edgelist()
                              (in
                                      module
         workx.readwrite.edgelist), 436
write yaml() (in module networkx.readwrite.nx yaml),
         455
Ζ
zipf rv() (in module networkx.utils.random sequence),
zipf_sequence()
                       (in
                                   module
                                                   net-
         workx.utils.random_sequence), 488
```