

## Summary

Executing `sna 6-18.py` will produce an author citation network named G1 along with variables listed in the table below. The edges of G1 are stored in a list of lists named `edge`, where each list `<- [Source ID, Destination ID, # of Citations, Edge ID]`.

A one hop network (G2) may also be produced by running function `onehop` and passing the returned object to function `onehopnetwork`. All nodes in network G1 are integer IDs and all google scholar (authors) have respective string IDs. Dictionaries `userdict` and `nodedict` will “translate” from an author’s string ID to the author’s integer ID (vice-versa). The list of lists named `edge` contains all the edges of the author citation network.

The contents of this repository (6-18-2019) include `sna 6-18.py`, the workspace of an execution in `spyder` (Python 2.7), dictionaries to go between string/integer IDs for authors, a dropbox link to a data dump of Breadth-First-Search graph files, author citation network edges, and `hopsatnodes`. `Hopsatnodes` contains for each node, the number of connections a hop distance away from that node.

Functions & network objects in `sna 6-18.py` rely on SNAP (Stanford Network Analysis Project) which requires Python 2.7.x and requirements may vary.

SNAP: <http://snap.stanford.edu/>

### **Generated Variables & Objects**

Name	Type	Size	Description
<code>DstNId</code>	<code>Int</code>	1	Destination Node
<code>SrcNId</code>	<code>Int</code>	1	Source Node
<code>destnodes</code>	<code>list</code>	794	nodes being cited
<code>directory</code>	<code>str</code>	1	dropbox path
<code>edge</code>	<code>list</code>	1395205	author citation network edges
<code>file</code>	<code>dict</code>	2892	current indexed dataset from jsons
<code>filepaths</code>	<code>list</code>	794	file names of datasets
<code>files</code>	<code>dict</code>	2892	---
<code>i</code>	<code>str</code>	1	index
<code>jsonindex</code>	<code>int</code>	1	index
<code>jsons</code>	<code>list</code>	794	author citation data
<code>key</code>	<code>unicode</code>	1	index
<code>line</code>	<code>str</code>	1	index
<code>node</code>	<code>int</code>	1	index
<code>nodedict</code>	<code>dict</code>	433638	dictionary
<code>nodeid</code>	<code>int</code>	1	number of unique authors

output	str	1	output path
path	str	1	index
reject	int	1	count of non-unique authors when building userdict
row	int	1	index
rowindex	int	1	index
self_cite	list	794	self citations
self_cite_ratio	list	794	self citation ratios
short	unicode	1	current indexed author ID
size	int	1	total number of edges
sizeindex	int	1	index
sum_cite	list	794	total citations of an author
userdict	dict	433638	dictionary
weights	list	2892	weights of currently indexed dataset

**Table 1**

## Functions

**self\_citation** :finds instances of self citations and return a list of lists where each list is of the form: [nodeid, # of self citations]

### Inputs

- destnodes: destination nodes, typically user IDs named in the file name of author citation datasets "jsons"
- edges: network edges (list of lists)
- jsons: list containing raw json data
- size: number of edges in network

### Function Call

self\_cite = self\_citation(destnodes,edge,jsons,size)

**sum\_citation**: finds the total number of citations by each author and returns a list of lists where each list is of the form: [nodeid, total # of citations]

### Inputs

- destnodes: destination nodes, typically user IDs named in the file name of author citation datasets "jsons"
- edges: network edges (list of lists)
- size: number of edges in network

### Function Call

sum\_cite = sum\_citation(destnodes,edge,size)

**self\_citation\_ratio**: returns the self citation ratio of each author in a list of lists where each list is of the form: [userid, self citation ratio]

### Inputs

- nodedict: dictionary of node IDs to user IDs
- self\_cite: returned list from self\_citation function
- sum\_cite: returned list from sum\_citation function

### Function Call

self\_cite\_ratio = self\_citation\_ratio(nodedict, self\_cite,sum\_cite)

**printOutDegV**: prints outdegree for each node in the network

Inputs: None

**printNumNodes**: this function prints the results of how many unique/non-unique authors are added to userdict (userID to nodeID dictionary) after building userdict

**sortmatrix**: returns a sorted matrix in ascending order by col, uses timsort

### Inputs

- matrix: typically a list of lists
- col: column of matrix to sort in ascending order

### Function Call

sortedmatrix = sortmatrix(matrix,col)

**asscedge** : creates an associative array mapping node IDS to edges associated with respective IDs

i.e. node ID 0 -> edge[0:5]

Inputs

- sorteditem: a presorted matrix with edges sorted in ascending order
- col: col number to use for mapping

Function Call

asscSrc = asscedge(edgeSrc,0)

**onehop** : generates a matrix containing one hop edges

Inputs

edge: matrix with a row composed of: Source node, Destination Node, No. of Citations, Edge ID

userdict: dictionary converting user IDs to integer node IDs

Function Call

\*onestep = onehop(edge, userdict)

**onehopnetwork** : generates a SNAP network object containing one hop edges

Inputs

- onestep: matrix containing onehop edges

function onehopnetwork must accept function onehopnetwork's return value

Function Call

G2 = onehopnetwork(onestep)

**getbfs** : Returns a directed Breadth-First-Search tree with root at startnode

Inputs

- G1: network
- nodekeys: keys of nodedict -> \*nodedict.keys()
- search(optional): if 0 -> generate 1 tree at startnode = nodeid  
if 1-> generate trees for all node ids
- startnode(optional) : if search = 0: choose a root for the bfs tree using startnode  
Do not use if search = 1
- FollowOut(Bool): graph constructed by following outward links
- FollowIn(Bool): graph constructed by following inward links

Function Call

bfstrees = getbfs(G1,nodekeys,search=0,startnode=0,FollowOut=True, FollowIn=False)  
nodekeys = \*nodedict.keys()

**getnodesathops**: generates a dictionary of dictionaries with the outer dictionary being a node key/ user id and the inner dictionary containing the number of nodes at hop distance i where i is an integer.

Inputs

- G1: G1 is a snap network object
- nodedict: dictionary of node IDs (int) to user IDs (string)
- keychoice: 0 for node integer IDs, 1 for user string IDs

#### Function Call

hopobject = getnodesathops(G1,nodedict)

**dictdump** : dumps the contents of a dictionary into a json file

#### Inputs

- item: should be a dictionary
- filename: should be a file name as a string with .json ending

#### Function Call

temp = dictdump(python\_dictionary, 'contents.json')

**matrixdump**: will dump the contents of a matrix (list of lists) into a csv file

#### Inputs

- matrix: item should be a list of lists or matrix
- filename: should be a file name as a string with .csv ending

example: temp = matrixdump(list of lists, 'contents.csv')

#### Note:

- *Functions typically accept generated variables, named in Table 1, as input*
- *Any mention of matrix refers to a list of lists.*
- *Some objects/variables produced by the functions above are not immediately generated and must be called separately.*
- *bfs trees dropbox link: [https://www.dropbox.com/sh/54jwmzw5o4vdqfd/AABOXP-W0bw0LI-XFNR\\_ui5Ja?dl=0](https://www.dropbox.com/sh/54jwmzw5o4vdqfd/AABOXP-W0bw0LI-XFNR_ui5Ja?dl=0)*
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