## Module IV - Basic Analysis

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Loading data from multiple sources

- Local network data files
- Connecting to a database
- Building directly from the Internet

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- Calculating degree distribution
- ▶ Plotting statistics using matplotlib
- Calculating cliques, clustering and transitivity

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#### Basic visualization

- Review of NetworkX's plotting algorithms
- Adding analysis to visualization

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### Let's try!

- ▶ We will load the edge list of Hartford drug users network
- ▶ Specify that the network be a directed graph, and the nodes be integers
- ▶ Use info() to check that data has been loaded correctly

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### Let's try!

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It's time to fire up your console and load Python!

### Starting NetworkX and loading data

```
>>> from networkx import *
>>> hartford=read_edgelist("../../data/hartford_drug.txt",create_using=DiGraph(),nodetype=int)
>>> info(hartford)
```

#### Name:

Type: DiGraph
Number of nodes: 212
Number of edges: 337
Average in degree: 1.5896
Average out degree: 1.5896

Local network data

## Loading the Hartford drug users network

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#### What did we just do?

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#### What did we just do?

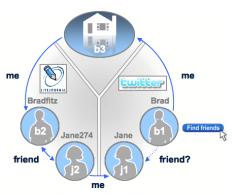
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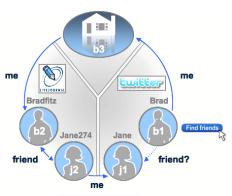
Some formats may have more or less options, always check the documentations!

## Building a network from a database

As data sets become larger and persistently changing, it may make more sense to store them in a database rather than a single file

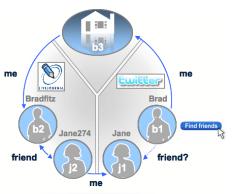
 As we have seen, Python provides binding to many modern database frameworks



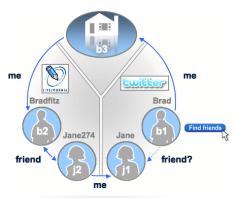


Perhaps the most powerful aspect of NetworkX is its ability to work in Python to generate networks from live-streaming data

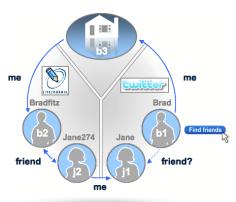
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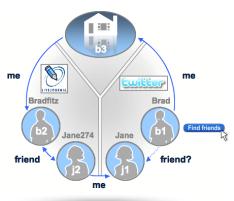
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    - Seed: imichaeldotorg.livejournal.com
    - k = 3



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- Through a process called "k-snowball searching" seed → friend → · · · → friend<sub>k</sub>
  - Seed: imichaeldotorg.livejournal.comk = 3
- ► Note the low value of k

### The code, part 1

#### Loading the libraries and setting things up

```
from cjson import *
from untlib import *
from networkx import *
from scipy import array,unique
...
if __name__ == "__main__":
seed_url='(http://imichaeldotorg.livejournal.com"
sg=get_sg(seed_url)
net_newnodes=create_egonet(sg)
info(net)
```

#### Get the JSON from SocialGraph

```
def get_sg(seed_url):
    sgapi_url="http://socialgraph.apis.google.com/lookup?q="+seed_url+"&edo=l&edi=l&fme=l&pretty=0"
    try:
        furl=urlopen(sgapi_url)
        fr=furl.read()
        furl.close()
        return fr
    except lOError:
        print "Could not connect to website"
        print sgapi_url
        return
```

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from cjson import *
from urllib import *
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    seed_url='('http://imichaeldotorg.livejournal.com"
    sgrget_sg(seed_url)
    net,newnodes=create_egonet(sg)
    info(net)
```

```
Name: ['http://imichaeldotorg.livejournal.com/']
Type: DiGraph
Number of nodes: 5
Number of edges: 5
Average in degree: 1.0
Average out degree: 1.0
```

#### Get the JSON from SocialGraph

```
def get_sg(seed_url):
    sgapi_url="http://socialgraph.apis.google.com/lookup?q="+seed_url+"&edo=1&edi=1&fme=1&pretty=0"
    try:
        furl=urlopen(sgapi_url)
        fr=furl.read()
        furl.close()
        return fr
    except IOError:
        print "Could not connect to website"
        print sgapi_url
    return
```

## Build egonet and snowball

#### Creating the egonet

```
def create_egonet(s):
    try:
        raw=decode(s)
        G=DiGraph()
        pendants=[]
        n=raw['nodes']
        nk=n.kevs()
        G.name=str(nk)
        pendants=[]
        for a in range(0.len(nk)):
            for b in range(0,len(nk)):
                if al=h·
                    G.add_edge(nk[a],nk[b])
        for k in nk:
            ego=n[k]
            ego out=ego['nodes referenced']
            for o in ego_out:
                 G.add_edge(k,o)
                pendants.append(o)
            ego_in=ego['nodes_referenced_by']
            for i in ego_in:
                G.add_edge(i,k)
                pendants.append(i)
        pendants=array(pendants,dtype=str)
        pendants.flatten()
        pendants=unique(pendants)
        return G, pendants
    except DecodeError:
    except KeyError:
```

#### Rolling the snowball

```
def snowball_round(G, seeds, myspace=False):
    t0=time()
    if myspace:
        seeds=get_myspace_url(seeds)
    sb data=∏
    for s in range(0,len(seeds)):
        s_sg=get_sg(seeds[s])
        new_ego,pen=create_egonet(s_sg)
        for p in pen:
                sb_data.append(p)
        if s<1:
            sb_net=compose(G,new_ego)
        else:
            sb_net=compose(new_ego,sb_net)
        del new_ego
        if s==round(len(seeds)*0.2):
            sb_net.name='20% complete'
            sb net.info()
            print 'AT: '+strftime('%m/%d/%Y, %H:%M:%S', gmtime())
            print ''
    # More time keeping, probably a MUCH better way to do this
    sb data=arrav(sb data)
    sb data.flatten()
    sb data=unique(sb data)
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    return sb net.sb data
```

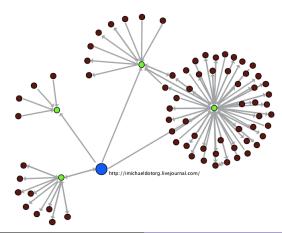
Step	Nodes	Edges	Mean Degree	Density
Seed	5	5	2.0	0.25
k=2	75	115	3.0	0.02
k = 3	4,938	8,659	3.5	3.6(10 <sup>-4</sup> )

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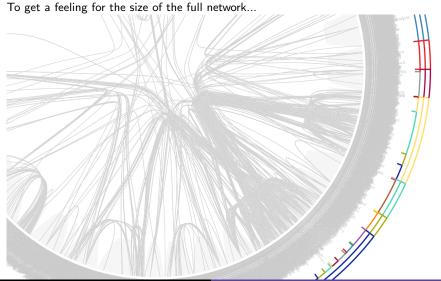
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- Our seed is abnormally isolated, with only four neighbors
- Large jump after first snowball
   Massive structural leap at k = 3
- http://imichaeldotorg.livejournal.com/

### The full network



# Python Dictionaries

The dict type is a data structure that represents a key—value mapping

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Now, try creating a dict of your own

# Using dictionaries for network analysis

#### From the documentation...

#### networkx.closeness\_centrality

closeness\_centrality(G, v=None, weighted\_edges=False)
Compute closeness centrality for nodes.

Closeness centrality at a node is 1/average distance to all other nodes.

Parameters: G : graph

A networkx graph

v : node, optional

Return only the value for node v.

weighted\_edges : bool, optional

Consider the edge weights in determining the shortest paths. If False, all edge weights are considered equal.

Returns: nodes : dictionary

Dictionary of nodes with closeness centrality as the value.

NetworkX's metric's make extensive use of the dict type

▶ In this case the key→value mapping is of the form: {node\_label: metric}

# Let's look at an example:

```
>>> in_cen=in_degree_centrality(hartford)
>>> in_cen
{1: 0.014218009478672987, 2: 0.018957345971563982,...
...
90: 0.0047393364928909956, 293: 0.0}
```

In-degree centrality of Hartford data

We can see that node #90 has in-degree centrality 0.0047

▶ But we can do so much more!

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For our first analysis in NetworkX, we will do some basic network manipulation, then run multiple measures to find highest centrality nodes

 First, we will need to convert to an undirected network, and extract the main component

#### Find main component & symmetrize

- # Many of the centrality metrics require undirected graphs, so we will symmetrize
  >>> hartford ud=hartford.to undirected()
- # The network also has many small components, but for
- # this analysis we are interested in the largest
- >>> hartford\_mc=hartford\_main=connected\_component\_subgraphs(hartford\_ud)[0]

Next, we will calculate multiple measures

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To find the most central actors we will use Python's list comprehension technique to do basic data manipulation on our centrality dictionaries

```
def highest_centrality(cent_dict):
    """Returns node key with largest value from
    NX centrality dict"""
    # Create ordered tuple of centrality data
    cent_items=cent_dict.items()
    # List comprehension!
    cent_items=[(b,a) for (a,b) in cent_items]
    # Sort in descending order
    cent_items.sort()
```

Function for finding most central actor

Now, just ask for the answer

cent\_items.reverse()
return cent items[0][1]

```
>>> print("Actor "+str(highest_centrality(bet_cen))+" has the highest Betweenness centrality")
Actor 82 has the highest Betweenness centrality
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    return cent_items[0][1]
```

# List comprehension

- Given a dict: d={1: 0.15, 2: 0.67}
  d.items() → [(1.0.15),(2.0.67)]
- d.1tems() → [(1,0.15),(2,0.07)
- ▶  $d=[(b,a) \text{ for } (a,b \text{ in } d)] \rightarrow [(0.15,1),(0.67,2)]$

Now, just ask for the answer

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    cent_items=cent_dict.items()
    # List comprehension!
    cent_items=[(b,a) for (a,b) in cent_items]
    # Sort in descending order
    cent_items.sort()
    cent_items.reverse()
    return cent_items[0][1]
```

### List comprehension

- Given a dict: d={1: 0.15, 2: 0.67}
- d.items()  $\rightarrow$  [(1,0.15),(2,0.67)]
- ▶  $d=[(b,a) \text{ for } (a,b \text{ in } d)] \rightarrow [(0.15,1),(0.67,2)]$

Here, we use list comprehension in order to use Python's built-in sort and reverse list functions

Now, just ask for the answer

#### Finding Most central actors

>>> print("Actor "+str(highest\_centrality(bet\_cen))+" has the highest Betweenness centrality")
Actor 82 has the highest Betweenness centrality

To find the most central actors we will use Python's list comprehension technique to do basic data manipulation on our centrality dictionaries

### Function for finding most central actor

```
def highest_centrality(cent_dict):
    """Returns node key with largest value from
    NX centrality dict"""
    # Create ordered tuple of centrality data
    cent_items=cent_dict.items()
    # List comprehension!
    cent_items=[(b,a) for (a,b) in cent_items]
    # Sort in descending order
    cent_items.sort()
    cent_items.reverse()
    return cent_items[0][1]
```

### List comprehension

- Given a dict: d={1: 0.15, 2: 0.67}
- d.items()  $\rightarrow$  [(1,0.15),(2,0.67)]
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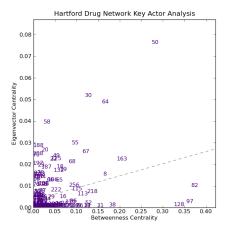
```
>>> print("Actor "+str(highest_centrality(bet_cen))+" has the highest Betweenness centrality")
Actor 82 has the highest Betweenness centrality
```

Recall Python's scientific computing trinity: NumPy, SciPy and matplotlib

► While NumPy and SciPy do most of the behind the scenes work, you will interact with matplotlib frequently for when doing network analysis

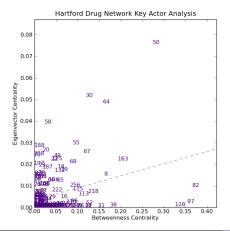
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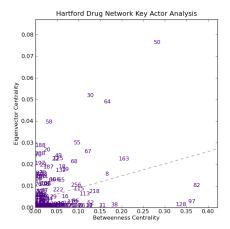
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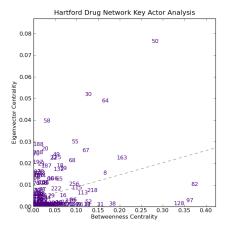


We will need to create a function that takes two centrality dict and generates this plot

1. Create a matplotlib figure

Recall Python's scientific computing trinity: NumPy, SciPy and matplotlib

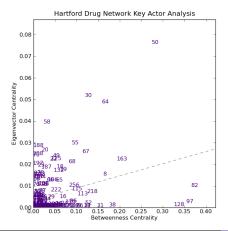
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- 1. Create a matplotlib figure
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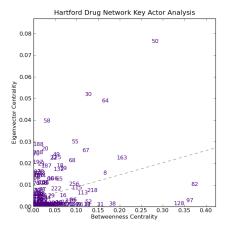
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- 1. Create a matplotlib figure
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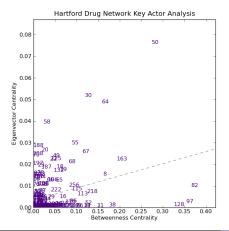
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- 4. Add axis and title labels

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- 1. Create a matplotlib figure
- 2. Plot each node label as a point
- 3. Add a "best fit" line
- 4. Add axis and title labels
- 5. Save figure as a PNG file

```
def centrality_scatter(met_dict1,met_dict2,path="",ylab="",xlab="",title="",reg=False):
    # Create figure and drawing axis
    fig=P.figure(figsize=(7,7))
    ax1=fig.add_subplot(111)
# Create items so actors can be sorted properly
    met_items1=met_dict1.items()
    met_items2=met_dict2.items()
    met_items1.sort()
    met_items2.sort()
# Grab data
    xdata=[(b) for (a,b) in met_items1]
    ydata=[(b) for (a,b) in met_items2]
# Add each actor to the plot by ID
    for p in xrange(len(met_items1)):
        ax1.text(x=xdata[p],y=ydata[p],s=str(met_items1[p][0]),color="indigo")
```

#### The centrality\_scatter function, part one

Create a canvas to draw on

```
def centrality_scatter(met_dict1,met_dict2,path="",ylab="",xlab="",title="",reg=False):
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- Create a canvas to draw on
- manipulate and store centrality data

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def centrality_scatter(met_dict1,met_dict2,path="",ylab="",xlab="",rittle="",reg=False):
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```

- Create a canvas to draw on
- manipulate and store centrality data
- ▶ Add points to plot as node labels

```
def centrality_scatter(met_dict1,met_dict2,path="",ylab="",xlab="",title="",reg=False):
    # If adding a best fit line, we will use NumPv to calculate the points.
    if reg:
        # Function returns y-intercept and slope. So, we create a function to
        # draw LOBF from this data
        slope, vint=polyfit(xdata, ydata, 1)
        xline=P.xticks()[0]
        vline=map(lambda x: slope*x+vint.xline)
        # Add line
        ax1.plot(xline,yline,ls='--',color='grey')
    # Set new x- and y-axis limits to data
    P.xlim((0.0,max(xdata)+(.15*max(xdata)))) # Give a little buffer
    P.ylim((0.0,max(ydata)+(.15*max(ydata))))
    # Add labels
    ax1.set_title(title)
    ax1.set_xlabel(xlab)
    ax1.set_vlabel(vlab)
    # Save figure
    P.savefig(path,dpi=100)
```

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Add a best fit line

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- Add a best fit line
- ▶ Resize figure to fit data

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    # Add labels
    ax1.set title(title)
    ax1.set_xlabel(xlab)
    ax1.set_vlabel(vlab)
    # Save figure
    P.savefig(path,dpi=100)
```

- Add a best fit line
- Resize figure to fit data
- Add labels, and save the figure as a PNG file

# Exporting network data and analytics

As powerful as NetworkX and the complementing scientific computing packages in Python are, it may often be useful or necessary to output your data for additional analysis

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Next, we will review how to save data in different formats and export metrics to a CSV file using the Hartford drug net data

```
NX syntax for loading a file

>>> write_format(G, "path/to/file.txt", ....options...)

↑ ↑ ↑

NX function, net variable File to be written Nodes/edge data, etc.
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The syntax for exporting network data follows exactly the syntax for loading it

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```

#### Let's try!

- Output the Hartford drug net data as an adjacency list
- Add metric data to each node of the network
- Output new network in Pajek format with node attributes

### Saving network data and adding node attributes

As shown, this is a simple one line operation

### Output Hartford drug net data as an adjacency list

```
write_adjlist(hartford_mc,"../../data/hartford_mc_adj.txt")
```

Next, we will add the Eigenvector centrality of each node to the graph object

#### Adding node attributes

```
def add_metric(G,met_dict):
    """Adds metric data to G from a dictionary keyed by node labels"""
    if(G.nodes().sort()==met_dict.keys().sort()):
        for i in met_dict.keys():
            G.add_node(i,metric=met_dict[i])
        return G
    else:
        raise ValueError("Node labels do not match")
```

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Quick error checking

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```

- Quick error checking
- Add node attribute as "metric"

Python has powerful built-in tools for reading and writing standard data formats

 One of the most useful, and frequently used, is the CSV library and the DictWriter

```
import csv
def csv_exporter(data_dict,path):
    """Takes a dict of centralities keved by column headers and exports
    data as a CSV file"""
    # Create column header list
    col headers=["Actor"]
    col headers.extend(data dict.kevs())
    # Create CSV writer and write column headers
    writer=csv.DictWriter(open(path."w").fieldnames=col headers)
    writer.writerow(dict((h,h) for h in col headers))
    # Write each row of data
    for j in data_dict[col_headers[1]].keys():
        # Create a new dict for each row
        row=dict.fromkeys(col_headers)
        row["Actor"]=j
        for k in data dict.kevs():
            row[k]=data dict[k][i]
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```

# The results of CSV export

We can now open the CSV file in our favorite spreadsheet program

- Perform traditional data exploration
- ► Load into other analytics platforms for additional analysis (e.g., R)
- Store for latter use

A B C  1 Actor Closeness Betweeness Eig	D
1 Actor Closeness Betweeness Eig	renvector
	genvector
2 1 0.12467532 0.0072576 0	0.00025176
3 2 0.12475634 0.01767427 0	0.00025964
	0.00023185
5 4 0.10223642 0.03108639	1.44E-05
6 5 0.1443609 0 0	0.00313152
7 6 0.09943035 0.01041667	1.49E-07
8 7 0.11340815 0.04362093	6.78E-05
9 8 0.20512821 0.16354003 0	0.01471888
10 9 0.11267606 0.00741624	0.0001101
	0.00095456
12 11 0.1703638 0.01250999	0.0032333
13 13 0.13892909 0	1.79E-05
14 14 0.17219731 0.11848775 0	0.00029737
15 15 0.13521127 0.00079897	2.11E-05
16 0.15907208 0.06203647 0	0.00432838