

Social Network Analysis Workshop

CIShell Powered Tools: Network Workbench (NWB) & Science of Science (Sci2) Tool

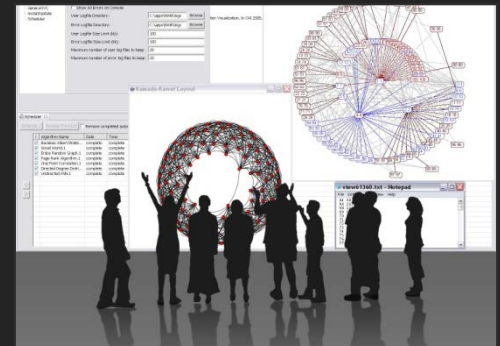
Dr. Katy Börner and Chin Hua Kong

Cyberinfrastructure for Network Science Center
Information Visualization Laboratory
School of Library and Information Science
Indiana University, Bloomington, IN
<http://cns.iu.edu>

With special thanks to Kevin W. Boyack, Micah Linnemeier,
Russell J. Duhon, Patrick Phillips, Joseph Biberstine, Chintan Tank
Nianli Ma, Scott Weingart, Hanning Guo, Mark A. Price, Angela M.
Zoss, Ted Polley, and Sean Lind

*Second Annual International Science of Team Science Conference
Chicago, IL*

Thursday, April 14, 2011 • 1:15 – 5:00 PM





Online Resources

- These slides
<http://sci2.cns.iu.edu/docs/2011-borner-SciTS-workshop.pdf>
- Sci2 Tool Manual v0.5 Alpha
<http://sci2.wiki.cns.iu.edu>
- Sci2 Tool v0.5 Alpha (April 4, 2011)
<http://sci2.cns.iu.edu>

- Additional Datasets
<http://sci2.wiki.cns.iu.edu/2.5+Sample+Datasets>
- Additional Plugins
<http://sci2.wiki.cns.iu.edu/3.2+Additional+Plugins>

Or copy them from the DVD or memory stick.



Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

- Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

- Load and clean a dataset as text file; process raw data into networks.
- Find basic statistics and run various algorithms over the network.
- Visualize as either a circular hierarchy or network

3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

4:45 Outlook and Discussion

5:00 Adjourn



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Macroscopes Serve the Changing Scientific Landscape



IS



CS



Bio



SNA



Physics

Different datasets/formats.
Diverse algorithms/tools written in
many programming languages.



The Changing Scientific Landscape

Star Scientist -> Research Teams: In former times, science was driven by key scientists. Today, science is driven by effectively collaborating co-author teams often comprising expertise from multiple disciplines and several geospatial locations (Börner, Dall'Asta, Ke, & Vespignani, 2005; Shneiderman, 2008).

Users -> Contributors: Web 2.0 technologies empower anybody to contribute to Wikipedia or to exchange information. WikiProfessionals, or WikiProteins combine wiki and semantic technology in support of real time community and research (Mons et al., 2008).

Cross-disciplinary: The best tools frequently borrow and synergistically combine methods and techniques from different disciplines of science and empower interdisciplinary and/or cross-disciplinary research to fine-tune and interpret data.

One Specimen -> Data Streams: Microscopes and telescopes were originally used to study one specimen at a time. Today, many researchers must make sense of massive streams of multiple types of data with different formats, dynamics, and origin.

Static Instrument -> Evolving Cyberinfrastructure (CI): The importance of hardware instruments that are rather static and expensive decreases relative to software infrastructures that are highly flexible and continuously evolving according to the needs of different sciences. Some of the most successful services and tools are decentralized increasing scalability and fault tolerance.

Cyberinfrastructure and Datasets for SciTS Research

Wednesday, April 13 • 12:00 PM – 1:15 PM



Macroscopic Design

Custom Tools for Different Scientific Communities

Information Visualization Cyberinfrastructure

<http://iv.cns.iu.edu>

Network Workbench Tool + Community Wiki

<http://nwb.cns.iu.edu>

Science of Science (Sci²) Tool and Portal

<http://sci2.cns.iu.edu>

Epidemics Cyberinfrastructure

Coming soon



180+ Algorithm Plugins and Branded GUIs

+

Core Architecture

Open Services Gateway Initiative (OSGi) Framework.

<http://orgi.org>

Cyberinfrastructure Shell (CIShell)

<http://cishell.org>



CIShell Powered Tools: Network Workbench (NWB)



Network Workbench Tool

<http://nwb.slis.indiana.edu>

The Network Workbench (NWB) tool supports researchers, educators, and practitioners interested in the study of biomedical, social and behavioral science, physics, and other networks.

In February 2009, the tool provides more 169 plugins that support the preprocessing, analysis, modeling, and visualization of networks.

More than 50 of these plugins can be applied or were specifically designed for S&T studies.

It has been downloaded more than 65,000 times since December 2006.



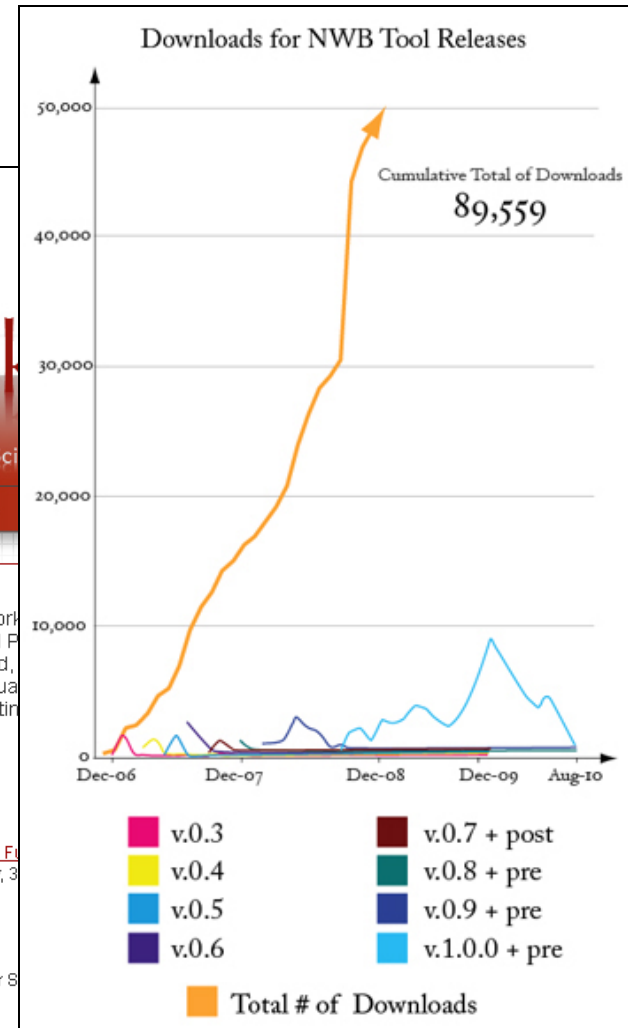
Summary

Network Workbench: A Large-Scale Network Toolkit for Biomedical, Social Science and P evaluate, and operate a unique distributed, scale network analysis, modeling, and visual (NWB). The envisioned data-code-computin

[more](#)
[How to cite this project](#)

News & Updates

- 5.1.09 Kaelble, Steve. 2009. [Mapping the Fu Knowledge](#). *Research & Creative Activity*, 3 ([website](#) accessed 5/1/09)
- 3.23.09 [1.0.0 beta 5](#) Released
- 1.23.09 Ann Mcranie's [tutorial abstract](#) for S 2009
- 11.4.08 Two NWB Pls featured in "[Connected—The Power of Six Degrees](#)." 2008. Anna Maria Talas, Director. Australian Broadcasting Corporation, Ltd. [\[YouTube\]](#) [\[Full Video\]](#) (300MB)



[Getting Started](#)
See more [documentation](#)

[Get Involved](#)

Herr II, Bruce W., Huang, Weixia (Bonnie), Penumarthy, Shashikant & Börner, Katy. (2007). *Designing Highly Flexible and Usable Cyberinfrastructures for Convergence*. In Bainbridge, William S. & Roco, Mihail C. (Eds.), *Progress in Convergence - Technologies for Human Wellbeing* (Vol. 1093, pp. 161-179), *Annals of the New York Academy of Sciences*, Boston, MA.

Investigators: Katy Börner, Albert-Laszlo Barabasi, Santiago Schnell, Alessandro Vespignani & Stanley Wasserman, Eric Wernert



Software Team: Lead: Micah Linnemeier
Members: Patrick Phillips, Russell Duhon, Tim Kelley & Ann McCranie
Previous Developers: Weixia (Bonnie) Huang, Bruce Herr, Heng Zhang, Duygu Balcan, Bryan Hook, Ben Markines, Santo Fortunato, Felix Terkhorn, Ramya Sabbineni, Vivek S. Thakre & Cesar Hidalgo



Goal: Develop a large-scale network analysis, modeling and visualization toolkit for physics, biomedical, and social science research.

Amount: \$1,120,926, NSF IIS-0513650 award

Duration: Sept. 2005 - Aug. 2009

Website: <http://nwb.slis.indiana.edu>

NWB Advisory Board:

James Hendler (Semantic Web) <http://www.cs.umd.edu/~hendler/>

Jason Leigh (CI) <http://www.evl.uic.edu/spiff/>

Neo Martinez (Biology) <http://online.sfsu.edu/~webhead/>

Michael Macy, Cornell University (Sociology) <http://www.soc.cornell.edu/faculty/macy.shtml>

Ulrik Brandes (Graph Theory) <http://www.inf.uni-konstanz.de/~brandes/>

Mark Gerstein, Yale University (Bioinformatics) <http://bioinfo.mbb.yale.edu/>

Stephen North (AT&T) <http://public.research.att.com/viewPage.cfm?PageID=81>

Tom Snijders, University of Groningen <http://stat.gamma.rug.nl/snijders/>

Noshir Contractor, Northwestern University <http://www.spcomm.uiuc.edu/nosh/>



Computational Proteomics

What relationships exist between protein targets of all drugs and all disease-gene products in the human protein–protein interaction network?

*Yildirim, Muhammed
A., Kwan-II Goh,
Michael E. Cusick,
Albert-László Barabási,
and Marc Vidal. (2007).
Drug-target Network.
Nature Biotechnology
25 no. 10: 1119-1126.*

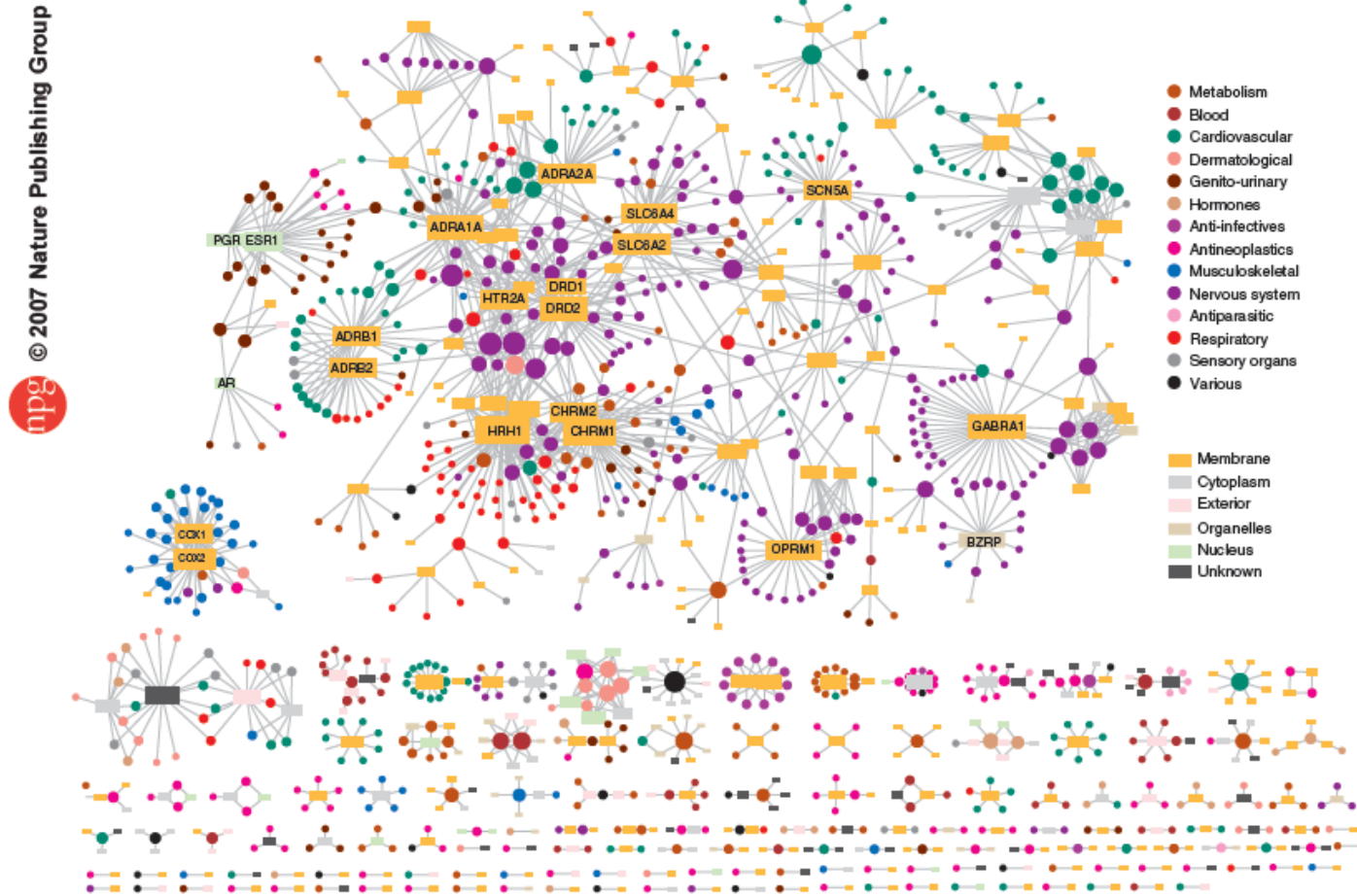


Figure 2 Drug-target network (DT network). The DT network is generated by using the known associations between FDA-approved drugs and their target proteins. Circles and rectangles correspond to drugs and target proteins, respectively. A link is placed between a drug node and a target node if the protein is a known target of that drug. The area of the drug (protein) node is proportional to the number of targets that the drug has (the number of drugs targeting the protein). Color codes are given in the legend. Drug nodes (circles) are colored according to their Anatomical Therapeutic Chemical Classification, and the target proteins (rectangular boxes) are colored according to their cellular component obtained from the Gene Ontology database.



Computational Economics

Does the type of product that a country exports matter for subsequent economic performance?

C. A. Hidalgo, B. Klinger,
A.-L. Barabási, R. Hausmann
(2007) *The Product Space
Conditions the Development
of Nations. Science* 317,
482 (2007).

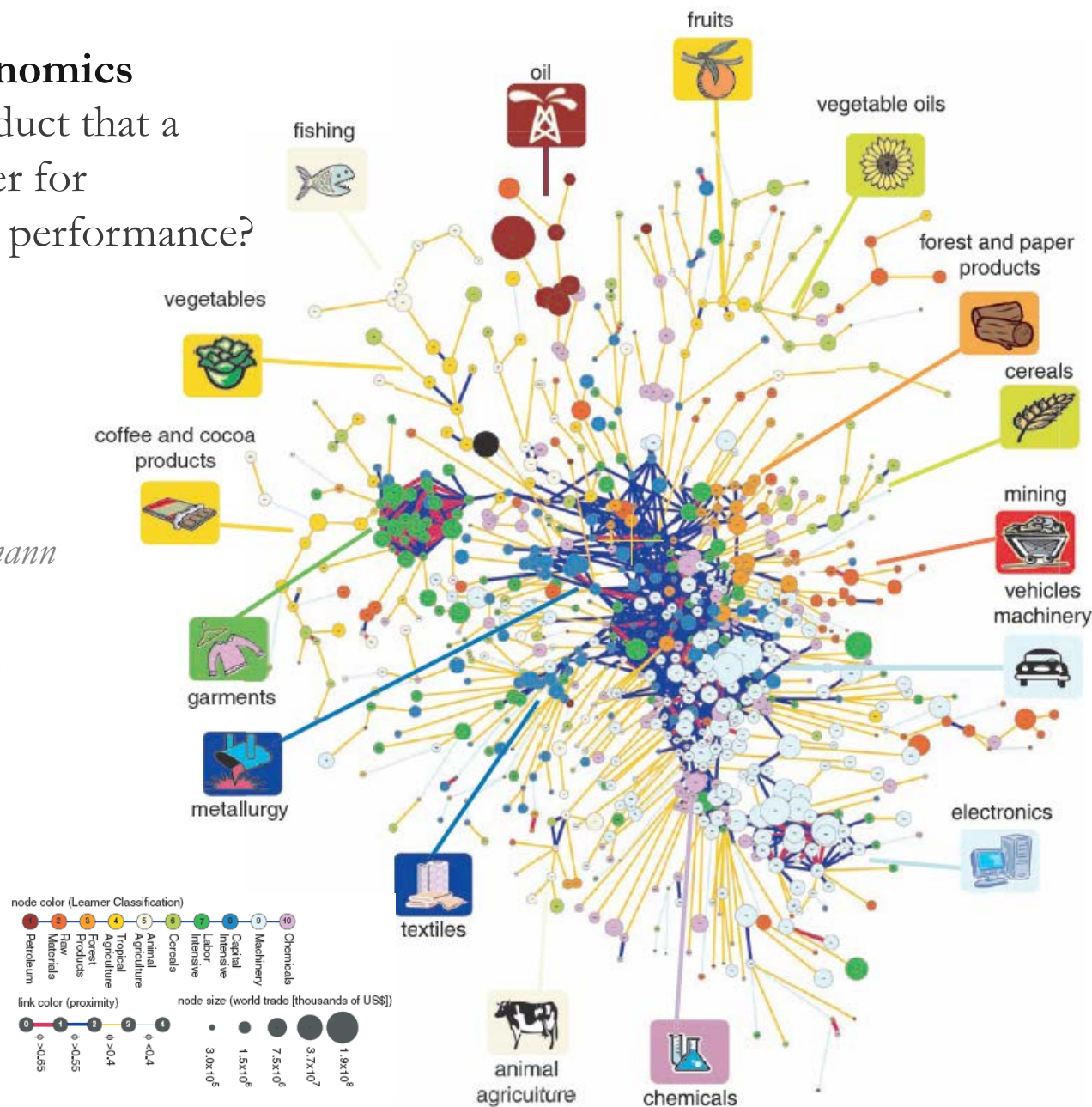


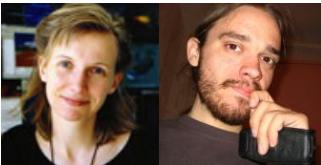
Fig. 1. The product space. (A) Hierarchically clustered proximity (ϕ) matrix representing the 775 SITC-4 product classes exported in the 1998–2000 period. (B) Network representation of the product space. Links are color coded

with their proximity value. The sizes of the nodes are proportional to world trade, and their colors are chosen according to the classification introduced by Leamer.

Computational Social Science

Studying large scale social networks such as Wikipedia

*Second Sight: An Emergent Mosaic of Wikipedian Activity,
The NewScientist, May 19, 2007*

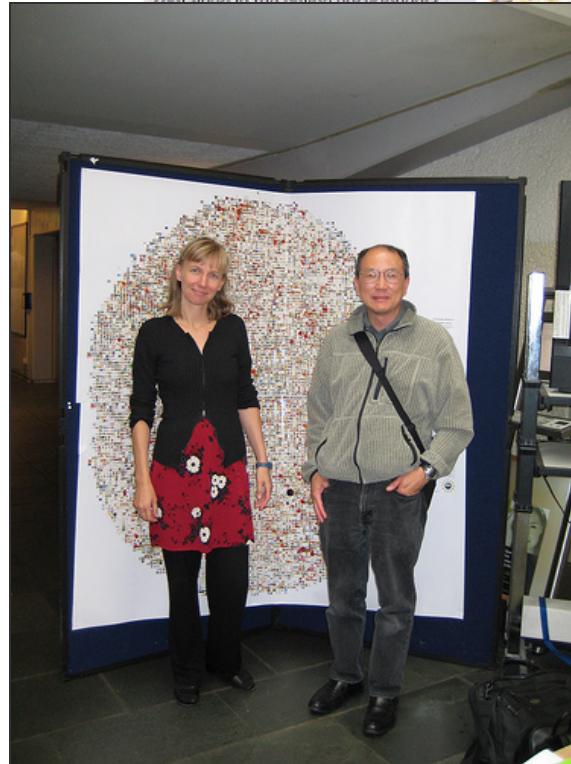


Second sight

Image: Bruce W. Herr and Todd M. Holloway

Power struggle

How do you keep track of the bubbling mass of information that is Wikipedia? This chaotic-looking mosaic is one attempt to show which topics are contained in the online encyclopedia.



locked until the mood tools (locked pages at the time of writing include entries on Sheffield Wednesday football club, Mikhail Gorbachev and pigs).

The mosaic has been commended in a competition for images that visualise network dynamics, coinciding with this week's International Workshop and Conference on Network Science in Bloomington.



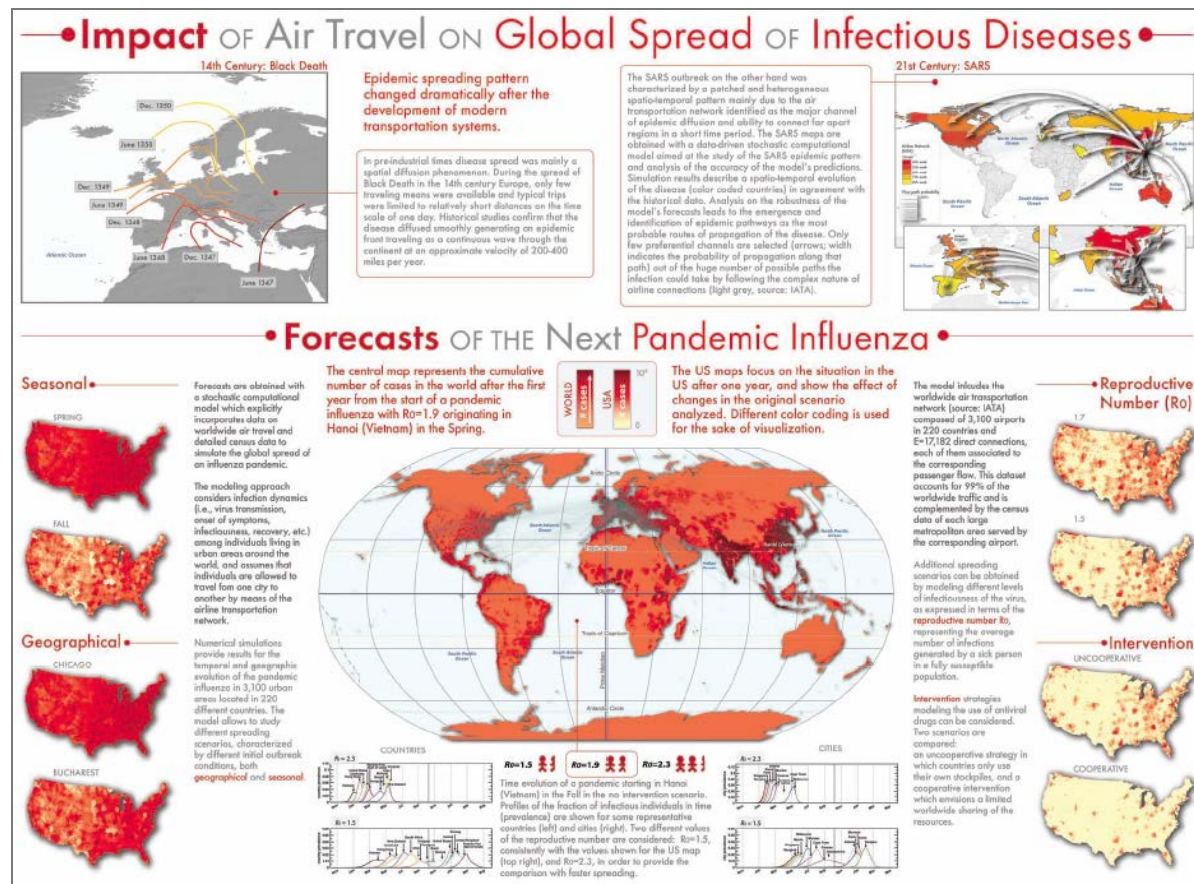
Computational Epidemics

Forecasting (and preventing the effects of) the next pandemic.

Epidemic Modeling in Complex realities, V. Colizza, A. Barrat, M. Barthelemy, A. Vespignani, *Comptes Rendus Biologie*, 330, 364-374 (2007).

Reaction-diffusion processes and metapopulation models in heterogeneous networks, V. Colizza, R. Pastor-Satorras, A. Vespignani, *Nature Physics* 3, 276-282 (2007).

Modeling the Worldwide Spread of Pandemic Influenza: Baseline Case and Containment Interventions, V. Colizza, A. Barrat, M. Barthelemy, A.-J. Valleron, A. Vespignani, *PLoS-Medicine* 4, e13, 95-110 (2007).



NWB Tool Download, Install, and Run

NWB Tool 1.0.0

Can be freely downloaded for all major operating systems from <http://nwb.cns.iu.edu>

Select your operating system from the pull down menu and download.

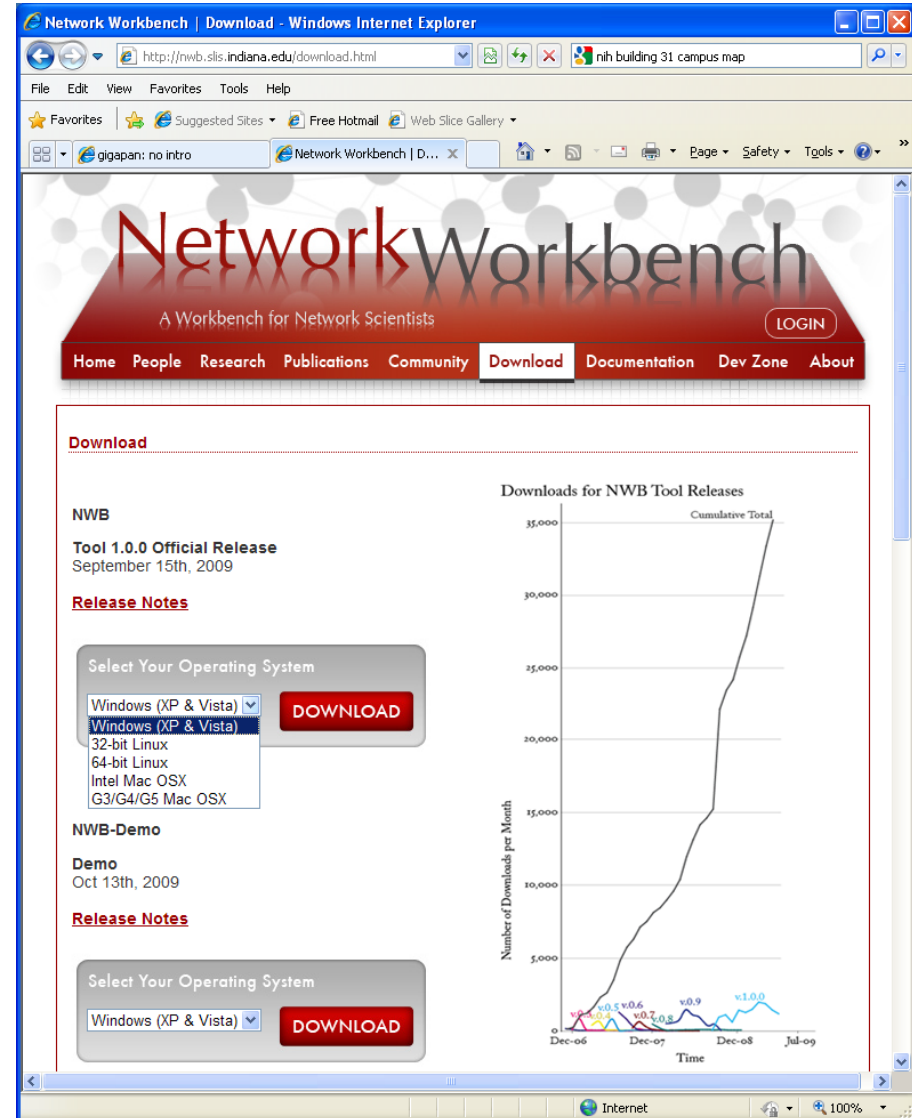
Unpack into a /nwb directory.

Run /nwb/nwb.exe

Session log files are stored in *'*yournwbdirectory*/logs'* directory.

Cite as

NWB Team. (2006). Network Workbench Tool. Indiana University, Northeastern University, and University of Michigan, <http://nwb.cns.iu.edu>.



NWB Tool Interface Components

Network Workbench Tool

File Preprocessing Modeling Analysis Visualization Scientometrics Help

Console

Console displays data operations (save, load, view, etc.) and algorithm input parameters, selection, & acknowledgements as well as error reporting.

Welcome to the Network Workbench Tool. This tool supports the preprocessing, modeling, analysis, and visualization of small, network datasets.

The Network Workbench Tool is supported in part by the NSF IIS-0513650 award. The primary investigators are Dr. Katy Börner, Dr. Albert-László Barabási, Dr. Santiago Schnell, Dr. Alessandro Vespignani, Dr. Stanley Wasserman, and Dr. Eric A. Wernert.

The NWB tool was developed by Weixia Huang, Russell Duhon, Micah Linnemeier, Timothy Kelley, Duygu Balcan, Mariano Beiró, Bruce Herr, Santo Fortunato, Ben Markines, Felix Terkhorn, Heng Zhang, Megha Ramawat, César Hidalgo, Ramya Sabbineni, Vivek Thakres, Soma Sanyal, Ann McCranie, Alessandro Vespignani, and Katy Börner. It uses the Cyberinfrastructure Shell (<http://cishell.org>) developed at the Cyberinfrastructure for Network Science Center (<http://cns.slis.indiana.edu>) at Indiana University.

Please cite as follows:
NWB Team. (2006). Network Workbench Tool. Indiana University and Northeastern University, <http://nwb.slis.indiana.edu>

Scheduler

Scheduler lists what algorithms you've used and displays algorithm progress.

Remove From List ☐ Remove completed automatically ☐ Remove all completed

!	Algorithm Name	Date	Time	% Complete

Data Manager

Data Manager keeps track of all datasets that are available for algorithmic visualization or manipulation.

- Table
- Matrix
- Plot
- Text
- GUESS
- Tree
- Network-

Console shows references to seminal works.
Workflows are recorded into a log file, and soon can be re-run for easy replication.
All algorithms are documented online; workflows are given in tutorials.

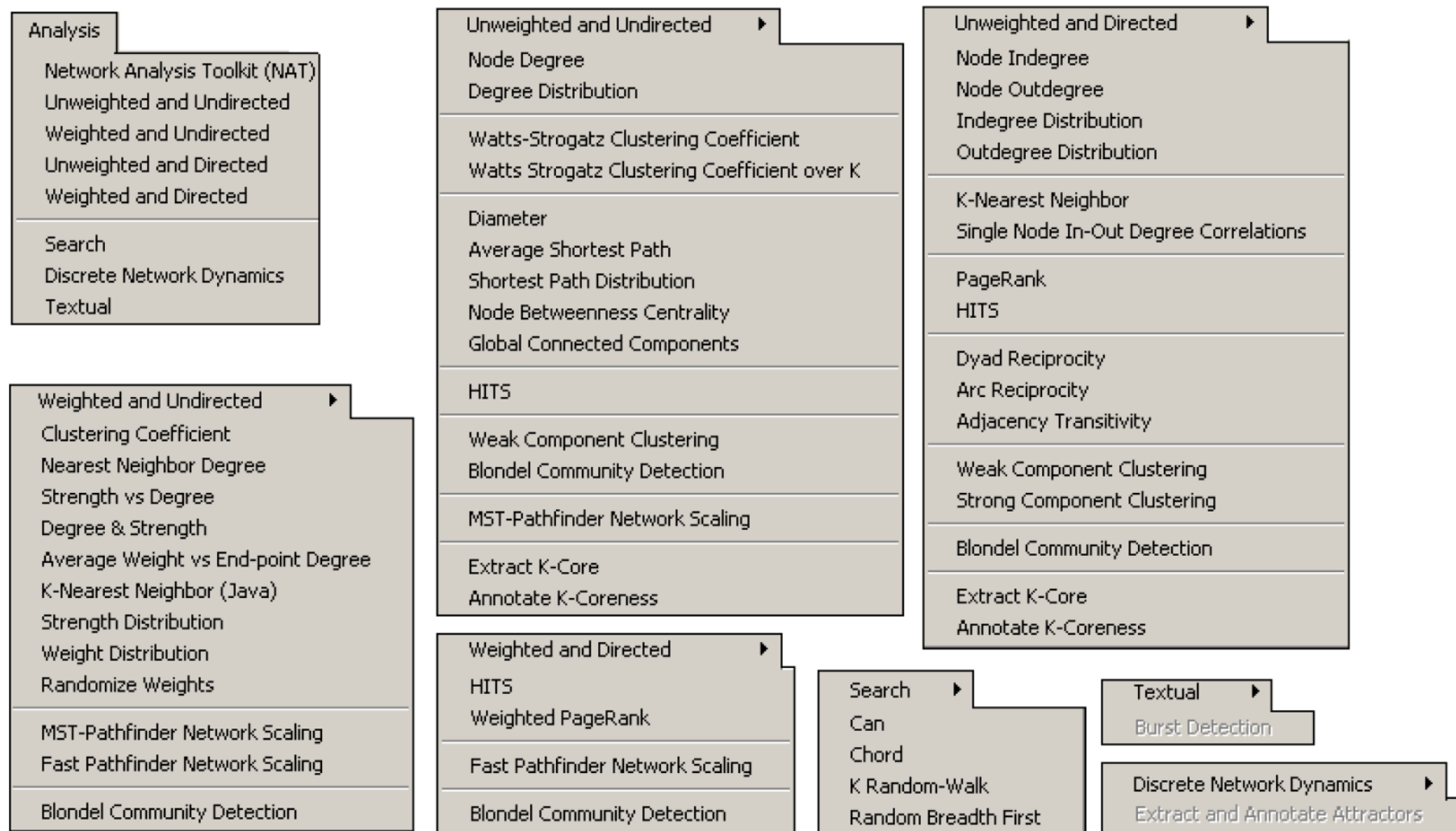
File, Preprocessing, Modeling, and Visualization Menus

File	Preprocessing	Modeling	Visualization
Load...	Extract Top Nodes	Random Graph	GUESS
Load and Clean ISI File	Extract Nodes Above or Below Value	Watts-Strogatz Small World	GnuPlot
Read Directory Hierarchy Datasets	Remove Node Attributes	Barabási-Albert Scale-Free	DrL (VxOrd)
Save...	Delete High Degree Nodes	Can	Specified (prefuse beta)
View...	Delete Random Nodes	Chord	Circular (JUNG)
View with...	Delete Isolates	Hypergrid	Radial Tree/Graph (prefuse alpha)
Merge Node and Edge Files	Extract Top Edges	PRU	Radial Tree/Graph with Annotation (prefuse beta)
Split Graph to Node and Edge Files	Extract Edges Above or Below Value	TARL	Tree Map (prefuse beta)
Tests	Remove Edge Attributes	Discrete Network Dynamics (DND)	Tree View (prefuse beta)
Preferences	Remove Self Loops	Evolving Network (Weighted)	Balloon Graph (prefuse alpha)
Exit	Trim by Degree		Force Directed with Annotation (prefuse beta)
	Snowball Sampling (n nodes)		Kamada-Kawai (JUNG)
	Node Sampling		Fruchterman-Reingold (JUNG)
	Edge Sampling		Fruchterman-Reingold with Annotation (prefuse beta)
	Symmetrize		Spring (JUNG)
	Dichotomize		Small World (prefuse alpha)
	Multipartite Joining		Parallel Coordinates (demo)
	Normalize Text		LaNet
	Slice Table by Time		Circular Hierarchy

Börner, Katy, Sanyal, Soma and Vespignani, Alessandro (2007). **Network Science**. In Blaise Cronin (Ed.), *ARIST*, Information Today, Inc./American Society for Information Science and Technology, Medford, NJ, Volume 41, Chapter 12, pp. 537-607.

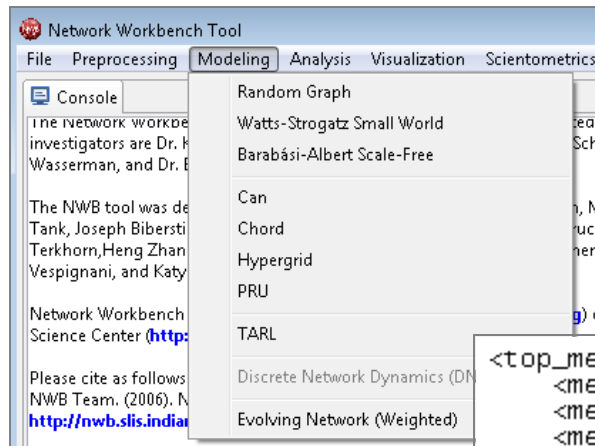
<http://ivl.slis.indiana.edu/km/pub/2007-borner-arist.pdf>

Analysis Menu and Submenus

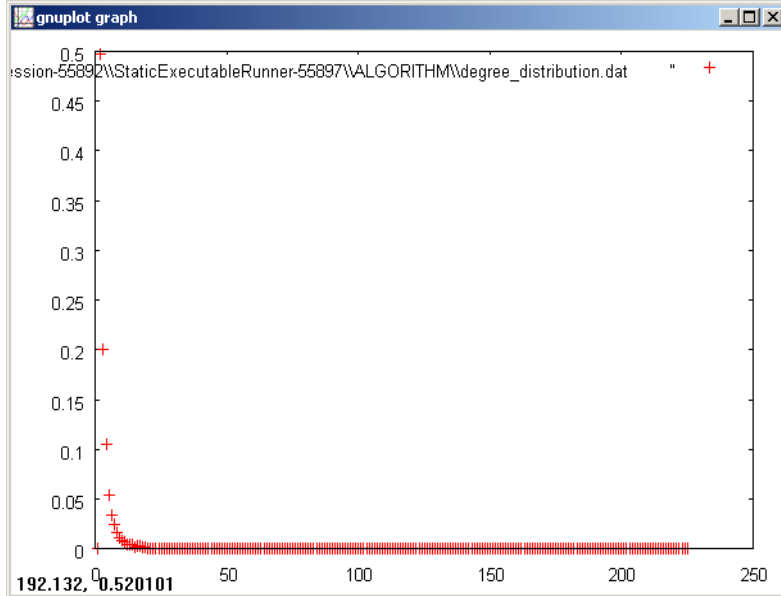


Börner, Katy, Sanyal, Soma and Vespignani, Alessandro (2007). **Network Science**. In Blaise Cronin (Ed.), *ARIST*, Information Today, Inc./American Society for Information Science and Technology, Medford, NJ, Volume 41, Chapter 12, pp. 537-607. <http://ivl.slis.indiana.edu/km/pub/2007-borner-arist.pdf>

- The file *yourtooldirectory/configuration/default_menu.xml* encodes the structure of the menu system.
- In NWB Tool, the Modeling menu (left) is encoded by the following piece of xml code:

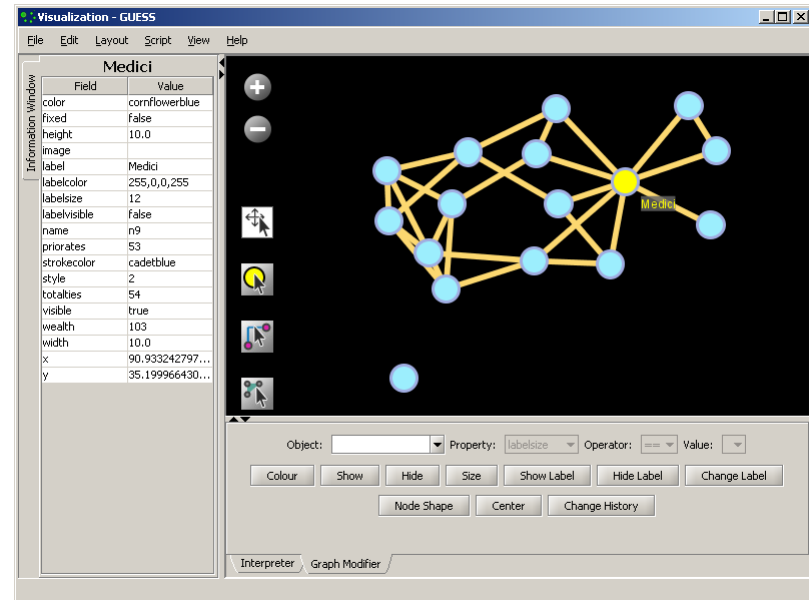


```
<top_menu name="Modeling">
  <menu pid="edu.iu.nwb.modeling.erdosrandomgraph"/>
  <menu pid="edu.iu.nwb.modeling.smallworld"/>
  <menu pid="edu.iu.nwb.modeling.barabasiAlbert"/>
  <menu type="break"/>
  <menu pid="edu.iu.iv.modeling.p2p.can.CanAlgorithm"/>
  <menu pid="edu.iu.iv.modeling.p2p.chord.ChordAlgorithm"/>
  <menu pid="edu.id.iv.modeling.p2p.hypergrid.Hypergrid"/>
  <menu pid="edu.iu.iv.modeling.p2p.pru.PruAlgorithm"/>
  <menu type="break"/>
  <menu pid="edu.iu.iv.modeling.tarl.TarlAlgorithm"/>
  <menu type="break"/>
  <menu pid="edu.iu.nwb.modeling.discretenetworkdynamics.DNAlgorithm"/>
  <menu type="break"/>
  <menu pid="edu.iu.nwb.modeling.weighted.evolvingnetwork"/>
</top_menu>
```



Gnuplot

portable command-line driven
interactive data and function plotting
utility <http://www.gnuplot.info/>.



GUESS

exploratory data analysis and visualization tool
for graphs and networks.

<https://nwb.slis.indiana.edu/community/?n=VisualizeData.GUESS>.

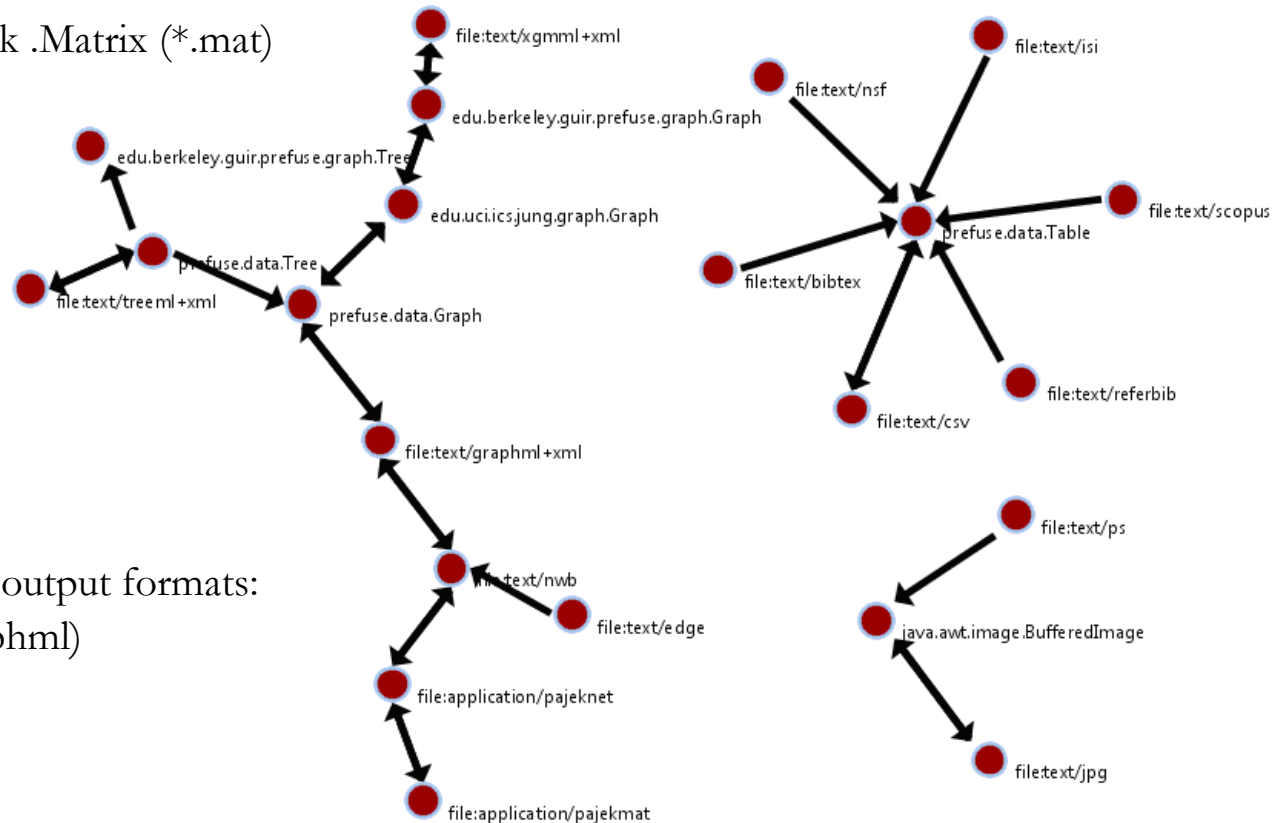
Supported Data Formats

The NWB tool supports loading the following input file formats:

- GraphML (*.xml or *.graphml)
- XGMML (*.xml)
- Pajek .NET (*.net) & Pajek .Matrix (*.mat)
- NWB (*.nwb)
- TreeML (*.xml)
- Edge list (*.edge)
- CSV (*.csv)
- ISI (*.isi)
- Scopus (*.scopus)
- NSF (*.nsf)
- Bibtex (*.bib)
- Endnote (*.enw)

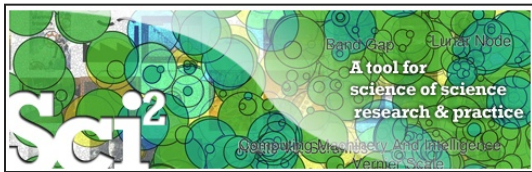
and the following network file output formats:

- GraphML (*.xml or *.graphml)
- Pajek .MAT (*.mat)
- Pajek .NET (*.net)
- NWB (*.nwb)
- XGMML (*.xml)
- CSV (*.csv)



Formats are documented at <https://nwb.slis.indiana.edu/community/?n=DataFormats.HomePage>.

CIShell Powered Tools:
Science of Science (Sci2) Tool



Science of Science (Sci2) Tool

<http://sci2.cns.iu.edu>

- Explicitly designed for SoS research and practice, well documented, easy to use.
- Empowers many to run common studies while making it easy for exports to perform novel research.
- Advanced algorithms, effective visualizations, and many (standard) workflows.
- Supports micro-level documentation and replication of studies.
- Is open source—anybody can review and extend the code, or use it for commercial purposes.

nature

OPINION

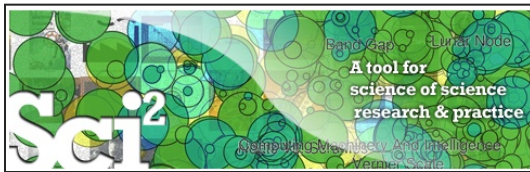
SUMMARY

- Existing metrics have known flaws
- A reliable, open, joined-up data infrastructure is needed
- Data should be collected on the full range of scientists' work
- Social scientists and economists should be involved

Vol 464|25 March 2010

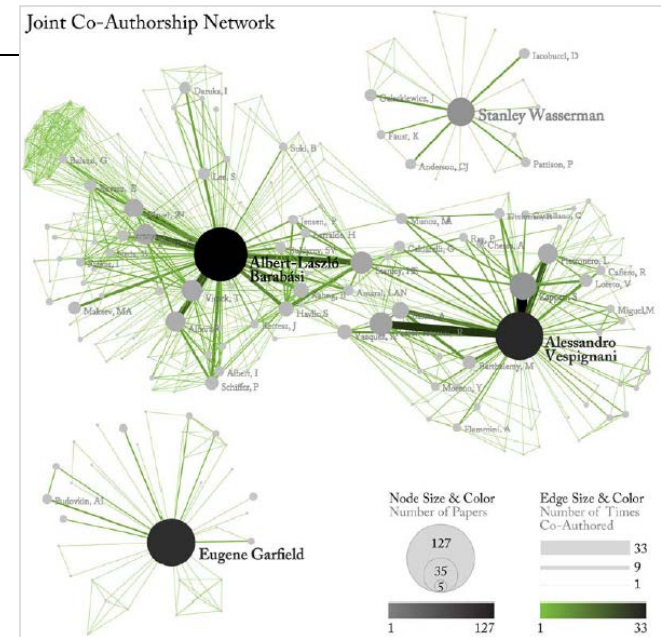
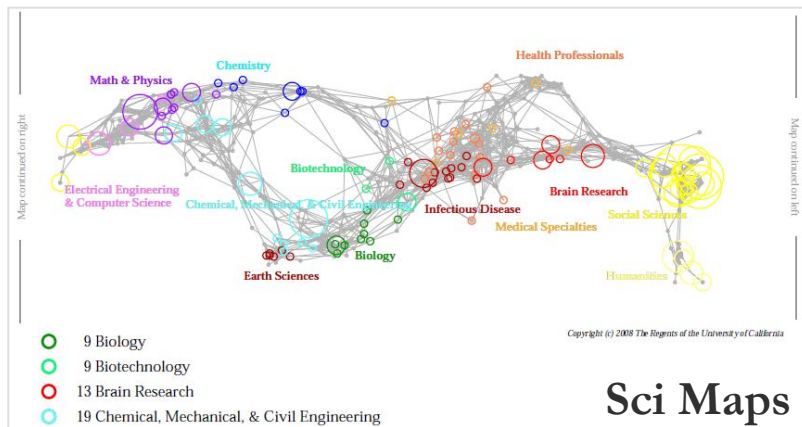
Let's make science metrics more scientific

To capture the essence of good science, stakeholders must combine forces to create an open, sound and consistent system for measuring all the activities that make up academic productivity, says **Julia Lane**.

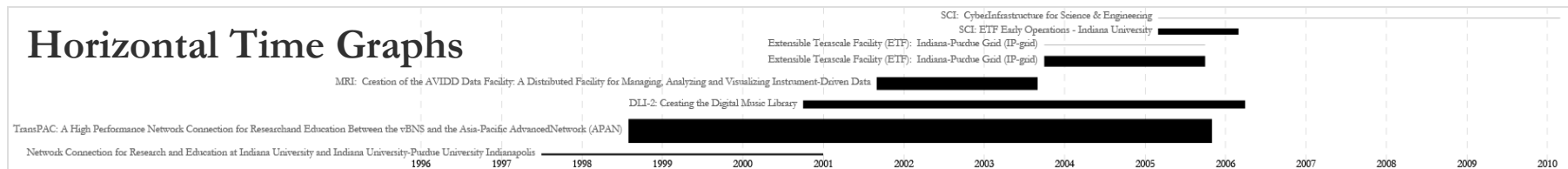


Sci² Tool – “Open Code for S&T Assessment”

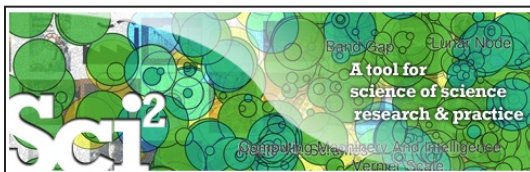
OSGi/CIShell powered tool with NWB plugins and many new scientometrics and visualizations plugins.



Horizontal Time Graphs



Börner, Katy, Huang, Weixia (Bonnie), Linnemeier, Micah, Dubon, Russell Jackson, Phillips, Patrick, Ma, Nianli, Zoss, Angela, Guo, Hanning & Price, Mark. (2009). *Rete-Netzwerk-Red: Analyzing and Visualizing Scholarly Networks Using the Scholarly Database and the Network Workbench Tool*. *Proceedings of ISSI 2009: 12th International Conference on Scientometrics and Informetrics, Rio de Janeiro, Brazil, July 14-17 . Vol. 2, pp. 619-630.*



Sci² Tool

Sci² Tool

File Preprocessing Modeling Analysis Visualization Scientometrics Help

Console

Welcome to the Science of Science Tool (Sci²). The development of this tool is supported in part by the National Science Foundation (NSF) Grant IRI-0715303, and the James S. McDonnell-Petersen Cyberinfrastructure portal (<http://sci.slis.indiana.edu>).

The primary investigators are Katy Börner, Ingrid Isenhardt, SciTech Strategies Inc. The Sci² tool was developed by J. Duhon, Patrick A. Phillips, Chintan Tank, a Cyberinfrastructure Shell (<http://cishell.org>) for Network Science Center (<http://cns.slis.indiana.edu>). Many algorithm plugins were derived from the Network Science Center (<http://nwb.slis.indiana.edu>).

Please cite as follows:
Sci² Team. (2009). Science of Science Tool. Ingrid Isenhardt, SciTech Strategies Inc., <http://sci.slis.indiana.edu>.

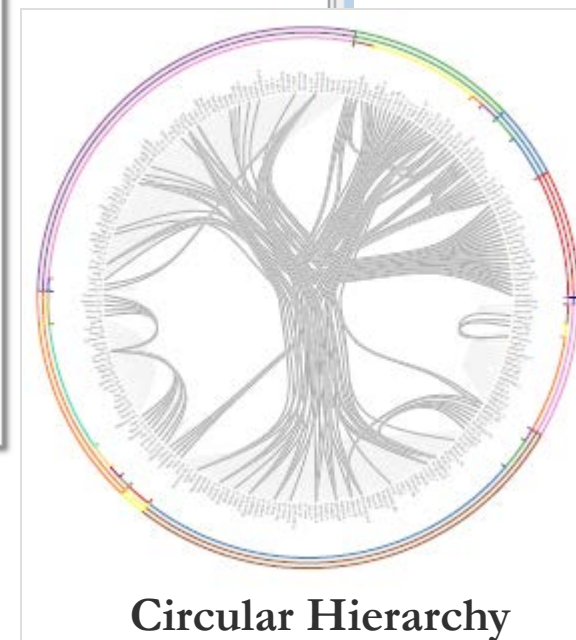
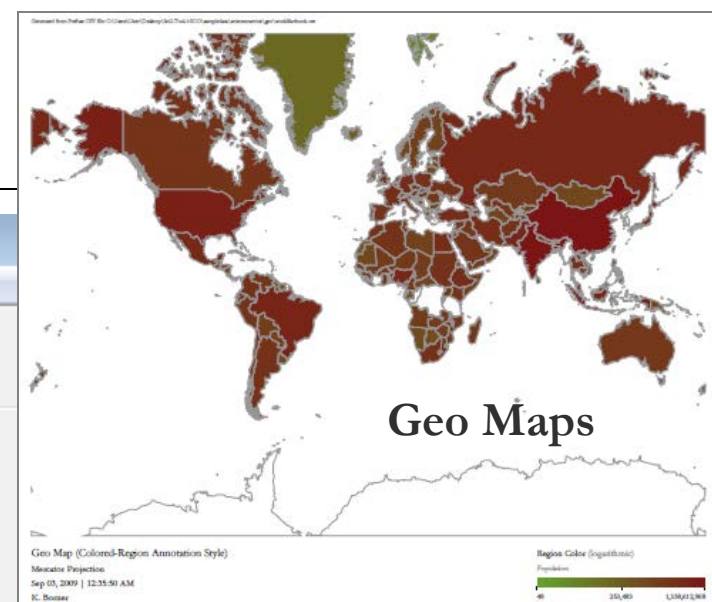
Scheduler

Remove From List ☐ Remove completed

!	Algorithm Name	Date	Time	% Complete
<input checked="" type="checkbox"/>	Extract Co-Author Network	09/03/2009	00:15:20 AM	100%
<input checked="" type="checkbox"/>	Load and Clean ISI File	09/03/2009	00:15:05 AM	100%

Visualization Menu:

- GUESS
- GnuPlot
- Radial Tree/Graph (prefuse alpha)
- Radial Tree/Graph with Annotation (prefuse beta)
- Tree View (prefuse beta)
- Tree Map (prefuse beta)
- Force Directed with Annotation (prefuse beta)
- Fruchterman-Reingold with Annotation (prefuse beta)
- DrL (VxOrd)
- Specified (prefuse beta)
- Horizontal Line Graph
- Circular Hierarchy
- Geo Map (circle annotations)
- Geo Map (region coloring annotations)
- Image Viewer
- RefMapper





Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

➤ **Download and run the tool.**

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

- Load and clean a dataset as text file; process raw data into networks.
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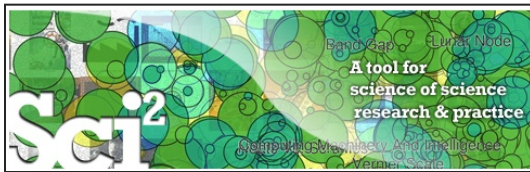
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

4:45 Outlook and Discussion

5:00 Adjourn



Sci² Tool: Download, Install, and Run

Sci² Tool v0.5 Alpha (April 4, 2011)

Can be freely downloaded for all major operating systems from

<http://sci2.cns.iu.edu>

Select your operating system from the pull down menu and download.

Unpack into a /sci2 directory.

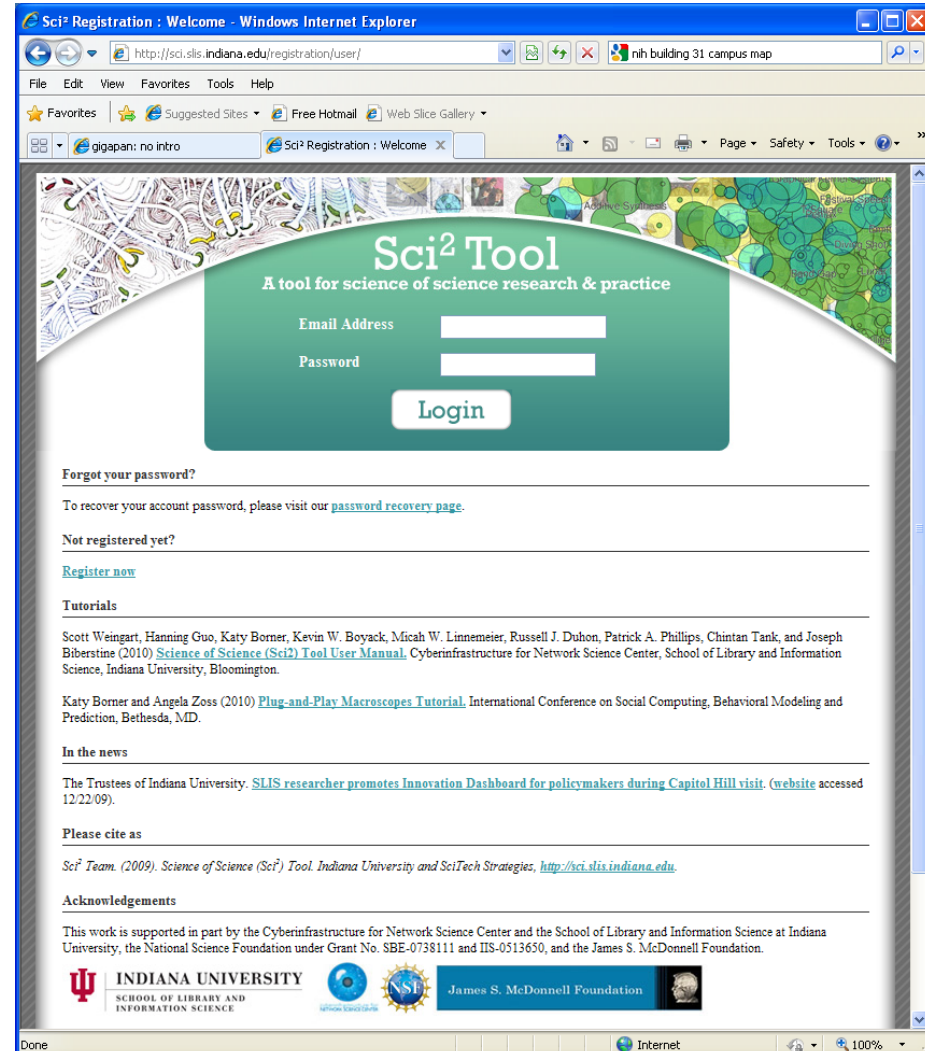
Run /sci2/sci2.exe

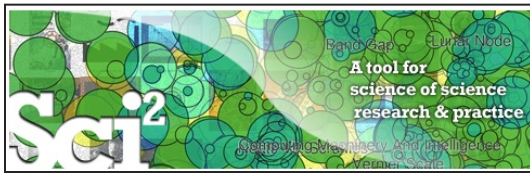
Sci² Manual is at

<http://sci2.wiki.cns.iu.edu>

Cite as

Sci² Team. (2009). Science of Science (Sci²) Tool. Indiana University and SciTech Strategies, <http://sci2.cns.iu.edu>

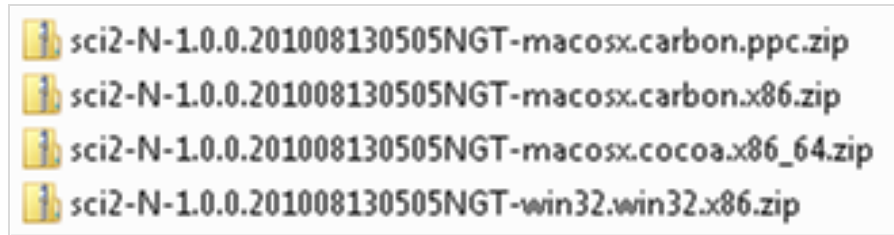




Sci² Tool: Download, Install, and Run

Sci2 Tool v0.5 Alpha (April 4, 2011)

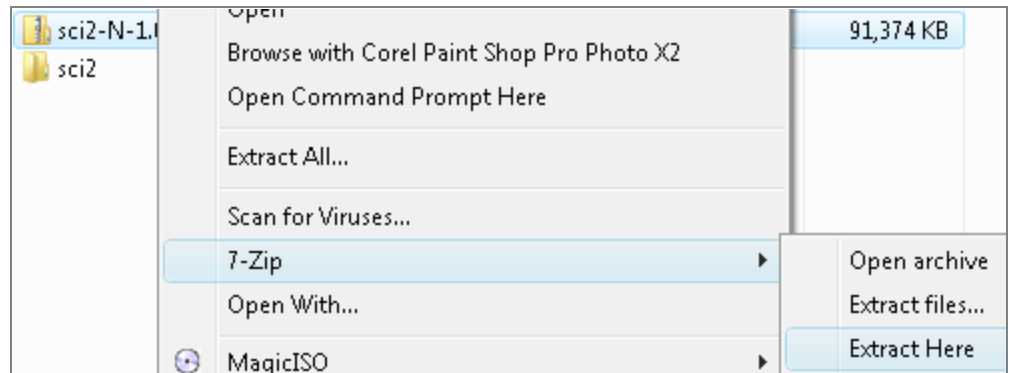
- Supports ASCII UTF-8 characters
- Web-based Yahoo! and desktop Geocoders
- U.S. and World geomapper
- Customizable stop word lists
- Merging of networks
- New home page, wiki-based tutorial
- Bug fixes, streamlined workflows

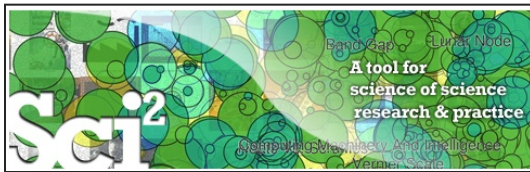


Sci2 Tool runs on Windows, Mac, and Linux.

Unzip.

Run /sci2/sci2.exe



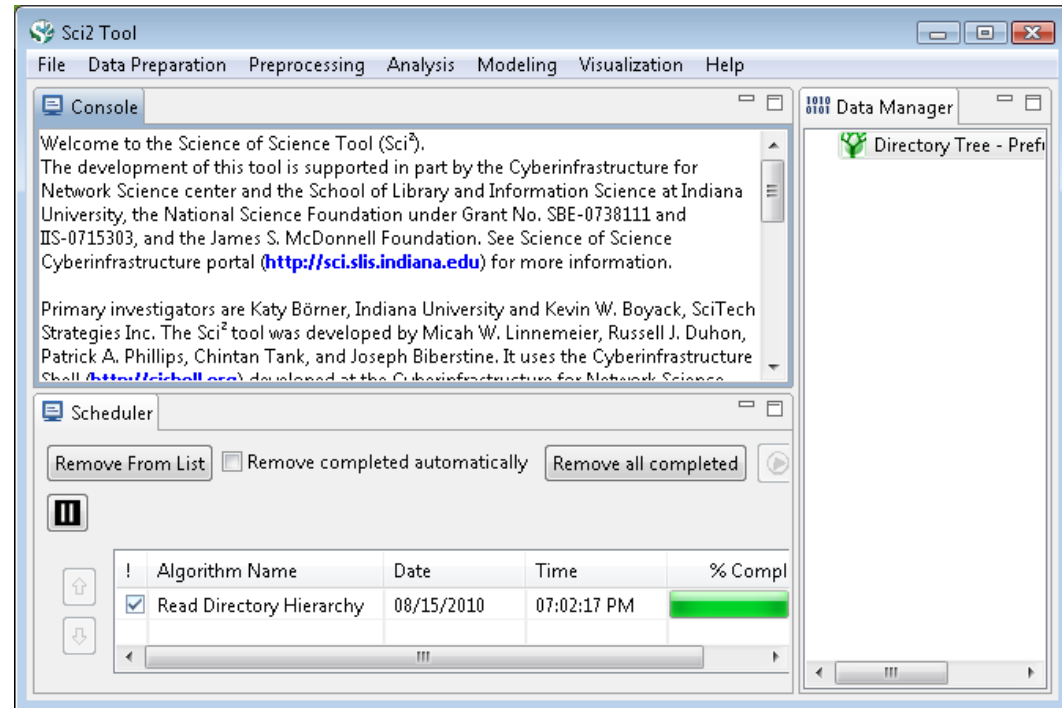


Sci2 Tool Interface Components

See also <http://sci2.wiki.cns.iu.edu/2.2+User+Interface>

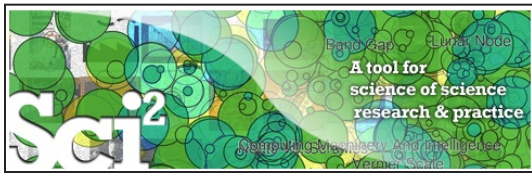
Use

- **Menu** to read data, run algorithms.
- **Console** to see work log, references to seminal works.
- **Data Manager** to select, view, save loaded, simulated, or derived datasets.
- **Scheduler** to see status of algorithm execution.



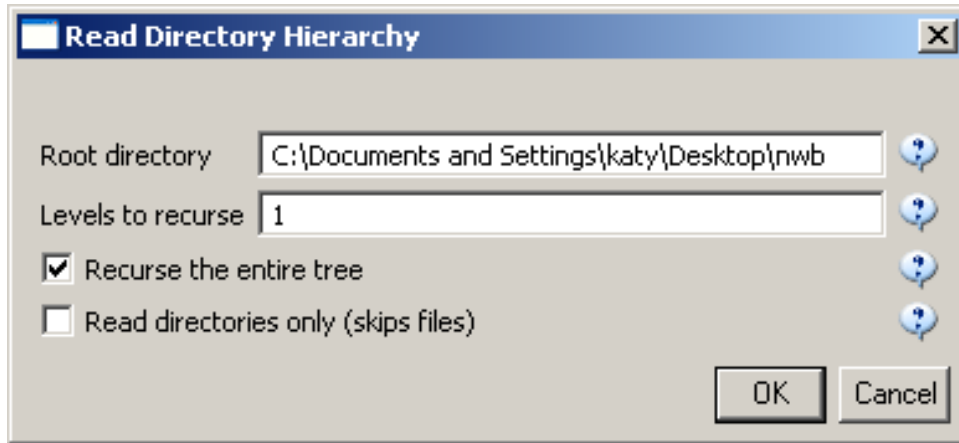
All workflows are recorded into a log file (see /sci2/logs/...), and soon can be re-run for easy replication. If errors occur, they are saved in a error log to ease bug reporting.

All algorithms are documented online; workflows are given in tutorials, see Sci2 Manual at <http://sci2.wiki.cns.iu.edu>



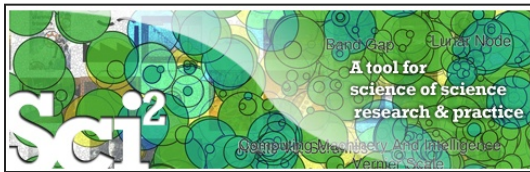
Sci2 Tool – Read+Visualize Sci2 Tool Directory Tree

Use '*File > Read Directory Hierarchy*' with parameters



Visualize resulting '*Directory Tree - Prefuse (Beta) Graph*' using

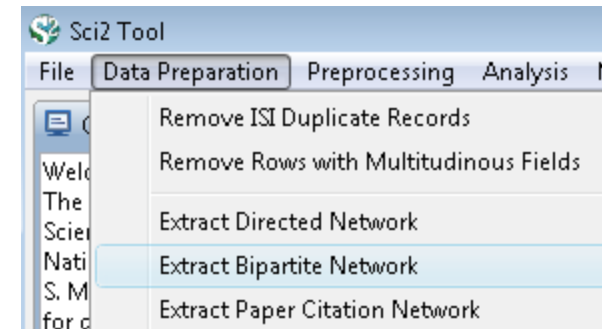
- '*Visualization > Tree View (prefuse beta)*'
- '*Visualization > Tree Map (prefuse beta)*'
- '*Visualization > Balloon Graph (prefuse alpha)*'
- '*Visualization > Radial Tree/ Graph (prefuse alpha)*'



Sci2 Tool – Visualize Workshop Attendees

Use *'File > Read'* to load *SciTS Conf SNA Registrants report 4.10.11-clean.csv*

	A	B	C
1	Last Name	Org	Organization-Cleaned
2	AgoulNIK	edu	Brigham and Women's Hospital, Harvard Medical School
3	Amaral	edu	Northwestern University
4	Bates	edu	University of Illinois at Chicago
5	Bennett	gov	NIH
6	Bietz	edu	University of California, Irvine
7	Bishop	edu	University of Tennessee
8	...		
9	Lotrecchiano	org,edu	George Washington University
10	Lusina	ca,edu	Centre for Hip Health & Mobility



Run *'Data Preparation > Extract Bipartite Network'*

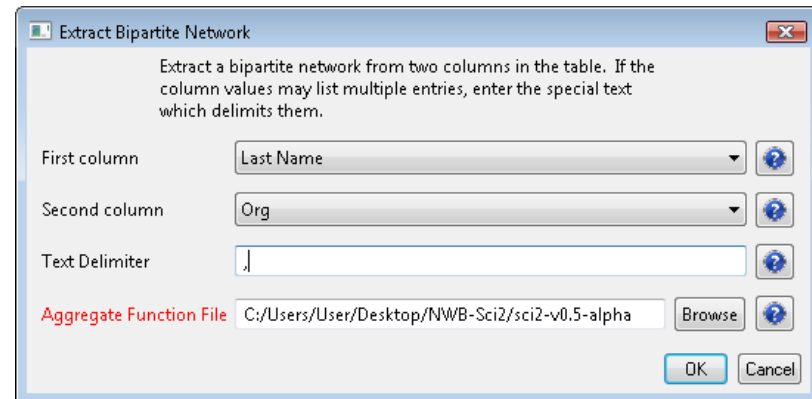
With parameter values:

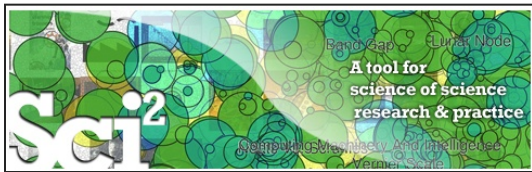
Optional: Calculate Node Degree

Visualize resulting *'Bipartite network*

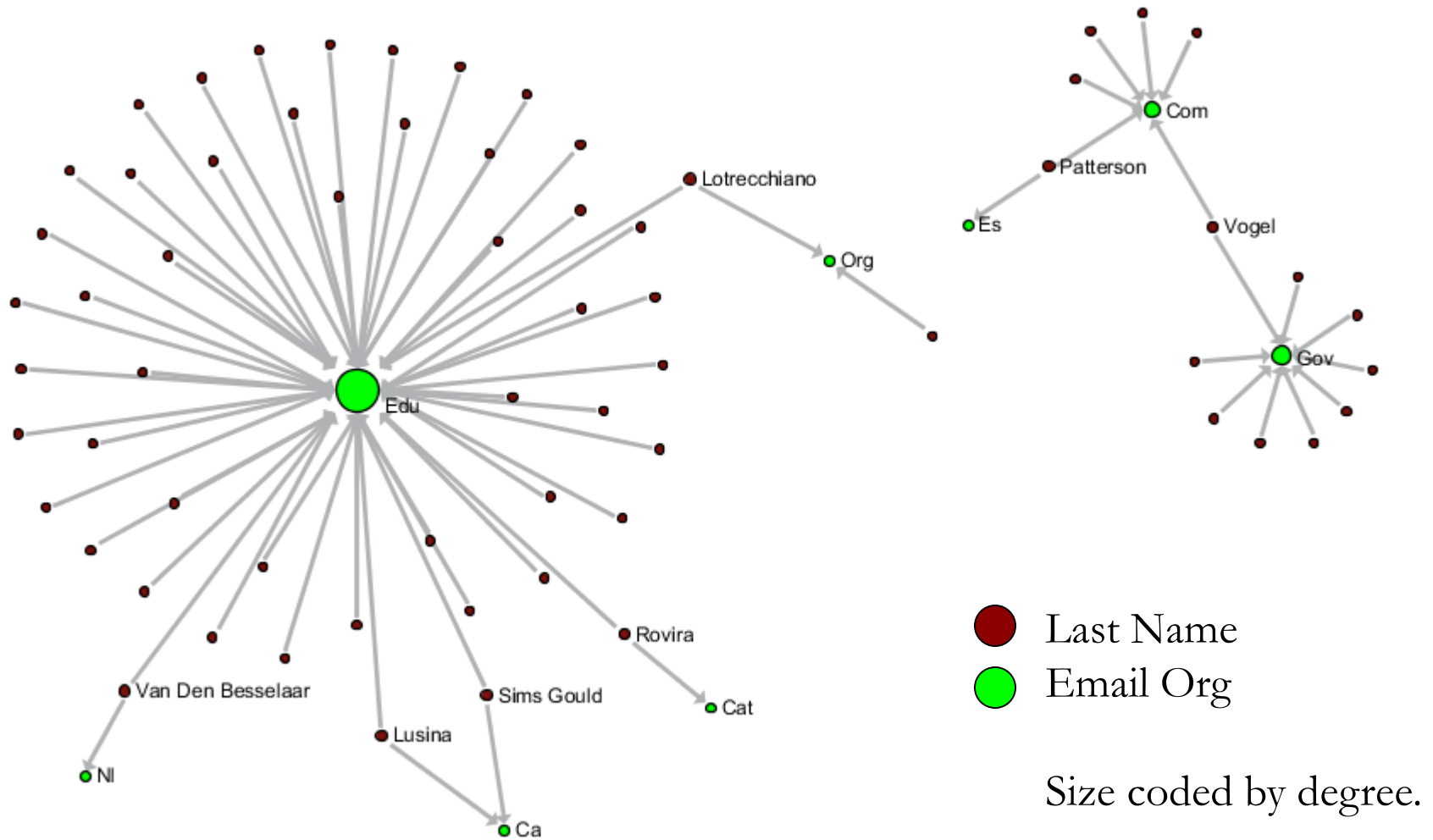
from Last Name and Org' using *'Visualization > Network > GUESS'* and

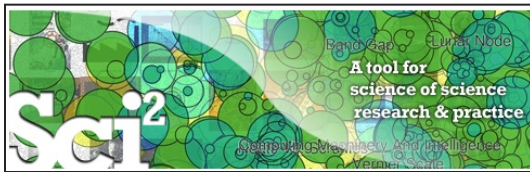
'Layout > GEM', 'Layout > Bin Pack'





Sci2 Tool – Visualize Workshop Attendees

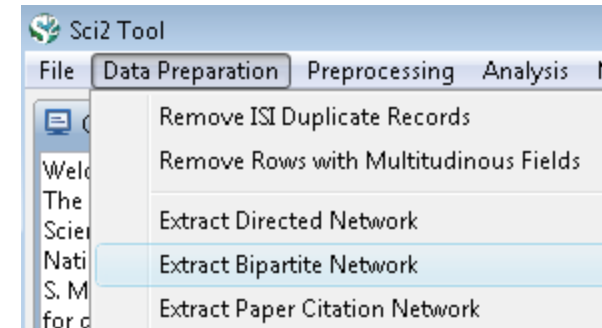




Sci2 Tool – Visualize Workshop Attendees

Use 'File > Read' to load *SciTS Conf SNA Registrants report 4.10.11-clean.csv*

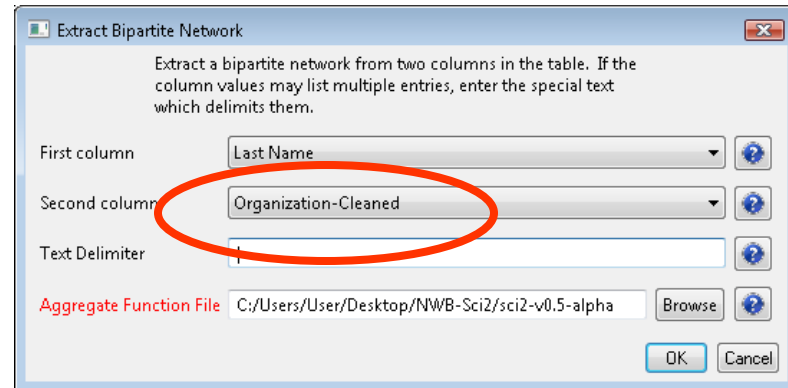
	A	B	C
1	Last Name	Org	Organization-Cleaned
2	Agoulnik	edu	Brigham and Women's Hospital, Harvard Medical School
3	Amaral	edu	Northwestern University
4	Bates	edu	University of Illinois at Chicago
5	Bennett	gov	NIH
6	Bietz	edu	University of California, Irvine
7	Bishop	edu	University of Tennessee
8	...		
9	Lotrecchiano	org,edu	George Washington University
10	Lusina	ca,edu	Centre for Hip Health & Mobility



Run 'Data Preparation > Extract Bipartite Network'

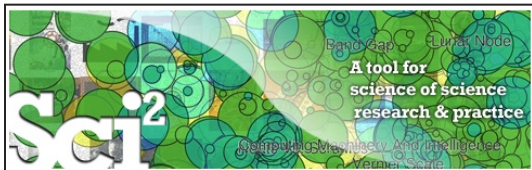
With parameter values:

Optional: Calculate Node Degree

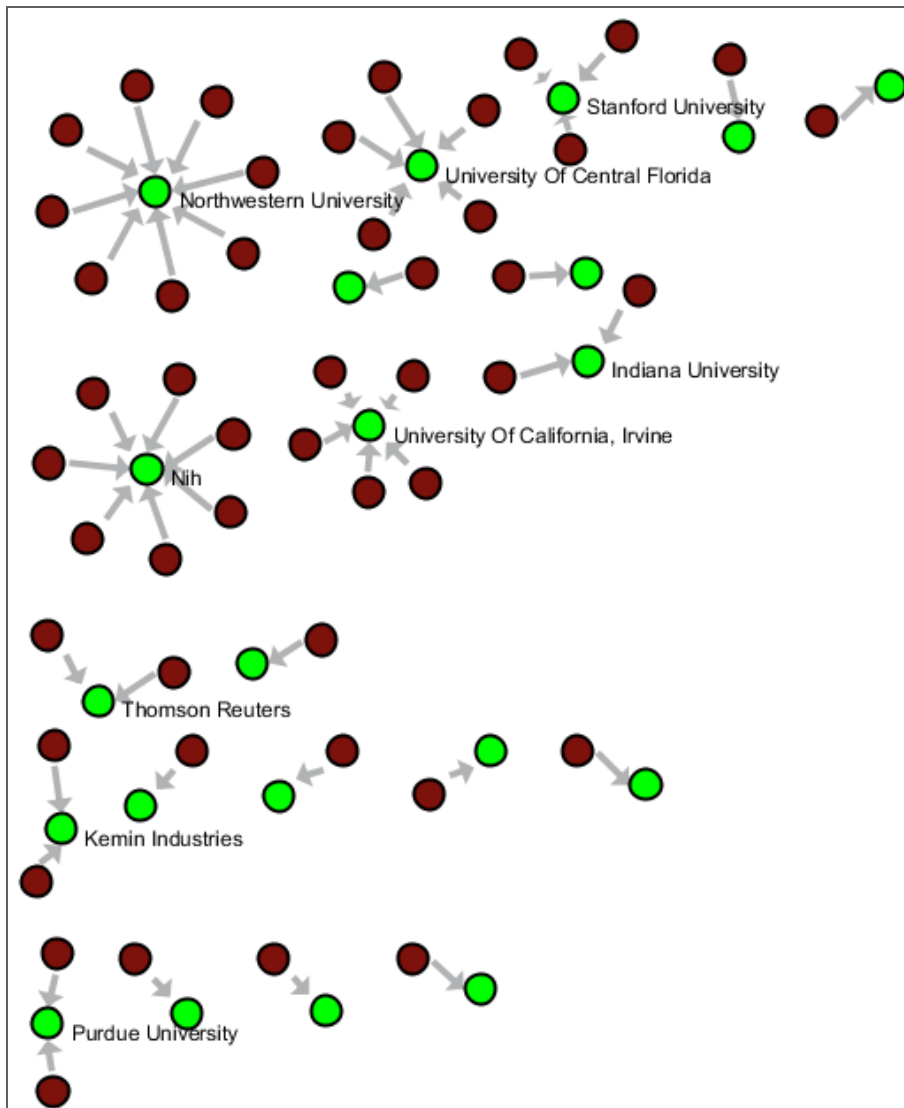
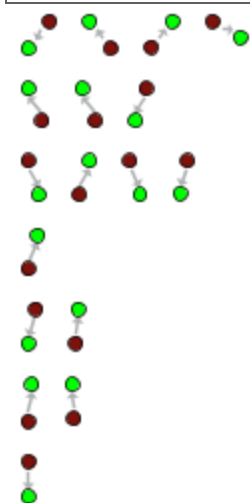
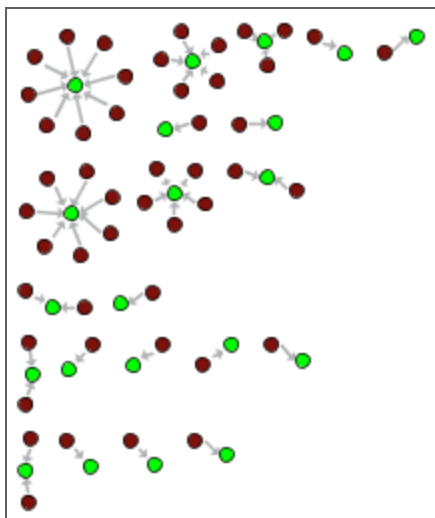


Visualize resulting *Bipartite network*

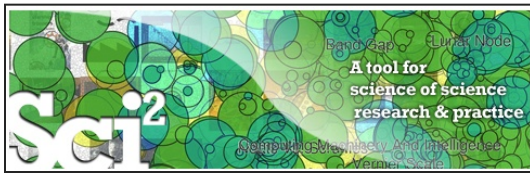
from *Last Name* and *Org*' using 'Visualization > Network > GUESS' and 'Layout > GEM', 'Layout > Bin Pack'



Sci2 Tool – Visualize Workshop Attendees



- Last Name
- Affiliation



Sci2 Tool – Visualize SciTS Co-Author Network Based on Holly’s EndNote File

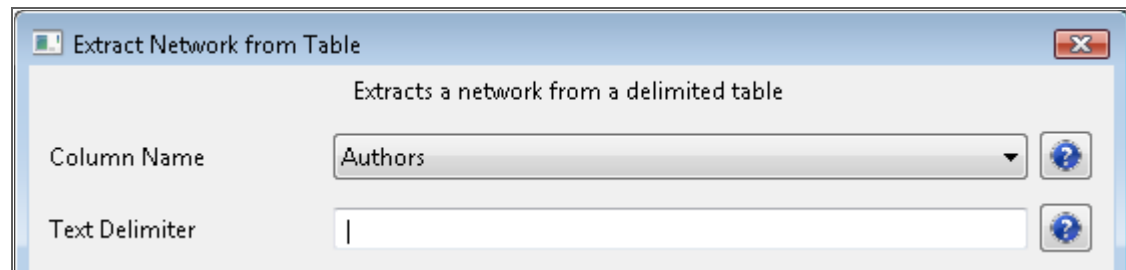
Open Holly’s ‘*SciTS-Library-03-04-2011.enl*’ in EndNote and save as ‘*SciTS-Library-03-04-2011.enw*’ following instructions on

<http://cishell.wiki.cns.iu.edu/Endnote+Export+Format>

Use ‘*File > Read*’ to load ‘*SciTS-Library-03-04-2011.enw*’

Run ‘*Data Preparation > Extract Co-Occurrence Network*’

With parameter values:

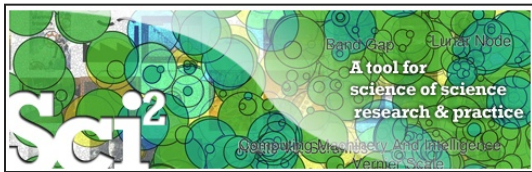


Optional: Calculate Node Degree

Visualize resulting ‘*Bipartite network*

from Last Name and Org’ using ‘*Visualization > Network > GUESS*’ and

‘*Layout > GEM*’, ‘*Layout > Bin Pack*’



Sci2 Tool – Visualize SciTS Co-Author Network Based on Holly's EndNote File

.....

Network Analysis Toolkit (NAT) was selected.

Implementer(s): Timothy Kelley

Integrator(s): Timothy Kelley

Reference: Robert Sedgewick. Algorithms in Java, Third Edition, Part 5 - Graph Algorithms. Addison-Wesley, 2002. ISBN 0-201-31663-3. Section 19.8, pp.205

Documentation:

<http://wiki.cns.iu.edu/display/CISHELL/Network+Analysis+Toolkit+%28NAT%29>

This graph claims to be undirected.

Nodes: 706

Isolated nodes: 100

Node attributes present: label

Edges: 1687

No self loops were discovered.

No parallel edges were discovered.

Edge attributes:

Did not detect any nonnumeric attributes.

Numeric attributes:

minmaxmean

weight 151.15412

This network seems to be valued.

Average degree: 4.779

This graph is not weakly connected.

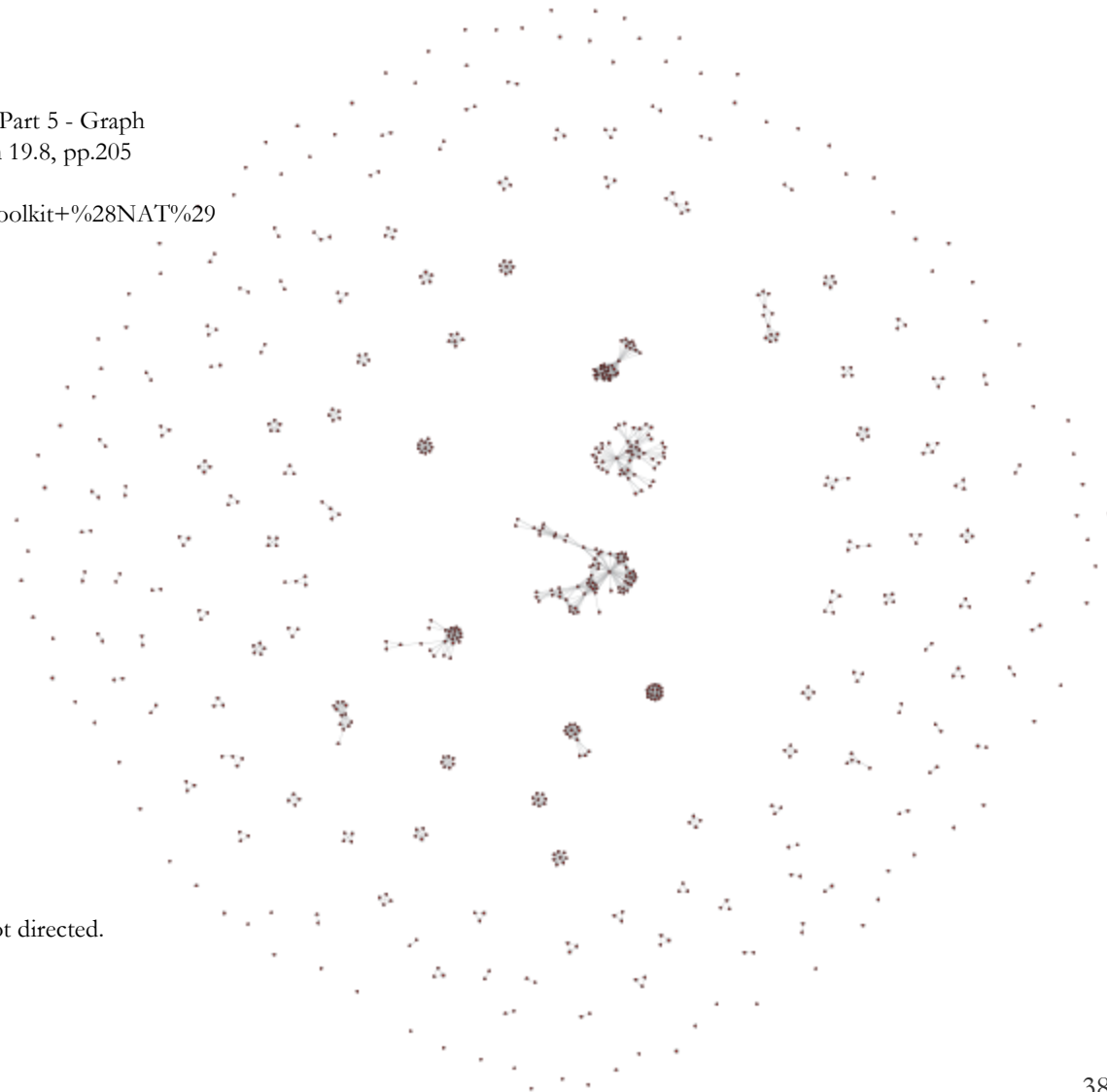
There are 223 weakly connected components. (100 isolates)

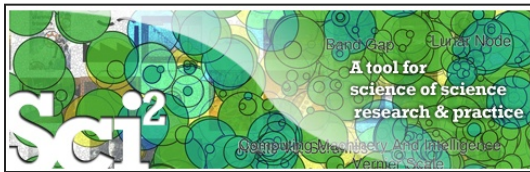
The largest connected component consists of 73 nodes.

Did not calculate strong connectedness because this graph was not directed.

Density (disregarding weights): 0.0068

Additional Densities by Numeric Attribute





Sci2 Tool – Visualize SciTS Co-Author Network Based on Holly's EndNote File

2nd Component



- Weak Component Cluster of 73 nodes
- Network with degree attribute added to node list.4
- Weak Component Cluster of 50 nodes
- Weak Component Cluster of 33 nodes
- Weak Component Cluster of 25 nodes
- Weak Component Cluster of 17 nodes
- Weak Component Cluster of 14 nodes
- Weak Component Cluster of 12 nodes
- Weak Component Cluster of 12 nodes.2
- Weak Component Cluster of 11 nodes
- Weak Component Cluster of 8 nodes

602	Stipelman, B.	589 *
603	Stokols, D.	200 *
604	Stokols, Daniel	75 *
605	Stone, Anthony R.	606 *
606	Stringer, M.j.	608 *
607	Subramanian, S.	559 *
608	Sundstrom, Eric	219 *
609	Sung, N.	611 *
610	Sung, N. S.	628 *
611	Syme,	638 *
638	Tress, G.	661 *
639	Trochim, W.	596 *
640	Trochim, W. M.	408 *
641	Trochim, W. M. K.	48 *
642	Trochim, W.m.	444 *
643	Trochim, William	76 *
644	Trochim, William M. K.	666 *
645	Unger, J.	598 *
646	Uzzi, B.	346 *
647	Uzzi, Brian	68 *



Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

- Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

- Load and clean a dataset as text file; process raw data into networks.
- Find basic statistics and run various algorithms over the network.
- Visualize as either a circular hierarchy or network

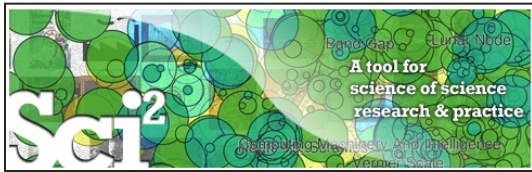
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

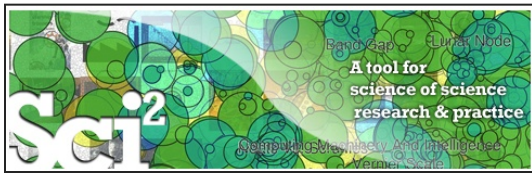
4:45 Outlook and Discussion

5:00 Adjourn



Padgett's Florentine Families - Compute Basic Network Properties & View in GUESS

- Florentine families related through business ties (specifically, recorded financial ties such as loans, credits and joint partnerships) and marriage alliances.
- Node attributes
 - Wealth: Each family's net wealth in 1427 (in thousands of lira)
 - Priorates: The number of priorates (seats on the civic council) held between 1282- 1344
 - Totalties: The total number of business or marriage ties in the total dataset of 116 families.
- “Substantively, the data include families who were locked in a struggle for political control of the city of Florence around 1430. Two factions were dominant in this struggle: one revolved around the infamous Medicis, the other around the powerful Strozzi.”
- <http://svitsrv25.epfl.ch/R-doc/library/ergm/html/florentine.html>



Padgett's Florentine Families - Compute Basic Network Properties & View in GUESS

- Load **yoursci2directory*/sampledata/socialscience/florentine.nwb*
- Run '*Analysis > Network Analysis Toolkit (NAT)*' to get basic properties.

This graph claims to be undirected.

Nodes: 16

Isolated nodes: 1

Node attributes present: label, wealth, totalities, priorates

Edges: 27

No self loops were discovered.

No parallel edges were discovered.

Edge attributes:

Nonnumeric attributes:

Example value

marriag...T

busines...F

Average degree: 3.375

There are 2 weakly connected components. (1 isolates)

The largest connected component consists of 15 nodes.

Did not calculate strong connectedness because this graph was not directed.

Density (disregarding weights): 0.225

- Select network and run '*Visualization > GUESS*' to open GUESS with file loaded.
- Apply '*Layout > GEM*'.

Network Workbench Tool
File Preprocessing Modeling Analysis Visualization Scientometrics Help

Console

```

.....
GUESS was selected.
Author(s): Eytan Adar
Implementer(s): Eytan Adar (GUESS), Russell Duhon (resizeLinear, colorize fix)
Integrator(s): Russell Duhon
Reference: Adar, Eytan, "GUESS: A Language and Interface for Graph Exploration," CHI 2006 (http://graphexploration.cond.org/)
Documentation: https://nwb.slis.indiana.edu/community/?n=VisualizeData.GUESS
ECHO is off.
Starting GUESS...
ECHO is off.
The initial layout for your visualization is random. For a clearer visualization, please run a layout from the Layout menu. (We recommend GEM.)
ECHO is off.
GUESS log file for this session can be found in
C:\DOCUME~1\katy\LOCALS~1\Temp\CIShell-Session-55892\StaticExecutableRunner-55904\ALGORITHM\guesslog.txt
.....
GUESS was selected.
Author(s): Eytan Adar
Implementer(s): Eytan Adar
Integrator(s): Russell Duhon
Reference: Adar, Eytan, "
Documentation: https://nwb.slis.indiana.edu/community/?n=VisualizeData.GUESS
ECHO is off.
Starting GUESS...
ECHO is off.
The initial layout for your visualization is random. For a clearer visualization, please run a layout from the Layout menu. (We recommend GEM.)
ECHO is off.
GUESS log file for this session can be found in
C:\DOCUME~1\katy\LOCALS~1\Temp\CIShell-Session-55892\StaticExecutableRunner-55904\ALGORITHM\guesslog.txt

```

Scheduler

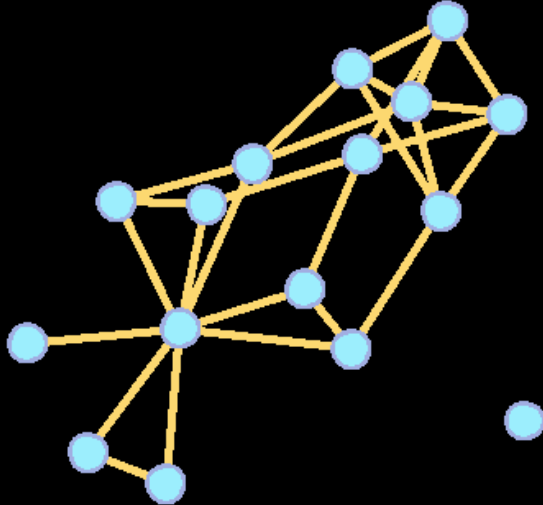
Remove From List ☐

Information window

Field	Value
__edgeid	0
business	F
color	dandelion
directed	false
label	
labelcolor	0,0,0,255
labelsize	12
labelvisible	false
marriage	T
node1	n9
node2	n1
visible	true
weight	1.0
width	2.0

Visualization - GUESS
File Edit Layout Script View Help

Medici-Acciaiuoli



Data Manager

- NWB file: C:\Documents and Settings\katy\Documents\Distribution of degree for network at stu...
- List of edges of network created through the...
- Distribution of degree for network at stu...
- Distribution of degree for network at stu...
- NWB file: C:\Documents and Settings\katy\Documents\Graph and Network Analysis Log
- Sequence of site betweennesses for network...
- Distribution of site betweenness for network...
- Distribution of site betweenness for network...
- NWB file with site betweenness attribute...

Interpreter **Graph Modifier**

Object: Property: Operator: Value:

Colour Show Hide Size Show Label Hide Label Change Label

Node Shape Center Change History

Visualization - GUESS

File Edit Layout Script View Help

Medici

Field	Value
color	cornflo...
fixed	false
height	10.0
image	
label	Medici
labelcolor	0,0,0,...
labelsize	12
labelvi...	false
name	n9
original...	Medici
priorates	53
stroke...	cadetb...
style	2
totalities	54
visible	true
wealth	103
width	10.0
x	90.625...
y	44.312...

Information Window

Graph visualization showing nodes (blue circles) and edges (yellow lines). A context menu is open over a node, listing options: Center On, Color..., Remove, Add, Modify Field..., Copy as Variable...

Graph Modifier

Object: Property: Operator: Value:

Buttons: Colour, Show, Hide, Size, Show Label, Hide Label, Change Label, Format Node Labels, Format Edge Labels, Node Shape, Center, Change History, Resize Linear, Colorize, Nodes, labels, From: To: Do Resize Linear


Interpreter

Pan:

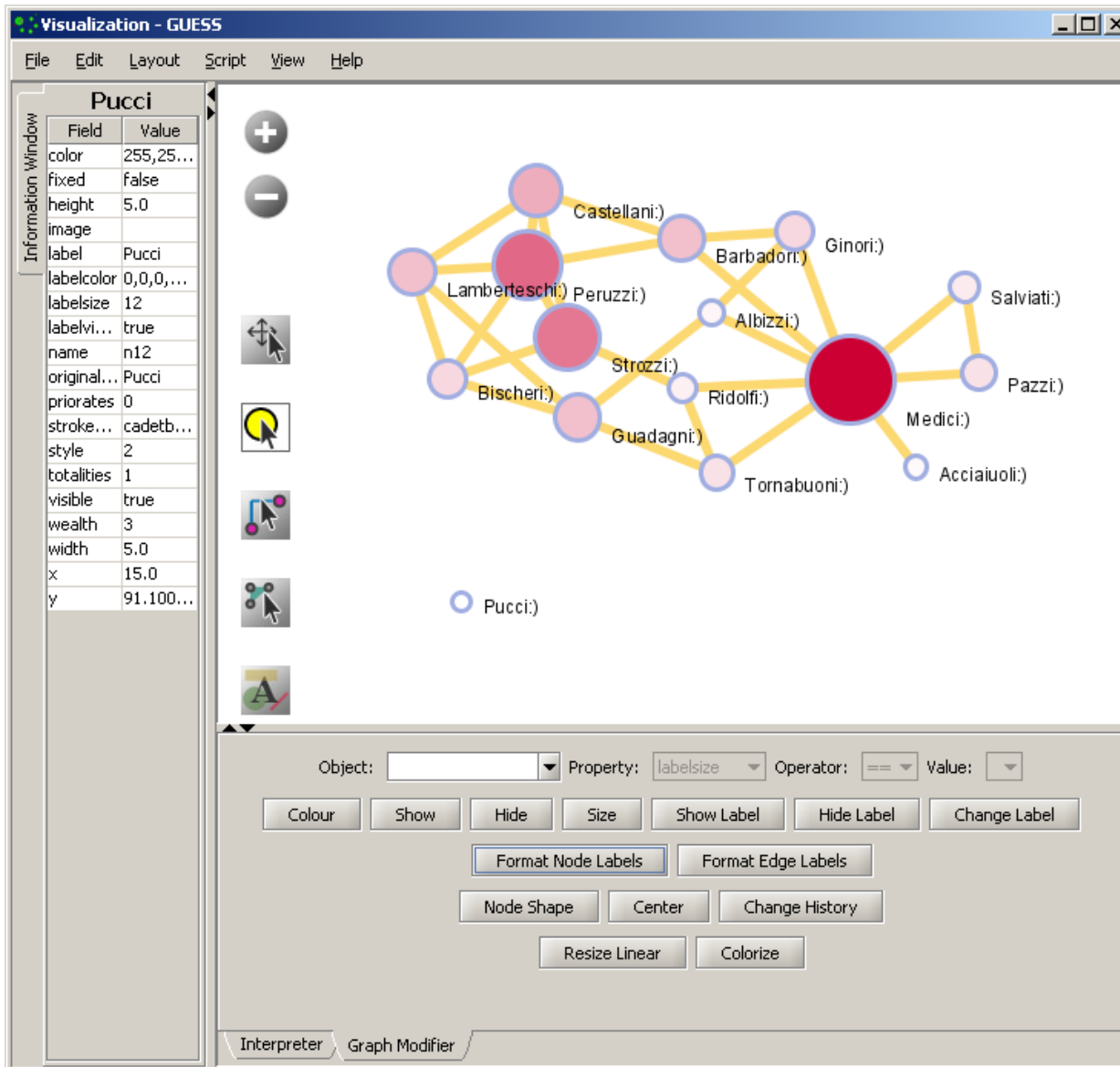
“grab” the background by holding left-click and moving your mouse.

Zoom:

Using scroll wheel, press the “+” and “-” buttons in the upper-left hand corner, or right-click and move the mouse left or right. Center graph by selecting ‘View -> Center’.

Select  to select/move single nodes. Hold down ‘Shift’ to select multiple.

Right click to modify Color, etc.



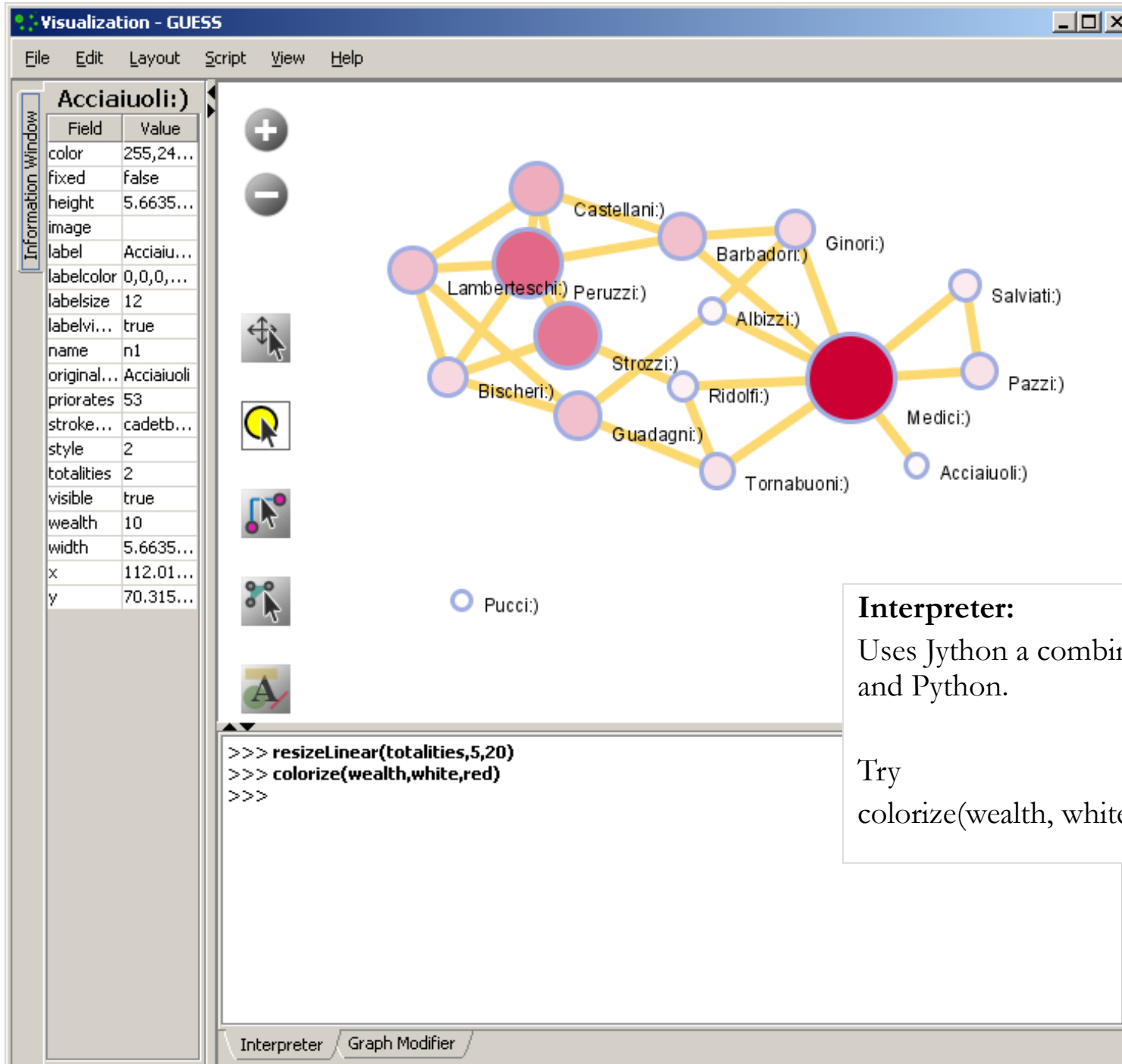
Graph Modifier:

Select “all nodes” in the Object drop-down menu and click ‘Show Label’ button.

Select ‘Resize Linear > Nodes > totalities’ drop-down menu, then type “5” and “20” into the From” and To” Value box separately. Then select ‘Do Resize Linear’.

Select ‘Colorize> Nodes>totalities’, then select white and enter (204,0,51) in the pop-up color boxes on in the “From” and “To” buttons.

Select “Format Node Labels”, replace default text {originallabel} with your own label in the pop-up box ‘Enter a formatting string for node labels.’



Interpreter:

Uses Jython a combination of Java and Python.

Try
colorize(wealth, white, red)



Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

➤ Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

➤ **Load and clean a dataset; process raw data into networks.**

➤ **Find basic statistics and run various algorithms over the network.**

➤ **Visualize as either a circular hierarchy or network.**

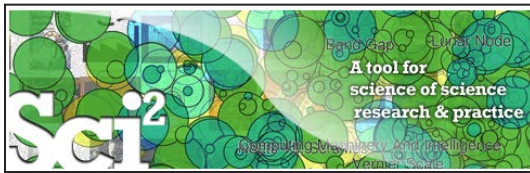
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

4:45 Outlook and Discussion

5:00 Adjourn



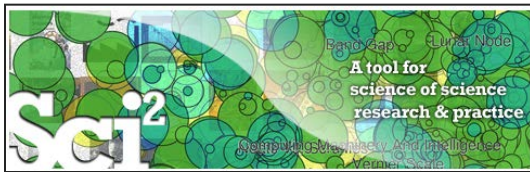
Studying Four Major NetSci Researchers (ISI Data) using Database (*section 5.1.4*)

FourNetSciResearchers.isi	
Time frame:	1955-2007
Region(s):	Miscellaneous
Topical Area(s):	Network Science
Analysis Type(s):	Paper Citation Network, Co-Author Network, Bibliographic Coupling Network, Document Co-Citation Network, Word Co-Occurrence Network

Thomson Reuter's Web of Knowledge (WoS) is a leading citation database. Access it via the "Web of Science" tab at <http://www.isiknowledge.com> (**note:** access to this database requires a paid subscription). Along with Scopus, WoS provides some of the most comprehensive datasets for scientometric analysis.

To find all publications by an author, search for the last name and the first initial followed by an asterisk in the author field.

[http://sci2.wiki.cns.iu.edu/5.1.4+Studying+Four+Major+NetSci+Researchers+\(ISI+Data\)](http://sci2.wiki.cns.iu.edu/5.1.4+Studying+Four+Major+NetSci+Researchers+(ISI+Data))



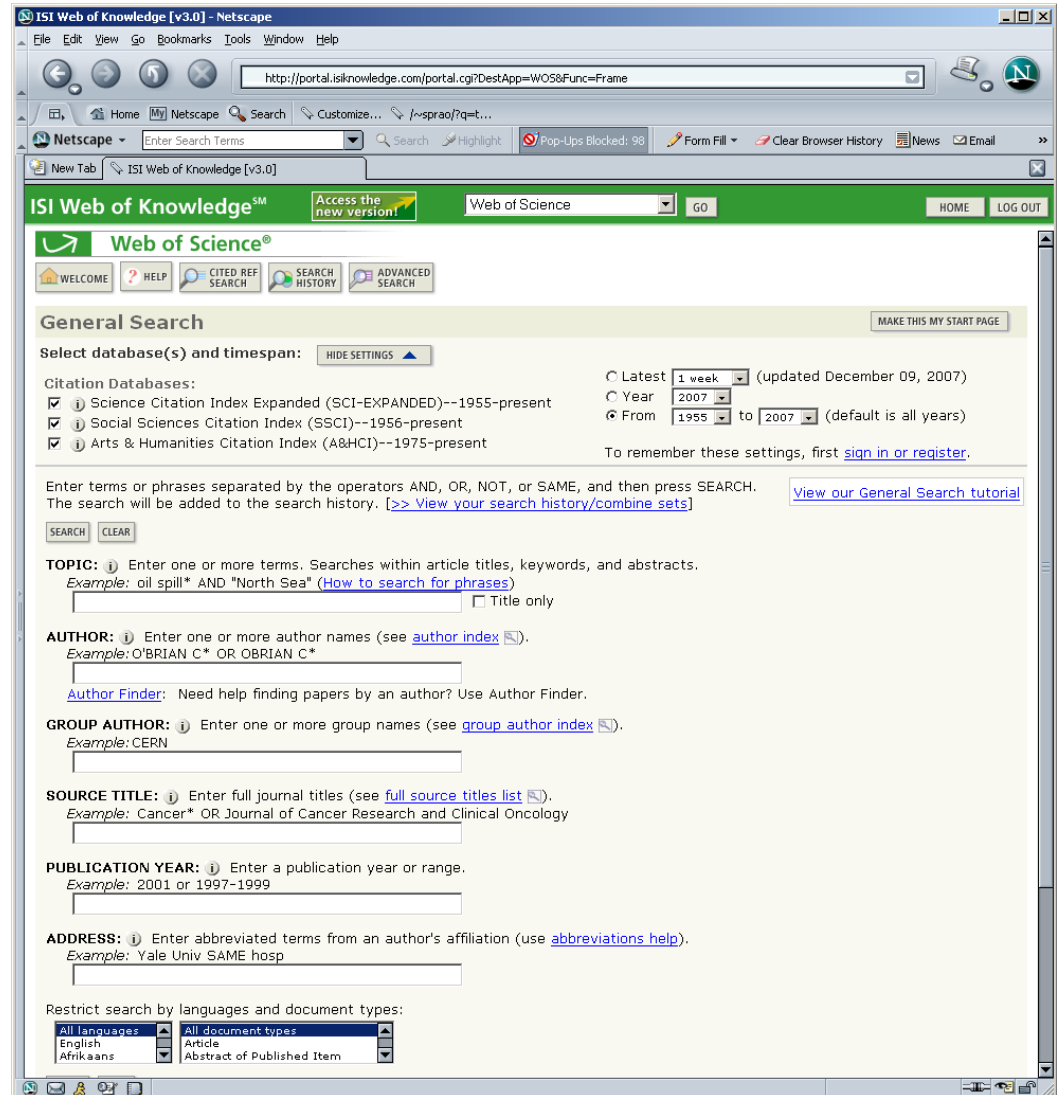
Data Acquisition from Web of Science

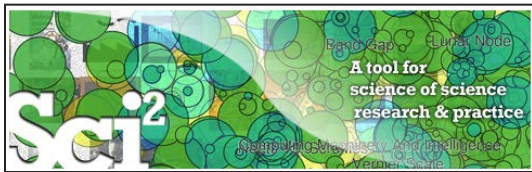
In Dec 2007, we downloaded
all papers by

- Eugene Garfield
- Stanley Wasserman
- Alessandro Vespignani
- Albert-László Barabási

from

- Science Citation Index
Expanded (SCI-EXPANDED)
--1955-present
- Social Sciences Citation Index
(SSCI)--1956-present
- Arts & Humanities Citation
Index (A&HCI)--1975-present

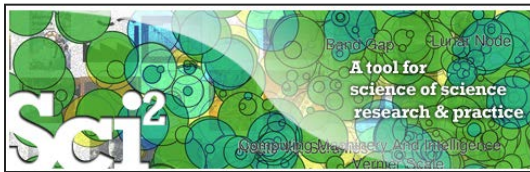




Comparison of Counts

No books and other non-WoS publications are covered.

	Age	Total # Cites	Total # Papers	H-Index
Eugene Garfield	82	1,525	672	31
Stanley Wasserman		122	35	17
Alessandro Vespignani	42	451	101	33
Albert-László Barabási	40	2,218	126	47 <i>(Dec 2007)</i>
	41	16,920	159	52 <i>(Dec 2008)</i>
	44	30,102	201	68 <i>(April 11)</i>



Extract Co-Author Network

Load **yoursci2directory*/sampledata/scientometrics/isi/FourNetSciResearchers.isi*
using *'File > Load ...'*

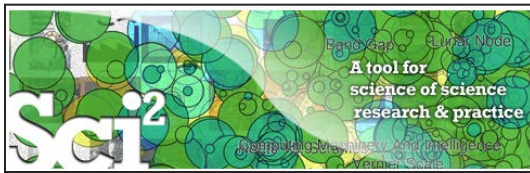
And file with 361 records appears in the Data Manager.

Duplicates were removed, author names normalized. Log file exists.

The screenshot shows the Sci2 Tool interface with the following components:

- Console:**
 - Loaded 361 records.
 - Removed 0 duplicate records.
 - Author names have been normalized.
 - 361 records with unique ISI IDs are available via Data Manager.
 - Wrote log to C:\Users\User\AppData\Local\Temp\isiduplicateremoverlog253473399342202281.txt
- Scheduler:**
 - Buttons: Remove From List, ☐ Remove completed automatically, Remove all complete
 - Table:
- Data Manager:**
 - ISI Data: C:\Users\User\Desktop\10-NEH-A&H-Workshop
 - 361 Unique ISI Records

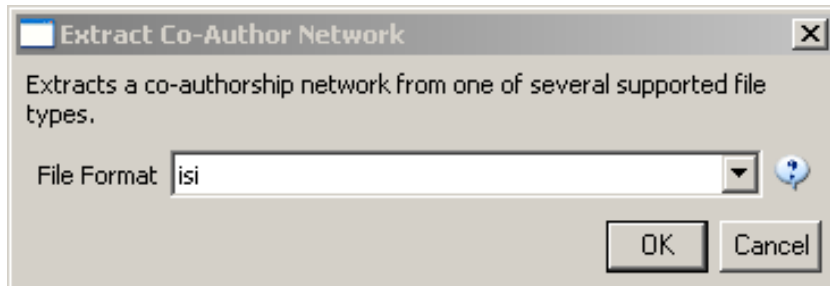
!	Algorithm Name	Date	Time	%
<input checked="" type="checkbox"/>	Load and Clean ISI File	08/15/2010	07:29:43 PM	100%
<input checked="" type="checkbox"/>	Load and Clean ISI File	08/15/2010	07:12:49 PM	100%



Extract Co-Author Network

(see section 5.1.4.2 on correcting duplicate/ misspelled author names)

To extract the co-author network, select the *'361 Unique ISI Records'* table and run *'Data Preparation > Extract Co-Author Network'* using isi file format:

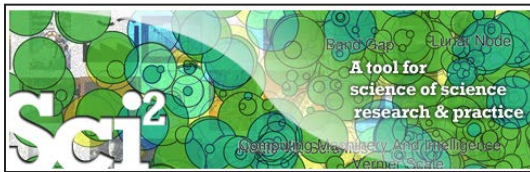


The result is an undirected but weighted network of co-authors in the Data Manager. Run *'Analysis > Network > Network Analysis Toolkit (NAT)'* to calculate basic properties: the network has 247 nodes and 891 edges.

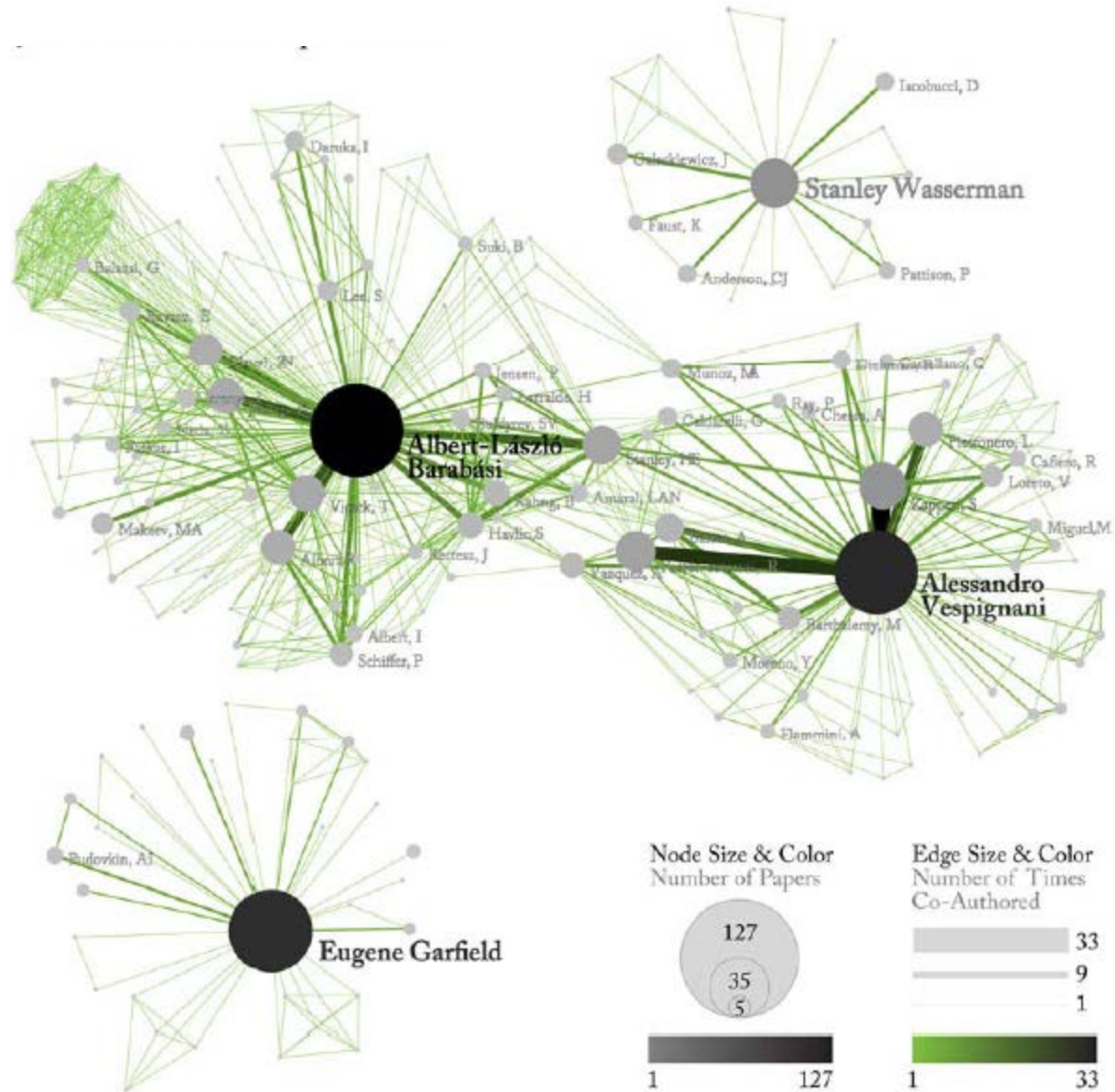
Use *'Analysis > Network > Unweighted and Undirected > Node Degree'* to calculate the number of neighbors for each node independent of co-authorship weight.

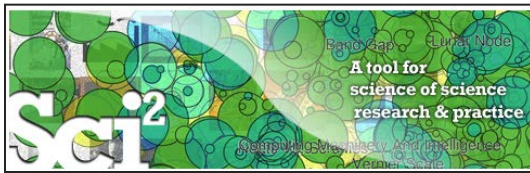
To view the complete network, select the *'Extracted Co-Authorship Network'* and run *'Visualization > Networks > GUESS'*.

Network is loaded with random layout. In GUESS, run *'Layout > GEM'* and *'Layout > Bin Pack'* to improve layout. Run *'Script > Run Script ...'* and select *'yoursci2directory/ scripts/ GUESS/ co-author-nw.py'*.



Co-Author Network of all Four NetsSci Researchers



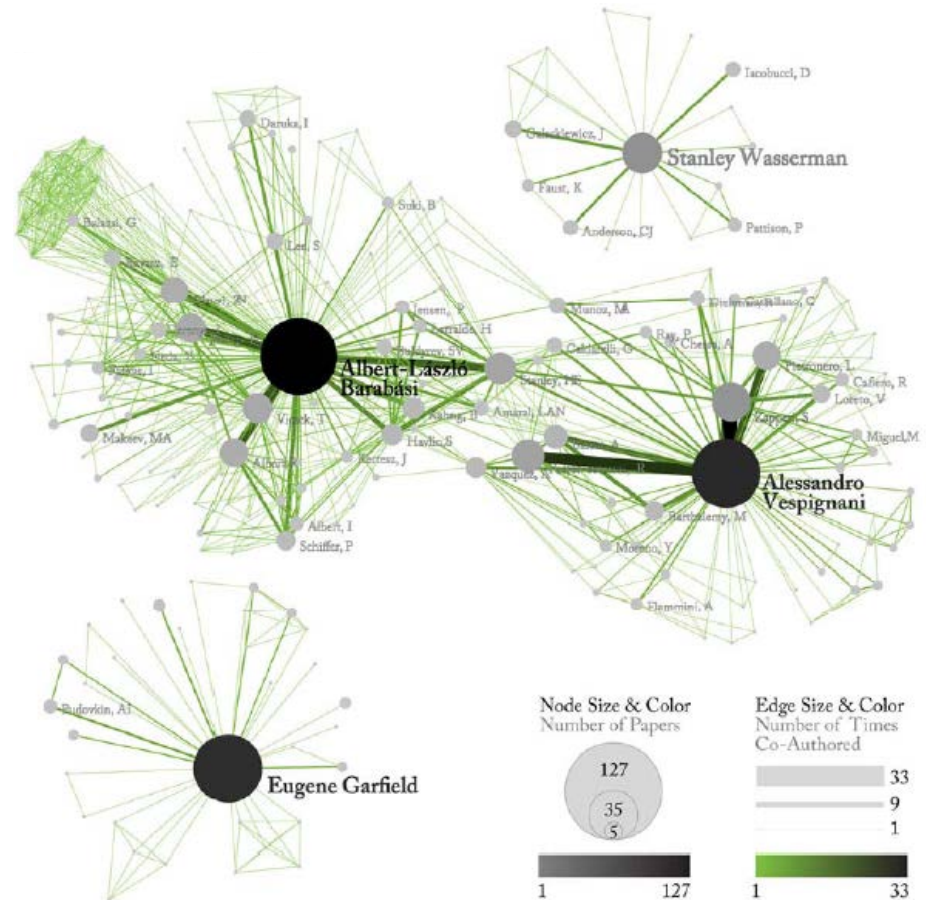


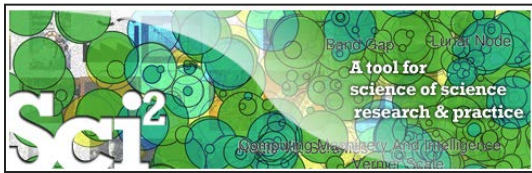
Co-Author Network of all Four NetsSci Researchers

Use the GUESS Graph Modifier to change
color and size coding.

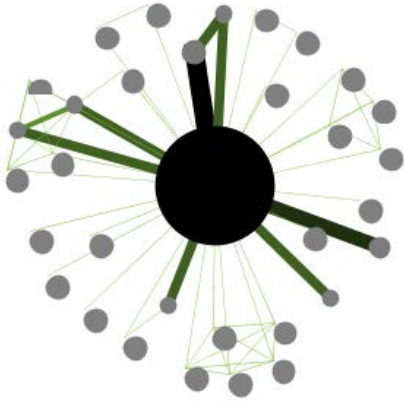
Calculate node degrees in Sci2 Tool.

Use a graphic program to add legend.

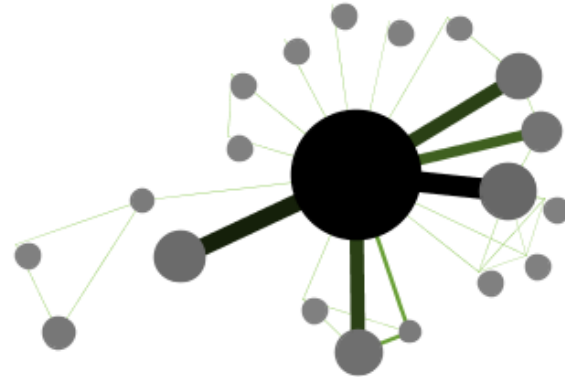




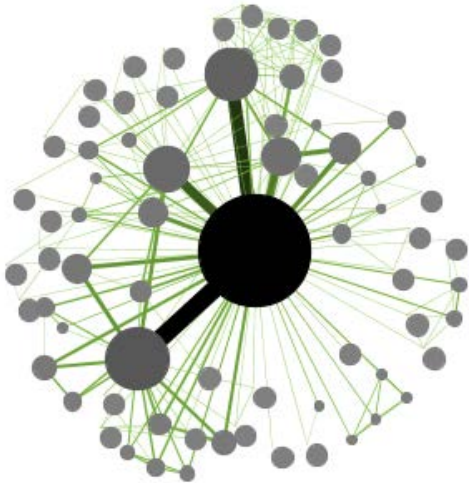
Individual Co-Author Networks (Read/map 4 files separately)



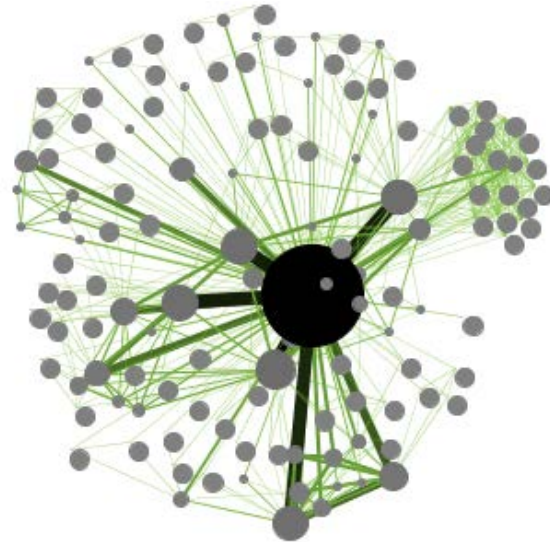
Eugene Garfield



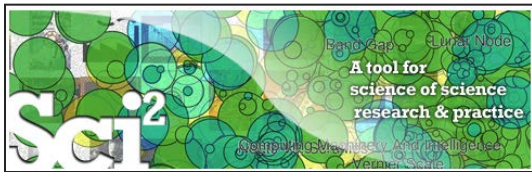
Stanley Wasserman



Alessandro Vespignani



Albert-László Barabási



Network Visualization: Node Layout

Load and Clean ISI File was selected.
Loaded 361 records.
Removed 0 duplicate records.
Author names have been normalized.
361 records with unique ISI IDs are available
via Data Manager.

.....

Extract Co-Author Network was selected.

Input Parameters:

File Format: isi

.....

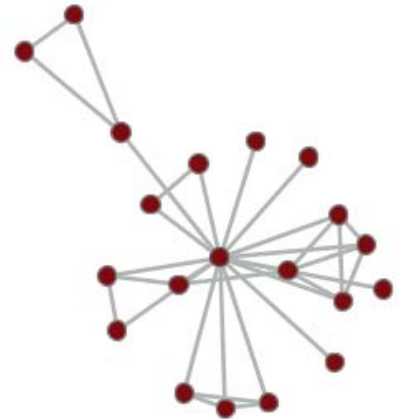
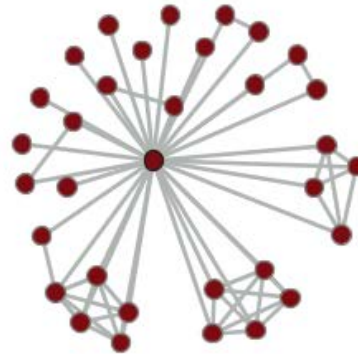
Network Analysis Toolkit (NAT) was selected.

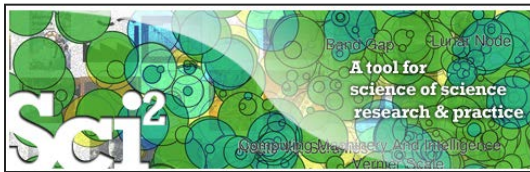
Nodes: 247

Edges: 891

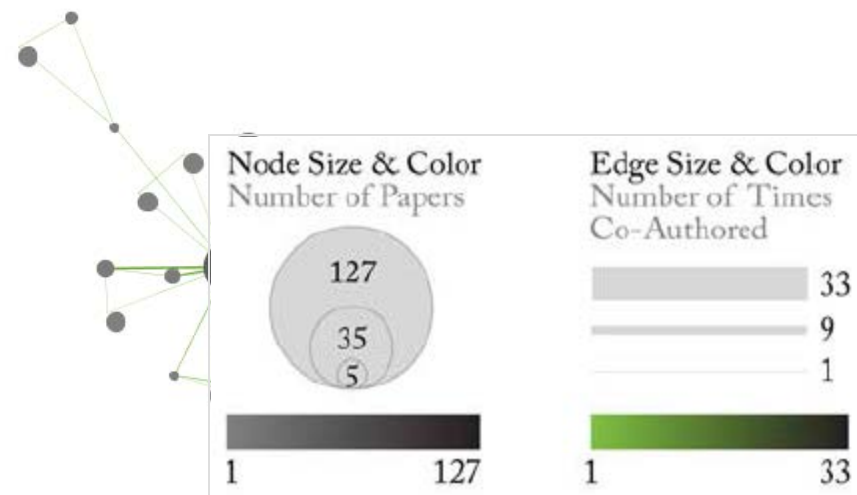
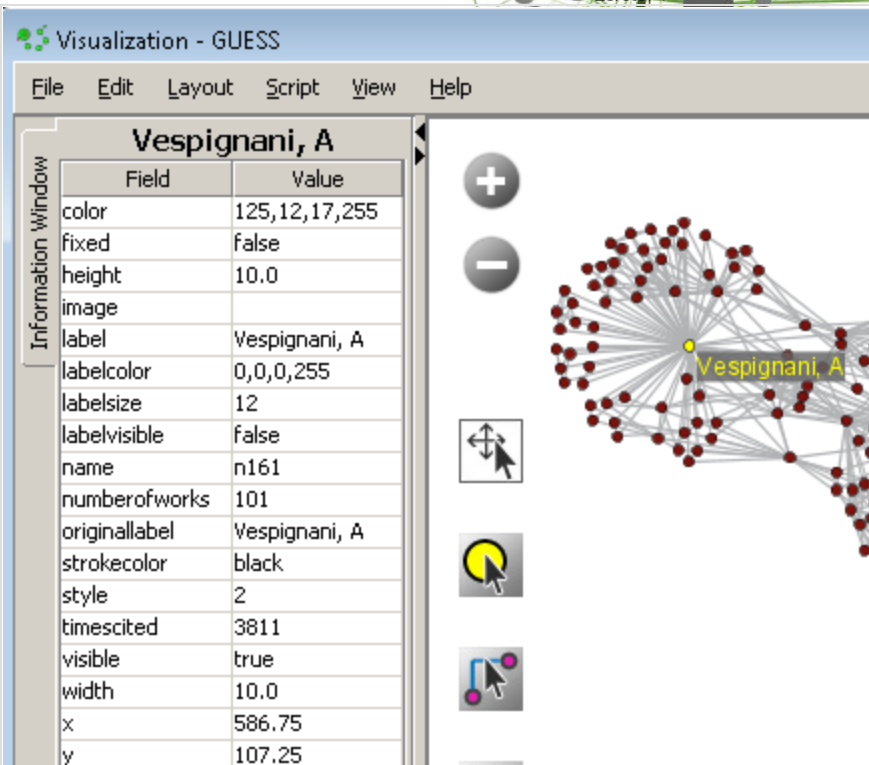
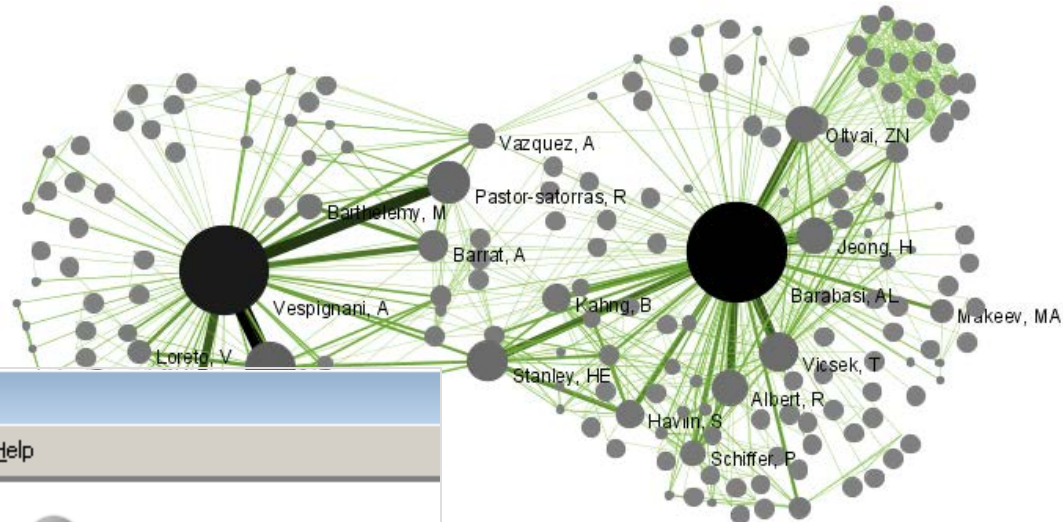
.....

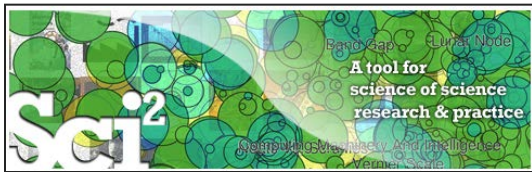
GUESS was selected.



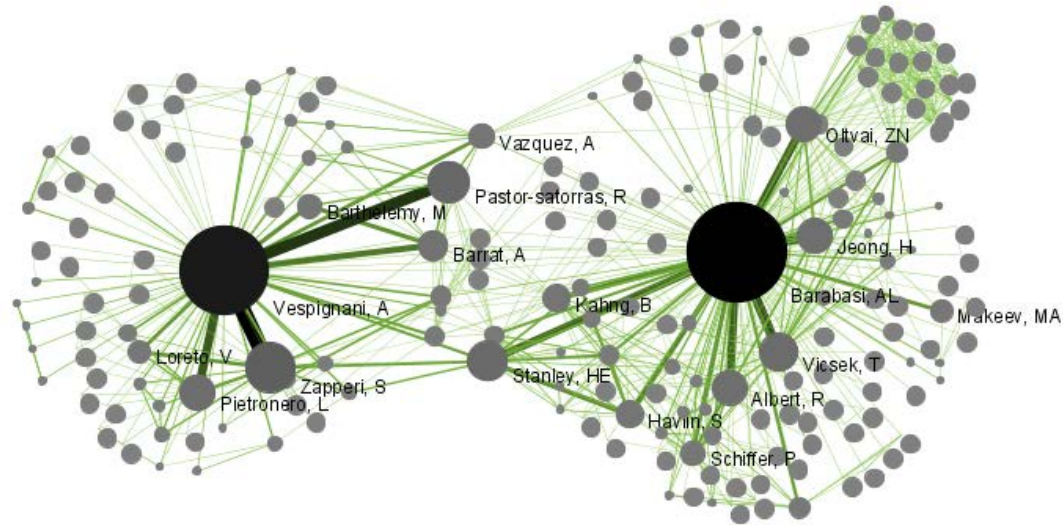


Network Visualization: Color/Size Coding by Data Attribute Values





Network Visualization: Giant Component



.....

Weak Component Clustering was selected.

Implementer(s): Russell Duhon

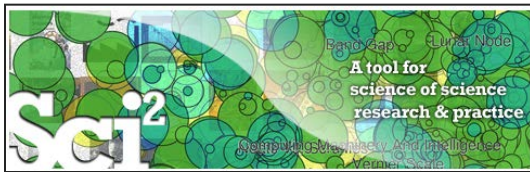
Integrator(s): Russell Duhon

Input Parameters:

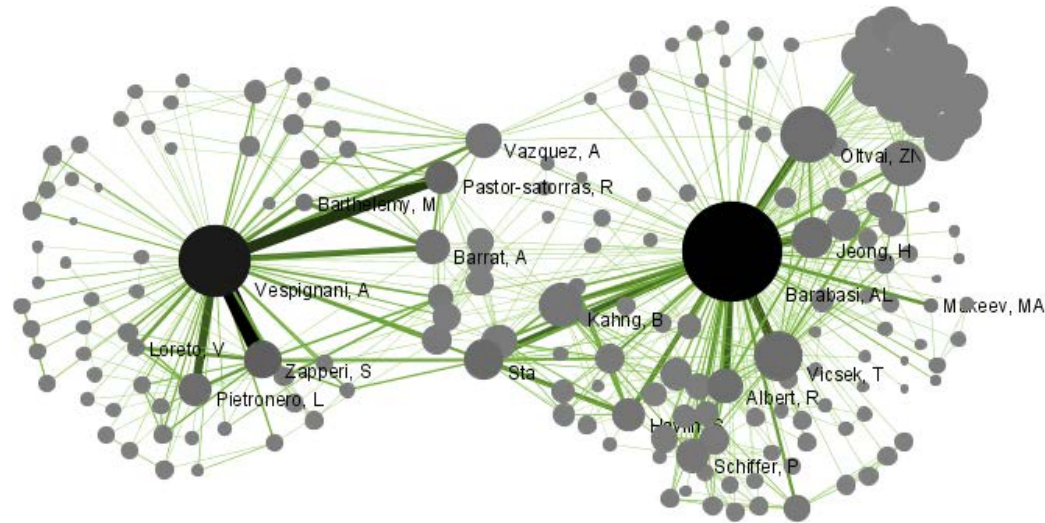
Number of top clusters: 10

3 clusters found, generating graphs for the top 3 clusters.

.....



Network Visualization: Color/Size Coding by Degree



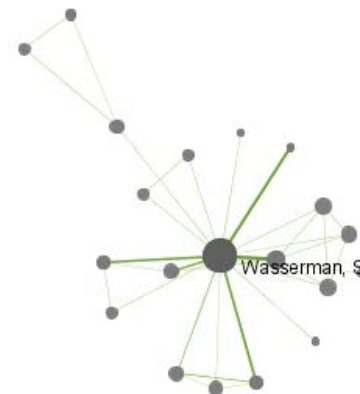
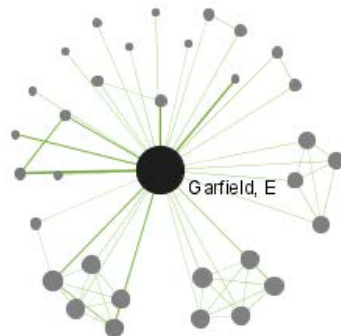
.....

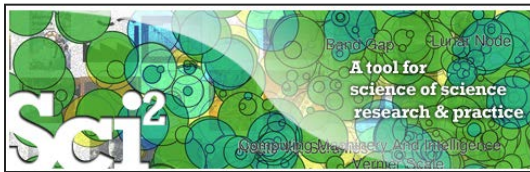
Node Degree was selected.

Documentation:

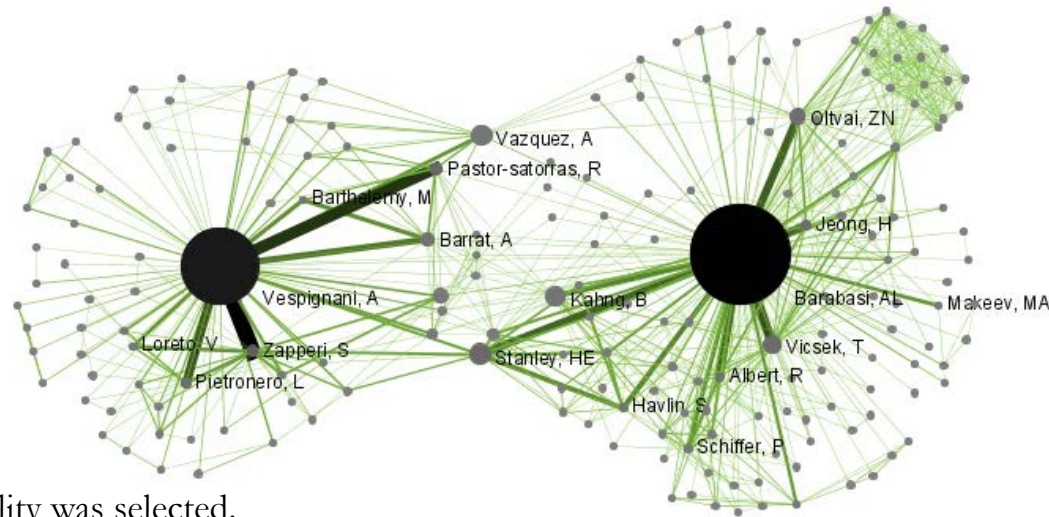
[https://nwb.slis.indiana.edu/community/?n=AnalyzeData.No deDegree](https://nwb.slis.indiana.edu/community/?n=AnalyzeData.No%20deDegree)

.....





Network Visualization: Color/Size Coding by Betweenness Centrality



.....

Node Betweenness Centrality was selected.

Author(s): L. C. Freeman

Implementer(s): Santo Fortunato

Integrator(s): Santo Fortunato, Weixia Huang

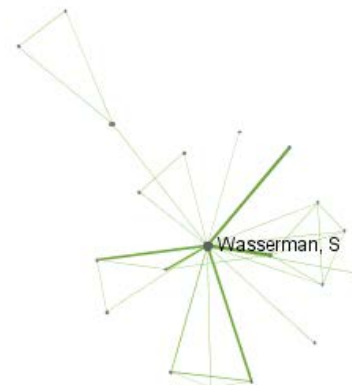
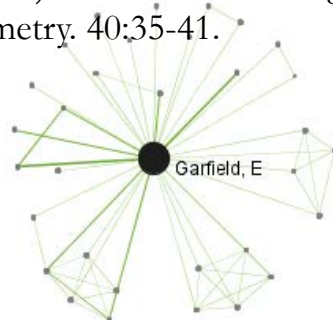
Reference: Freeman, L. C. (1977). A set of measuring centrality based on betweenness. Sociometry. 40:35-41.

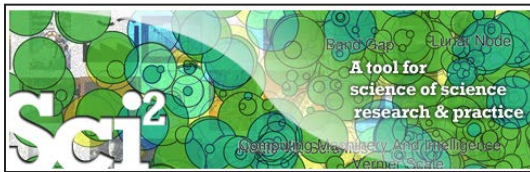
Input Parameters:

Number of bins: 10

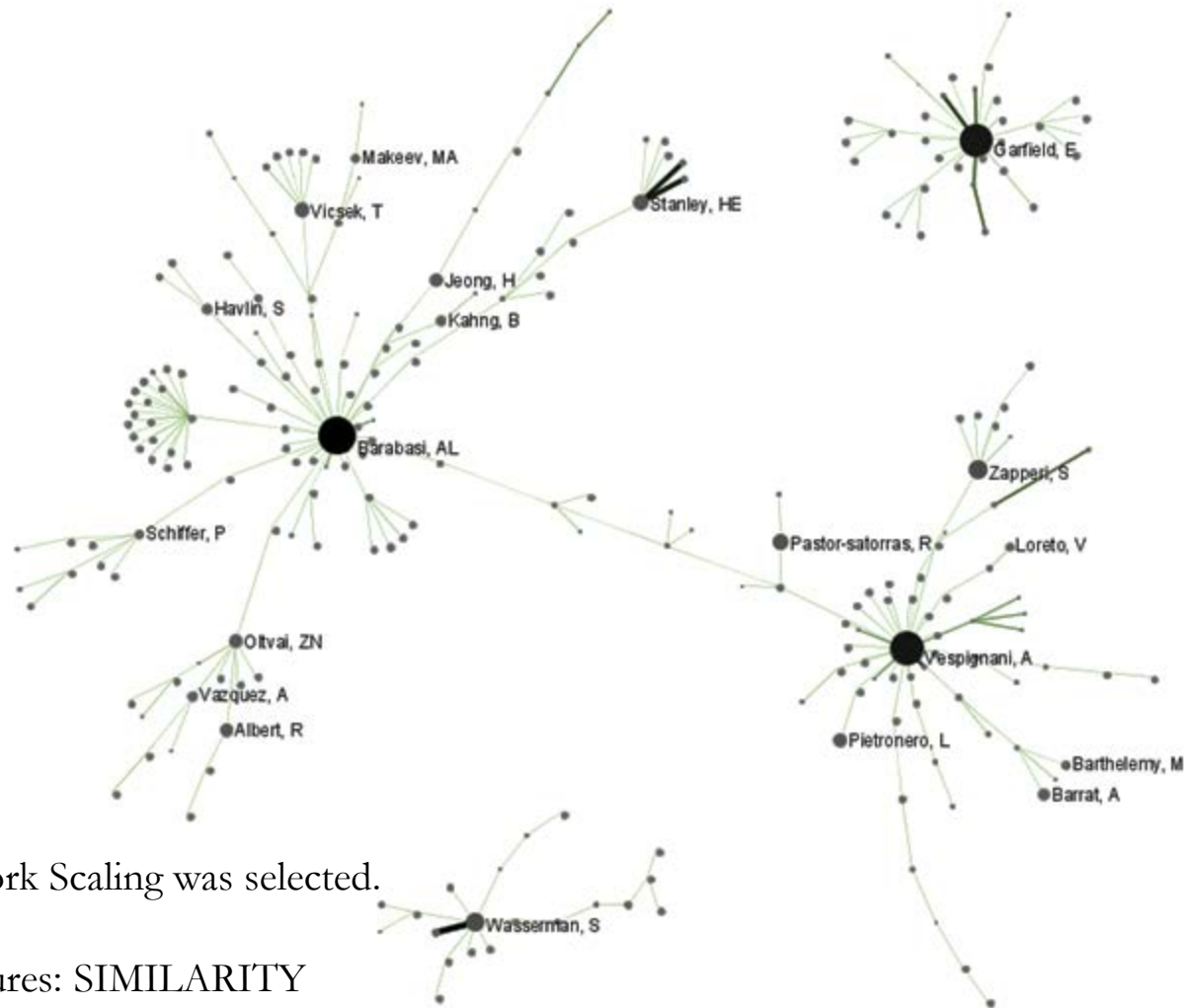
umber of bins: 10

.....





Network Visualization: Reduced Network After Pathfinder Network Scaling



.....

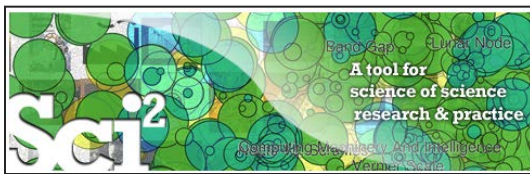
MST-Pathfinder Network Scaling was selected.

Input Parameters:

Weight Attribute measures: SIMILARITY

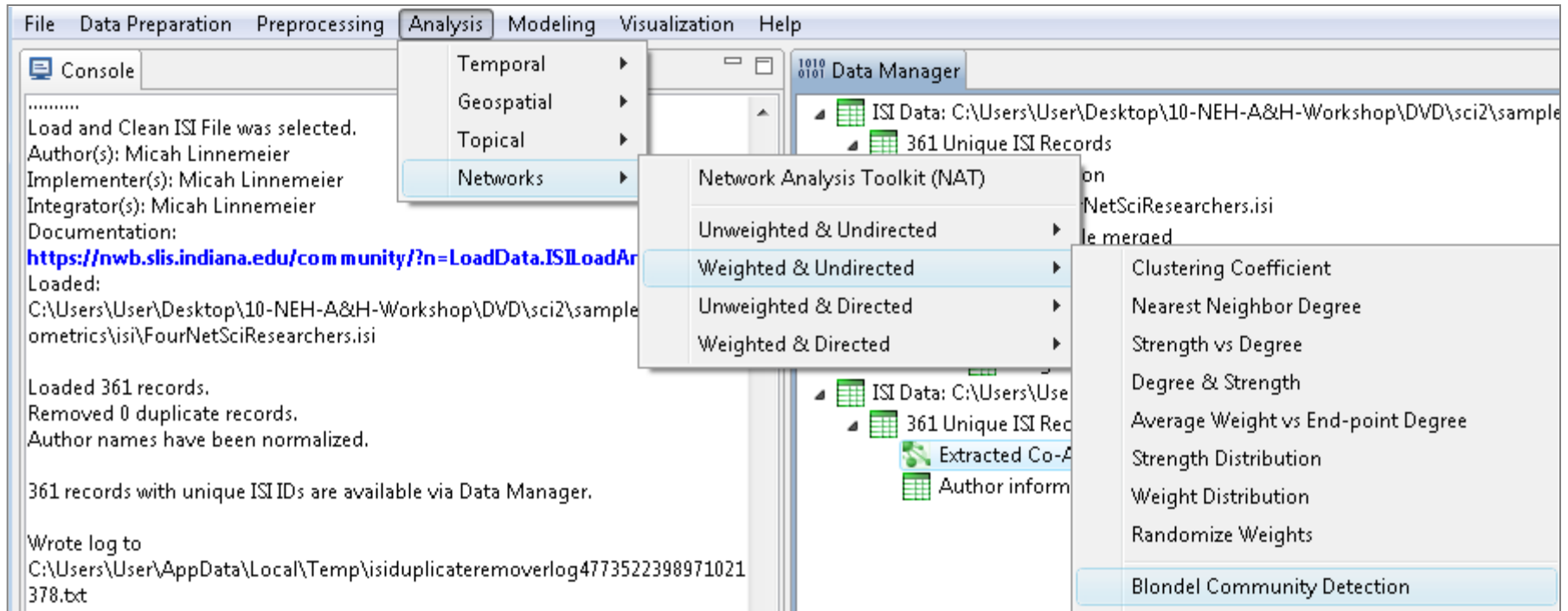
Edge Weight Attribute: weight

.....

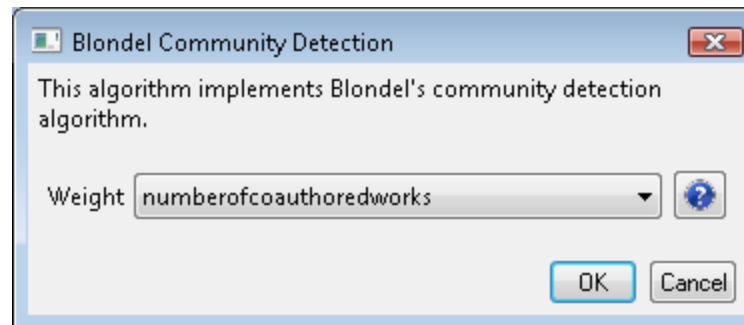


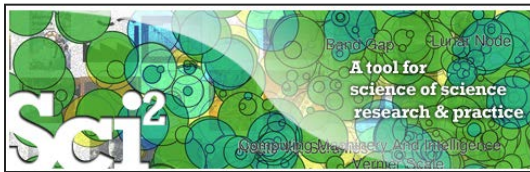
Network Visualization: Circular Hierarchy Visualization

Select Co-Author Network and run Blondel Community detection:



With parameter values





Network Visualization: Circular Hierarchy Visualization

Visualize resulting file using '*Visualization > Networks > Circular Hierarchy*' with parameter values

The image is a screenshot of a software dialog box titled "Circular Hierarchy". The dialog box has a light blue header bar with a close button (X) in the top right corner. Below the header, the text "Provides Circular Hierarchy Visualization on the network." is displayed. The main area of the dialog contains several settings, each with a label, a control element (text box or dropdown), and a help button (question mark icon). The settings are: "Degree of Edge Bundling" with a text box containing "0.75"; "Node Strength Column" with a dropdown menu showing "timescited"; "Level 0" with a dropdown menu showing "blondel_community_level_0"; "Level 1" with a dropdown menu showing "blondel_community_level_1"; "Level 2" with a dropdown menu showing "blondel_community_level_2"; "Level 3" with a dropdown menu showing "No Level"; "Edge Weight Column" with a dropdown menu showing "numberofcoauthoredworks"; "Node Color Column" with a dropdown menu showing "numberofworks"; and "Node Color Range" with a dropdown menu showing "Green to red". At the bottom right of the dialog are "OK" and "Cancel" buttons.

Circular Hierarchy

Provides Circular Hierarchy Visualization on the network.

Degree of Edge Bundling 0.75 ?

Node Strength Column timescited ?

Level 0 blondel_community_level_0 ?

Level 1 blondel_community_level_1 ?

Level 2 blondel_community_level_2 ?

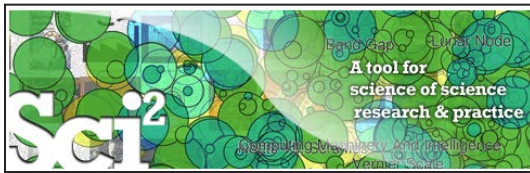
Level 3 No Level ?

Edge Weight Column numberofcoauthoredworks ?

Node Color Column numberofworks ?

Node Color Range Green to red ?

OK Cancel



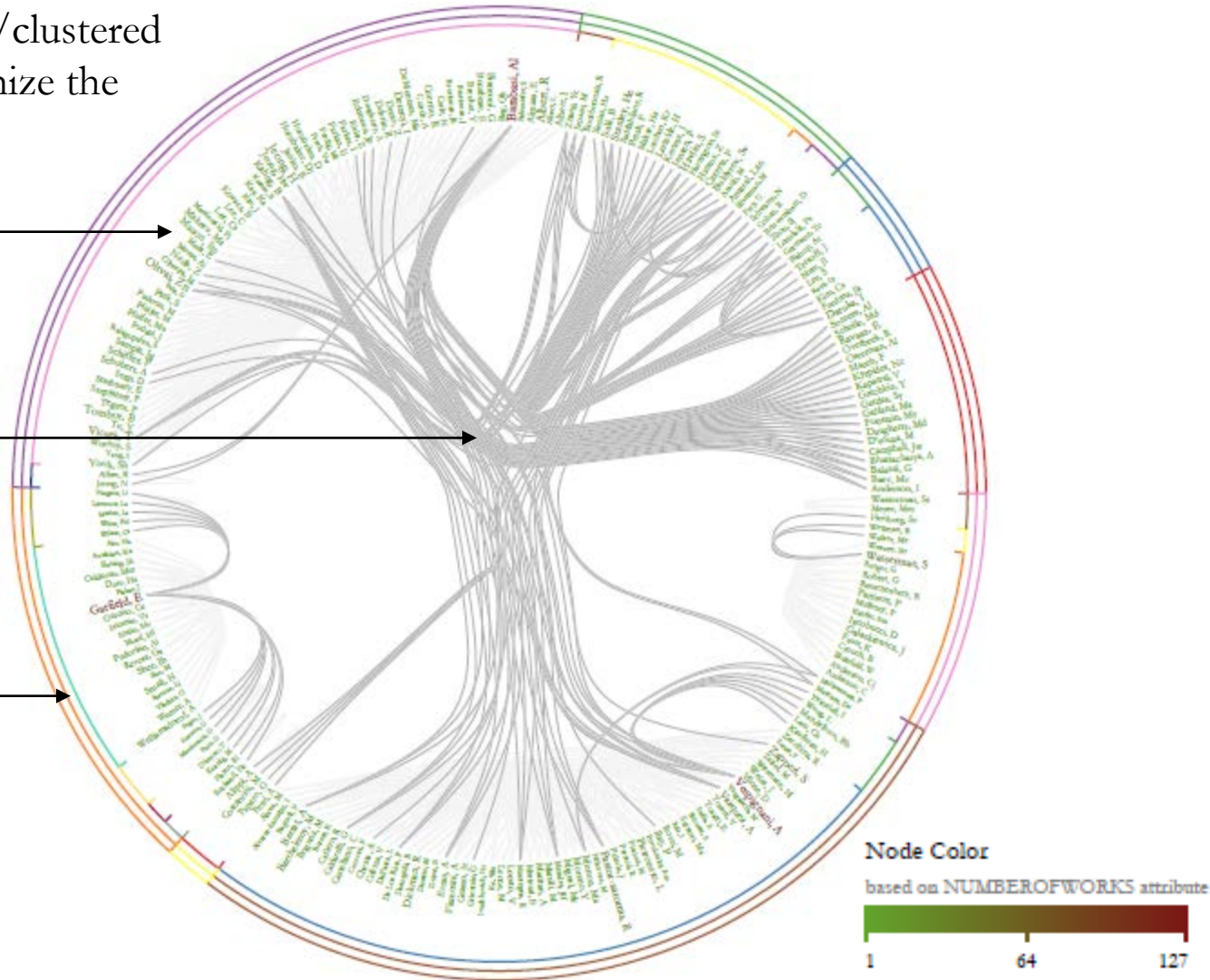
Network Visualization: Circular Hierarchy Visualization

Nodes that are interlinked/clustered are spatially close to minimize the number of edge crossings.

Node labels, e.g.,
author names.

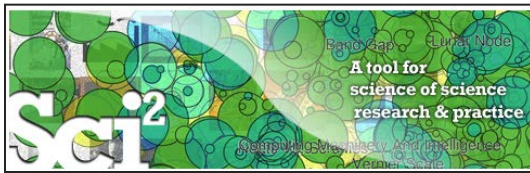
Network structure
using edge bundling.

Color coded cluster
hierarchy according to
Blondel community
detection algorithm.



Note:

Header/footer info, legend, and more meaningful color coding are under development.



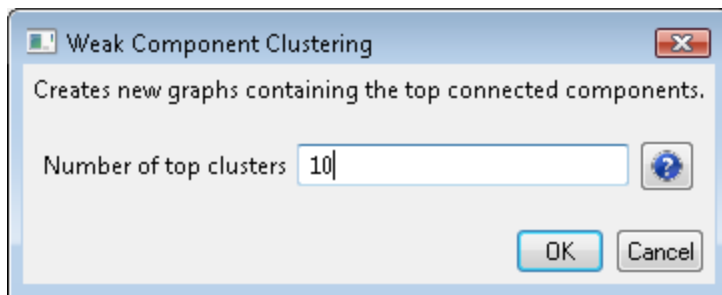
Paper-Citation Network Layout

To extract the paper-citation network, select the '*361 Unique ISI Records*' table and run '*Data Preparation > Extract Paper Citation Network.*'

The result is a unweighted, directed network of papers linked by citations, named *Extracted paper-citation network* in the Data Manager.

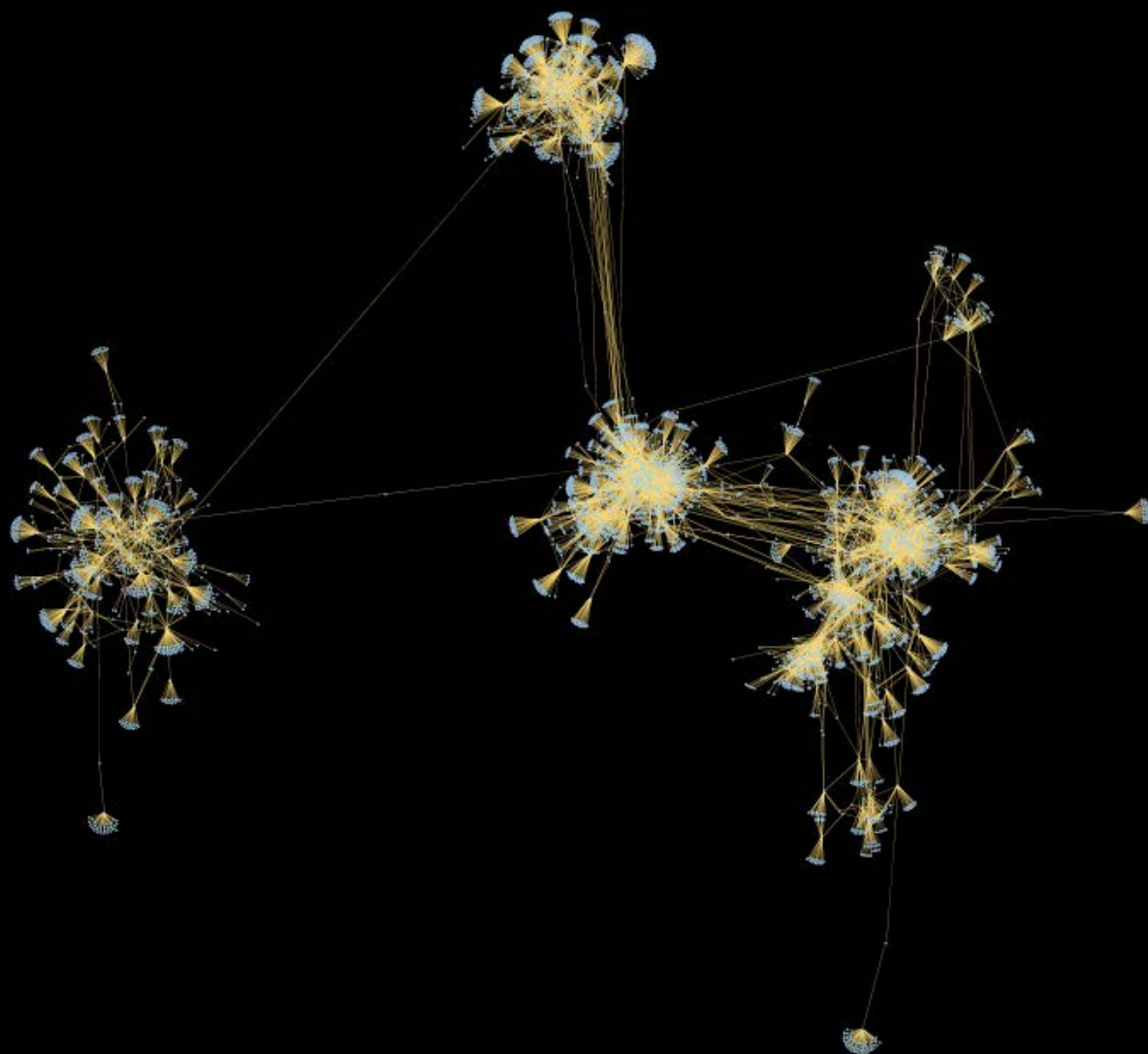
Run *NAT* to calculate that the network has 5,342 nodes and 9,612 edges. There are 15 weakly connected components. (0 isolates)

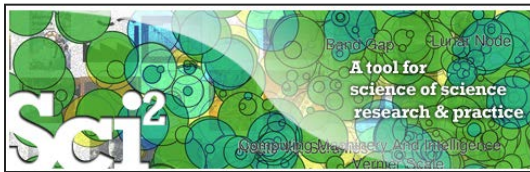
Run '*Analysis > Networks > Unweighted and Directed > Weak Component Clustering*' with parameters



to identify top-10 largest components. The largest (giant) component has 5,151 nodes.

To view the complete network, select the network and run '*Visualization > GUESS*'.



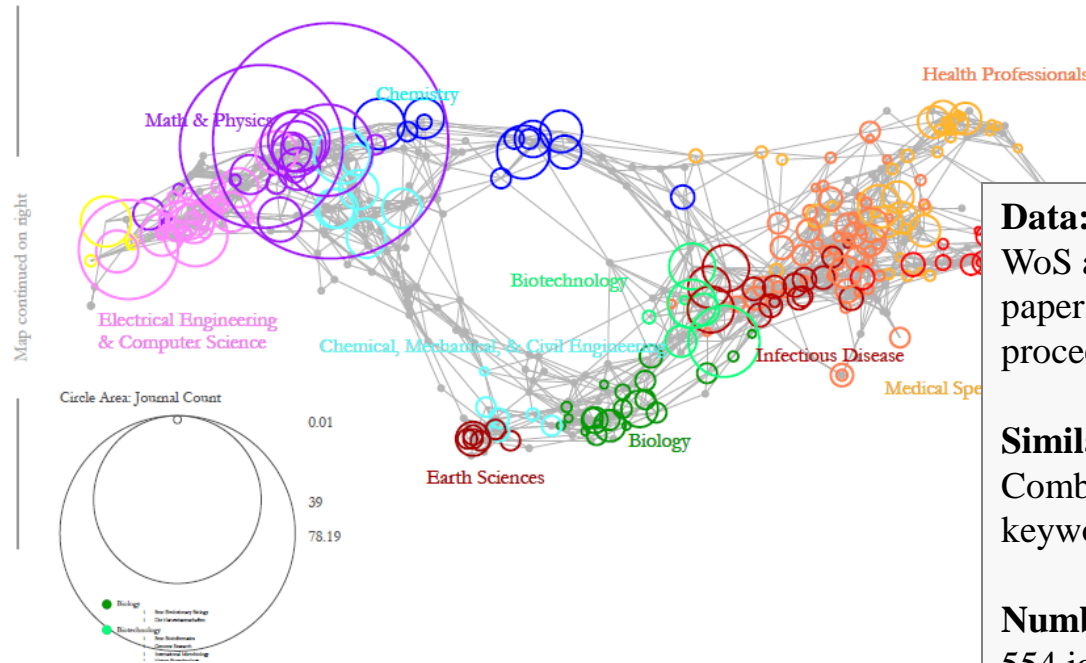


Topic Mapping: UCSD Science Map

Science Map via Journals for FourNetSciResearchers.isi

314 journal references matched out of 361 found.

These 314 references are associated with 13 of 13 disciplines of science and 255 of 554 research specialties in the UCSD Map of Science.



Data:

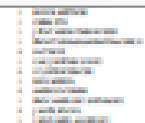
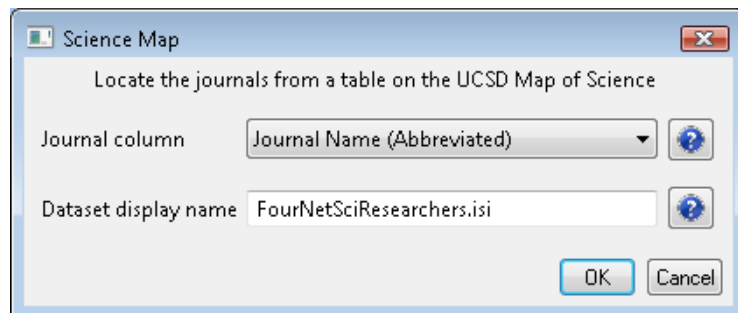
WoS and Scopus for 2001–2005, 7.2 million papers, more than 16,000 separate journals, proceedings, and series

Similarity Metric:

Combination of bibliographic coupling and keyword vectors

Number of Disciplines:

554 journal clusters further aggregated into 13 main scientific disciplines that are labeled and color coded in a metaphorical way, e.g., Medicine is blood red and Earth Sciences are brown as soil.

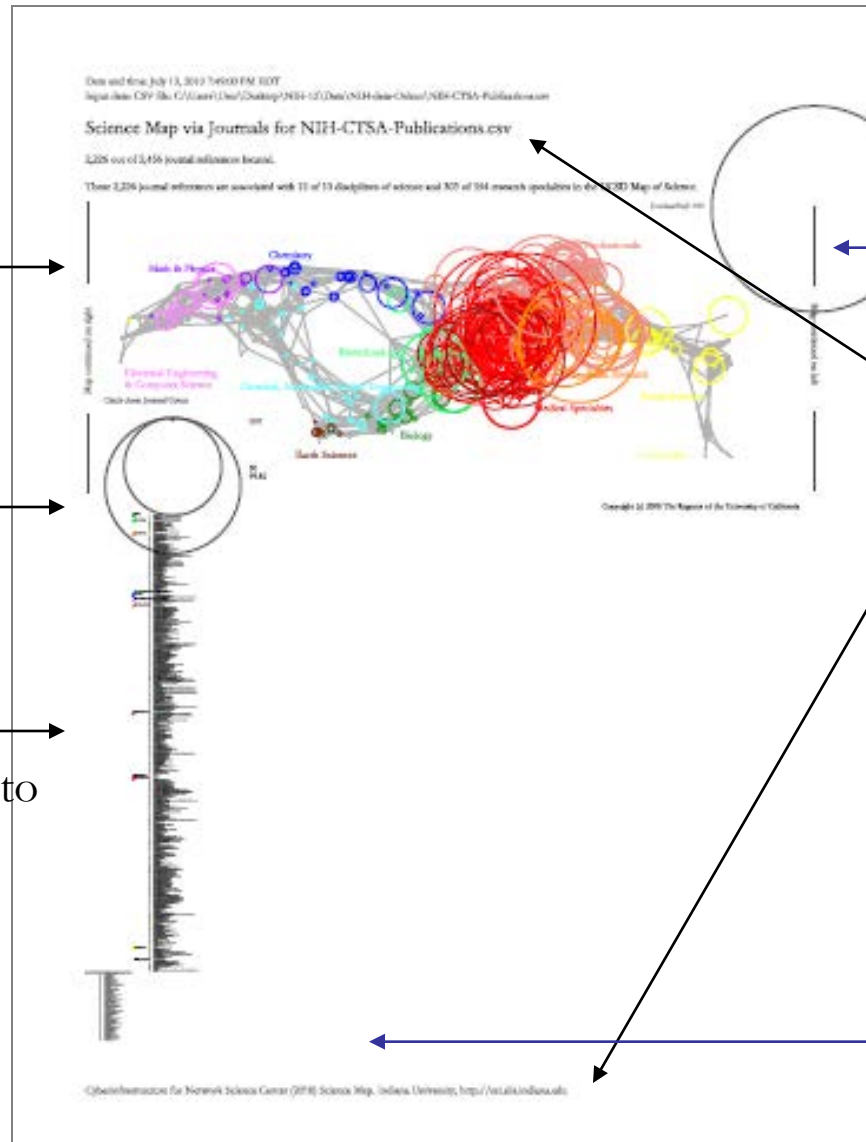


How to Read the UCSD Map

UCSD Science Map
with data overlay.

Map legend of
circle area size
coding

Listing of all data
records organized into
UCSD science areas.



Circle of non-located,
e.g., 'Unclassified'
records.

Header and footer with
information when this
map was created, by
whom and using what
data set.

Listing and circle of
non-located, e.g.,
'Unclassified' records.

Break



Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

➤ Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

➤ Load and clean a dataset as text file; process raw data into networks.

➤ Find basic statistics and run various algorithms over the network.

➤ Visualize as either a circular hierarchy or network

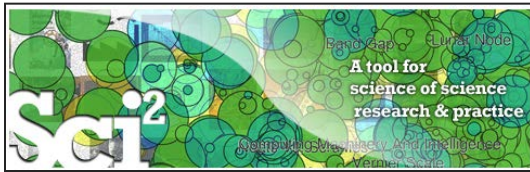
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

4:45 Outlook and Discussion

5:00 Adjourn



Sci2 Demo I: Geospatial maps with congressional districts

	A
1	Zip code
2	90095
3	4672
4	232980568
5	10032
6	10039242
7	46091500
8	191112434
9	27705
10	981959472
11	10065
12	10065



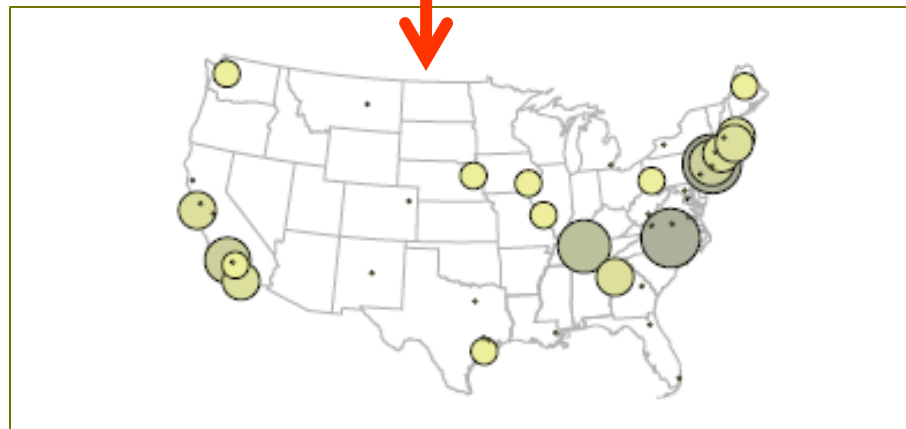
Identify Congressional District, Latitude, Longitude

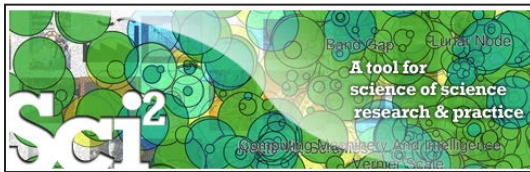
	A	B	C	D
1	Zip code	Congressional District	Latitude	Longitude
2	90095	CA-30	34.0735035	-118.6645815
3	4672	ME-02	45.818717	-69.0290345
4	232980568	VA-03	37.270472	-77.0699835



Aggregate/Count identical Congressional Districts

	A	B	C	D
1	Congressional District	Latitude	Longitude	Count
2	CA-30	34.0735035	-118.6645815	4
3	ME-02	45.818717	-69.0290345	2
4	VA-03	37.270472	-77.0699835	1
5	NY-15	40.8341475	-73.9342095	4





Relevant Sci2 Manual entry

- ☐ Home
- ☐ 1 Introduction
- ☒ 2 Getting Started
- ☒ 3 Algorithms, Tools, and Plugins
- ☒ 4 Workflow Design
- ☒ 5 Sample Workflows
 - ☒ 5.1 Individual Level Studies - Micro
 - ☒ 5.2 Institution Level Studies - Meso
 - ☒ 5.3 Global Level Studies - Macro
 - ☐ 5.3.1 Geo USPTO (SDB Data)
 - ☐ 5.3.2 Congressional District Geocoder
- ☒ 6 Sample Science Studies & Online Services
- ☒ 7 Extending the Sci2 Tool
- ☒ 8 Relevant Datasets and Tools
- ☐ 9 References
- ☐ Appendix 1 Glossary
- ☐ Appendix 2 CShell Algorithms
- ☐ Appendix 3 Sci2 Release Notes v0.5 alpha



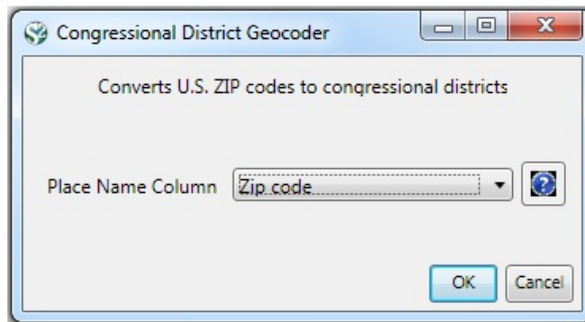
5.3.2 Congressional District Geocoder

14 Added by Scott Weingart, last edited by Ted Polley on Mar 28, 2011 (view change)

Tools ▾

zip code.csv	
Region(s):	United States
Analysis Type(s):	Geospatial Analysis

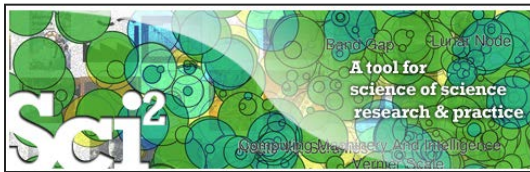
To visualize Congressional Districts you must first extract that data from a dataset containing either ZIP codes or addresses. You can download the Congressional District Geocoder plugin [here](#). You can load any file that contains 9-digit U.S. ZIP codes to be geocoded. A sample file can be loaded by using 'File > Load' and following this path: '[yoursic2directory/sampledata/geo/zipcode.csv](#)'. Load the file in Standard csv format. Then select the file in the data manager and use 'Analysis > Geospatial > [Congressional District Geocoder](#)' with the following parameters:



5-digits ZIP codes with multiple congressional districts, empty entries and invalid ZIP codes that failed to be geocoded will list in warning messages on the console. The output table contains all columns of the input table with three additional columns appended: Congressional district, latitude, and longitude. To view the output table save the file using 'File > Save...' and selecting the desired save location (to view the file in Excel save it as a csv file). Once the file has been saved it can be viewed with your choice of program. Below the file has been opened as a [csv file](#):

	A	B	C	D
1	Zip code	Congressional District	Latitude	Longitude
2	90095	CA-30	34.0735035	-118.6645815

<http://sci2.wiki.cns.iu.edu/5.3.2+Congressional+District+Geocoder>



Relevant CShell plugin



Congressional District Geocoder



1 Added by [Ted Polley](#), last edited by [Chin Hua Kong](#) on Mar 29, 2011 ([view change](#))

Description

This algorithm converts the given **9-digits U.S. ZIP codes (ZIP+4 codes)** into its congressional districts and geographical coordinates (latitude and longitude). The Benchmark is 50,000 ZIP codes per second. Download the plugin [here](#).

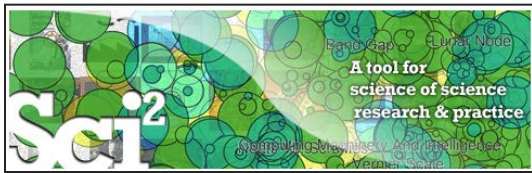
Pros & Cons

1. The algorithm is using a local database mapping with 25MB file size. It will increase the application size dramatically. So it is build as an external plugin
2. For first execution in the same application window, the plugin required 5 seconds to load the database. The consequent execution will not required the pre-loading phase.
3. Since some 5-digits ZIP codes contain multiple districts, the 9-digits ZIP codes is required for the conversion. Warning message will be printed to notice user if the given 5-digits ZIP codes contain multiple districts
4. Congressional district might be varied by each election. The database would need to be maintained and updated relatively.

Applications

This plugin only support U.S. ZIP codes. It convert 9-digits ZIP codes to their belonging congressional district. It is an external plugin since the data size is so large. The dataset is based on the year 2008 election.

<http://cishell.wiki.cns.iu.edu/Congressional+District+Geocoder>



Console Messages

Load... was selected.

Documentation: <http://wiki.cns.iu.edu/display/CISHELL/Data+Formats>

Loaded: C:\Users\katy\Desktop\NWB-SCI2\sci2-2011.04.04-v0.5a\sampladata\geo\zip code.csv

.....

Congressional District Geocoder was selected.

Implementer(s): Chin Hua Kong

Integrator(s): Chin Hua Kong

Documentation: <https://nwb.cns.iu.edu/community/?n=SampleData.CongressionalDistrictGeocoder>

Input Parameters:

Place Name Column: Zip code

District values added to Congressional District, Latitude and Longitude respectively.

There are 2 rows with "33612" ZIP code, which could not been given a congressional district.

There are 1 rows with "2472" ZIP code, which could not been given a congressional district.

There are 3 rows with "10016" ZIP code, which could not been given a congressional district.

There are 1 rows with "11203" ZIP code, which could not been given a congressional district.

There are 1 rows with "60637" ZIP code, which could not been given a congressional district.

There are 1 rows with "70118" ZIP code, which could not been given a congressional district.

There are 1 rows with "60612" ZIP code, which could not been given a congressional district.

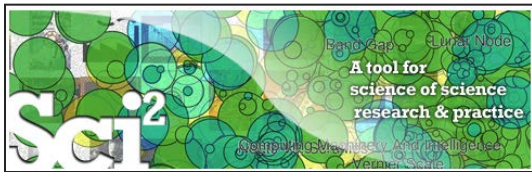
There are 3 rows with "21205" ZIP code, which could not been given a congressional district.

There are 1 rows with "2467" ZIP code, which could not been given a congressional district.

5-digit ZIP codes may often be insufficient, as many zip codes contain multiple congressional districts. 9-digit zip codes may be required. If a zip code was recently created, it may also not be contained in our database.

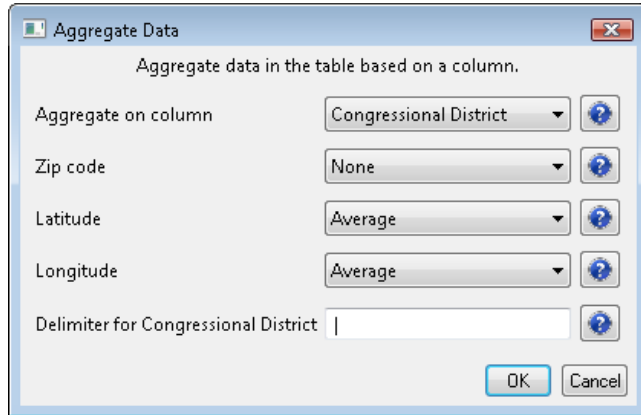
Successfully converted 86 out of 100 ZIP codes to congressional districts.

.....



Sci2 Demo I: Geospatial maps with congressional districts

Run *Preprocessing > General > Aggregate Data*
using parameter values



Note: Need lat/long for geomap.

Input Parameters:

Aggregate on column: Congressional District

Longitude: AVERAGE

Latitude: AVERAGE

Delimiter for Congressional District: |

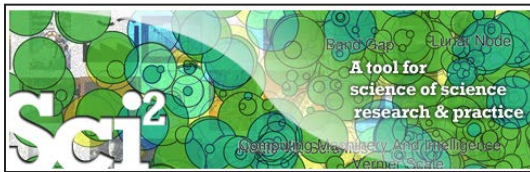
Zip code: NONE

Aggregated by ": All rows of Latitude column were skipped due to no non-null, non-empty values.

Aggregated by ": All rows of Longitude column were skipped due to no non-null, non-empty values.

Frequency of unique "Congressional District" values added to "Count" column.

"Zip code" column has been deleted from the output. Since No aggregation was mentioned for it.



Create Geo Map (Circle Annotation)

.....

Geo Map (Circle Annotations) was selected.

Author(s): Joseph R. Biberstine

Implementer(s): Joseph R. Biberstine

Integrator(s): Joseph R. Biberstine

Documentation: <http://wiki.cns.iu.edu/display/CISHELL/Geo+Map>

Input Parameters:

Longitude: Longitude

Size Circles By: CircleSize

Color Circle Exteriors By: None (no outer color)

Color Circle Interiors By: CircleSize

Exterior Color Scaling: Linear

Exterior Color Range: Yellow to Blue

Interior Color Range: Blue to Red

Size Scaling: Linear

Map: US States

Author Name:

Interior Color Scaling: Linear

Latitude: Latitude

14 rows in the table did not specify all values needed to make a circle; those rows were skipped.

Printing PostScript..

Done.

Saved: C:\Users\katy\Desktop\geoMaps2903082942930990749.ps

Save ps file, convert to pdf, view.



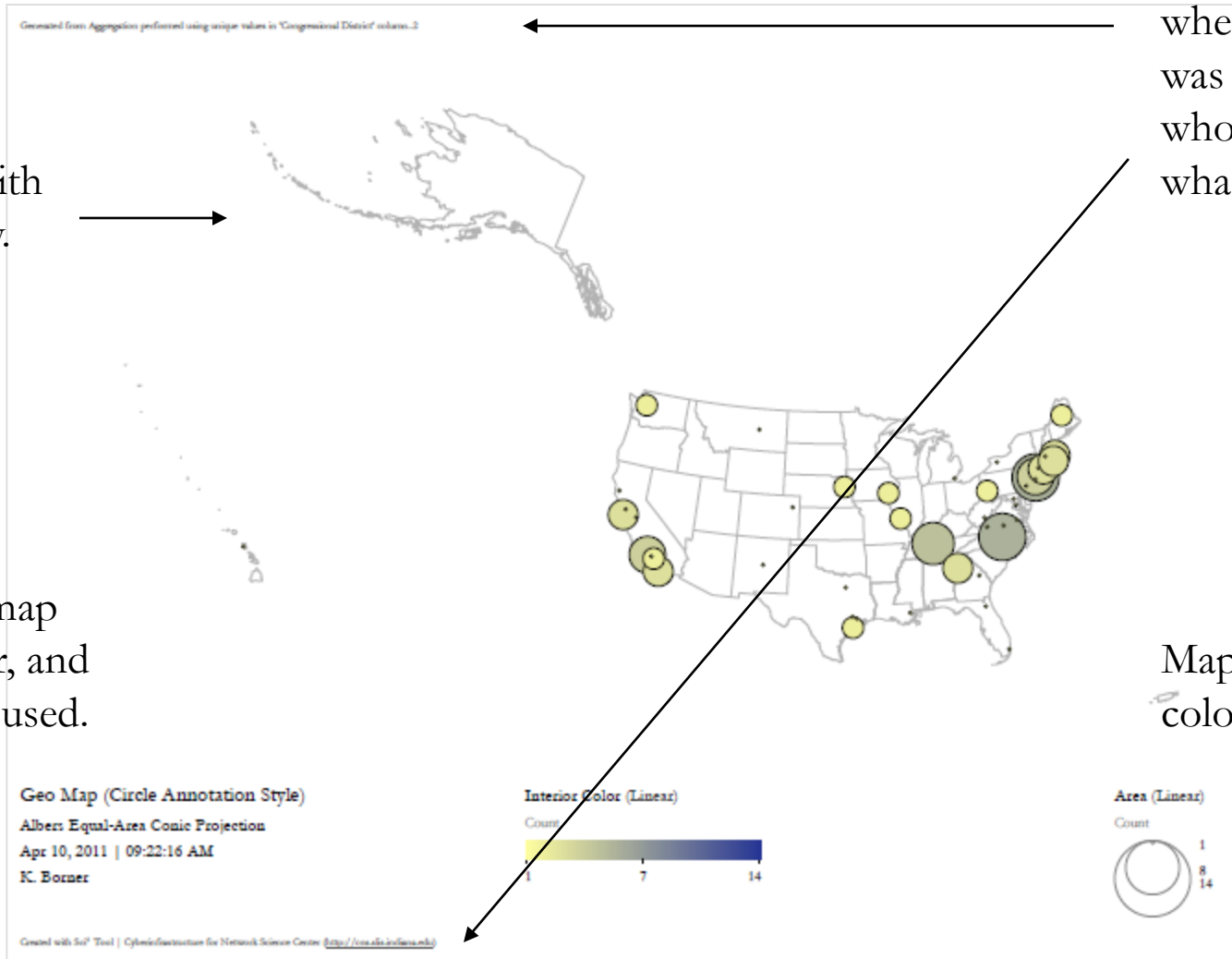
How to Read the Geo Map

Header and footer with information when this map was created, by whom and using what data set.

U.S. Map with data overlay.

Listing of map type, author, and parameters used.

Map legend with color coding.





Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

➤ Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

2:30 Sci2 Sample Workflow: Studying Four Major NetSci Researchers.

➤ Load and clean a dataset as text file; process raw data into networks.

➤ Find basic statistics and run various algorithms over the network.

➤ Visualize as either a circular hierarchy or network

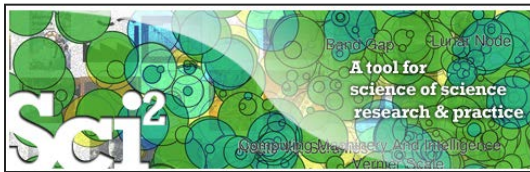
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

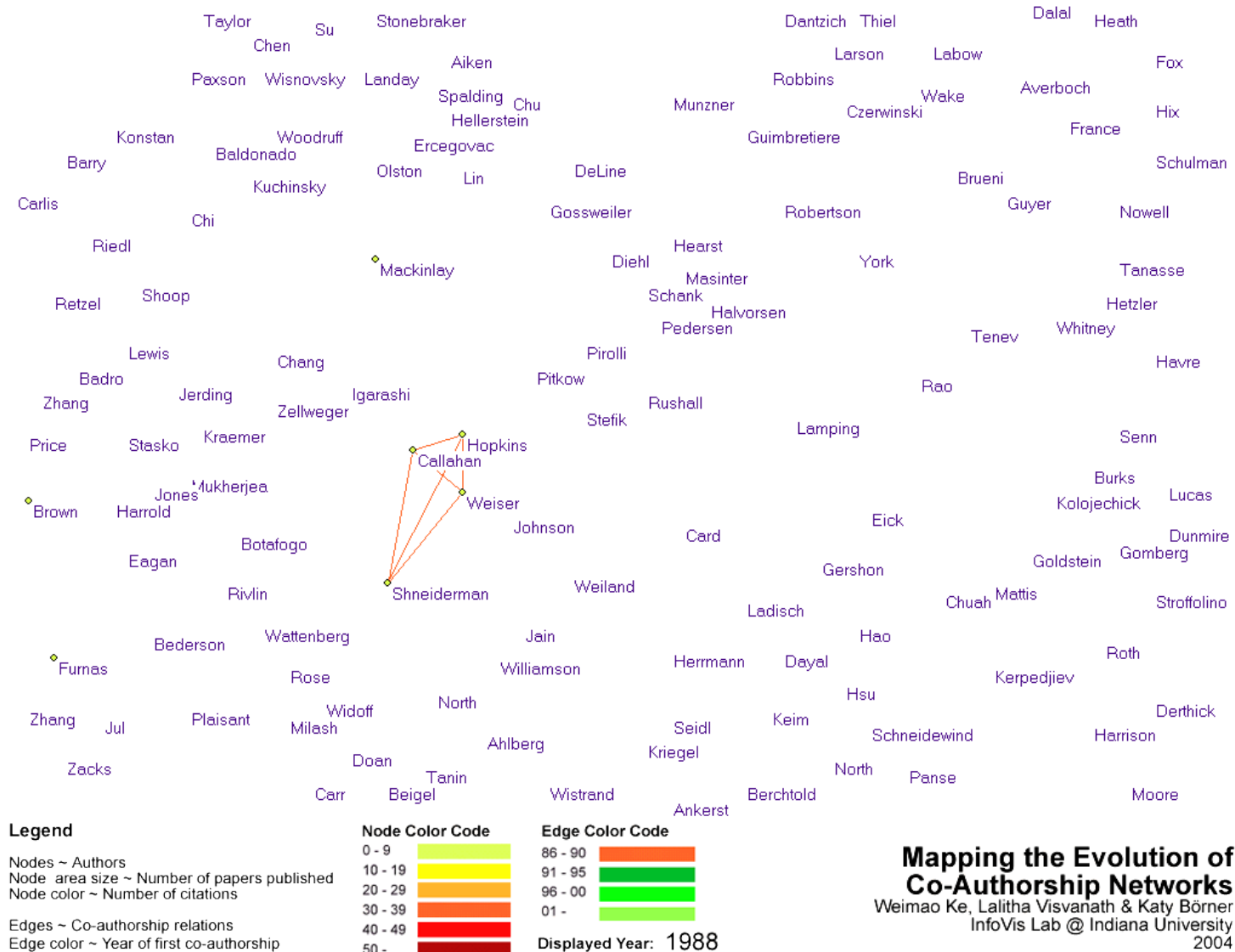
4:30 Sci2 Demo II: Evolving collaboration networks

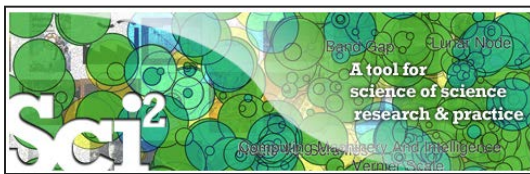
4:45 Outlook and Discussion

5:00 Adjourn



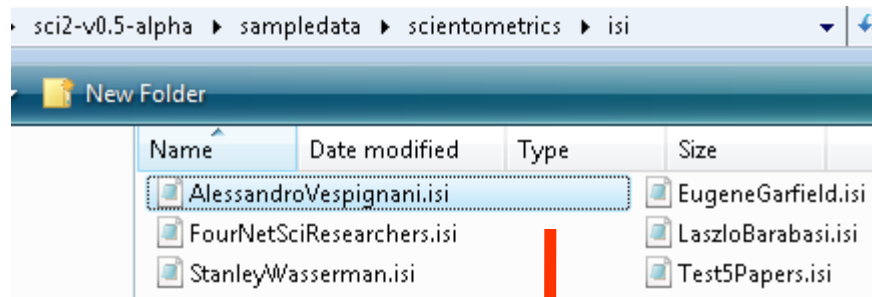
Sci2 Demo II: Evolving collaboration networks





Sci2 Demo II: Evolving collaboration networks

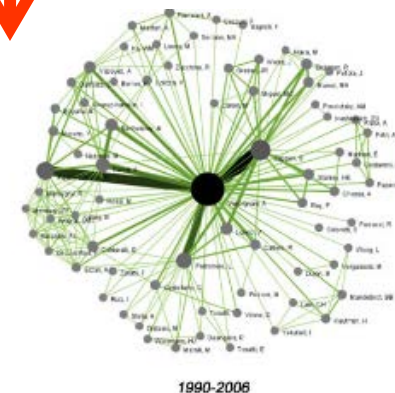
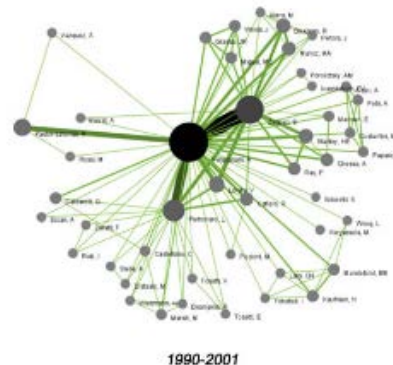
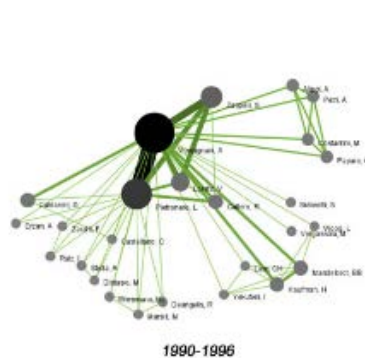
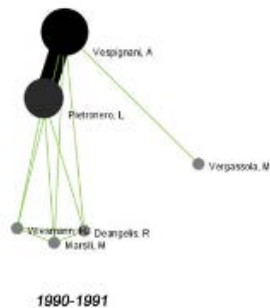
Load isi formatted file

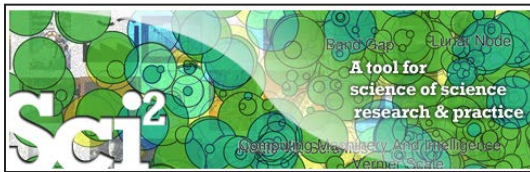


As csv, file looks like:

	A	B	C	D	E	F	G
1	Abstract	Authors	Authors (Full Names)	Beginning	Book Series	Book Series	Cited Page
2	The systematic study of	Colizza, V Barrat, A Barthelemy, M Vespignani, A		2015			
3	Uncovering the hidden re	Colizza, V Flammini, A Serrano, MA Vespignani, A		110			
4	Computer viruses can s	Vespignani, A		135			
5	Mapping the Internet ge	Dall'Asta, L Alvarez-Hamelin, I Barrat, A Vazquez, A Vespignani, A		140		LECTURE NOTES IN	

Visualize each time slide separately:





Relevant Sci2 Manual entry

- ☐ Home
- ☐ 1 Introduction
- ☒ 2 Getting Started
- ☒ 3 Algorithms, Tools, and Plugins
- ☒ 4 Workflow Design
- ☒ 5 Sample Workflows
 - ☒ 5.1 Individual Level Studies - Micro
 - ☐ 5.1.1 Mapping Collaboration, Publication, and Funding Profiles of One Researcher (EndNote and NSF Data)
 - ☒ 5.1.2 Time Slicing of Co-Authorship Networks (ISI Data)
 - ☐ 5.1.3 Funding Profiles of Three Researchers at Indiana University (NSF Data)
 - ☐ 5.1.4 Studying Four Major NetSci Researchers (ISI Data)
 - ☒ 5.2 Institution Level Studies - Meso
 - ☒ 5.3 Global Level Studies - Macro
- ☒ 6 Sample Science Studies & Online Services
- ☒ 7 Extending the Sci2 Tool
- ☒ 8 Relevant Datasets and Tools
- ☐ 9 References



5.1.2 Time Slicing of Co-Authorship Networks (ISI Data)

Tools ▾

6 Added by Ted Polley, last edited by Scott Weingart on Mar 16, 2011 (view change)

AlessandroVespignani.isi	
Time frame:	1990-2006
Region(s):	Indiana University, University of Rome, Yale University, Leiden University, International Center for Theoretical Physics, University of Paris-Sud
Topical Area(s):	Informatics, Complex Network Science and System Research, Physics, Statistics, Epidemics
Analysis Type(s):	Co-Authorship Network

The Sci2 Tool supports the analysis of evolving networks. For this study, load Alessandro Vespignani's publication history from ISI, which can be downloaded from Thomson's Web of Science or loaded using 'File > Load' and following this path: '**yoursci2directory**/sampledata/scientometrics/isi/AlessandroVespignani.isi' using.' Slice the data into five year intervals from 1990-2006 using 'Preprocessing > Temporal > **Slice Table by Time**' and the following parameters:

Slice Table by Time

Slice a table into groups of rows by time.

Date/Time Column
Publication Year

Date/Time Format
yyyy

Slice Into
Years

How Many?
5

From Time
1990

To Time
2006

☒ Cumulative?

☐ Align With Calendar

Week Starts On
Sunday

[http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+\(ISI+Data\)](http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+(ISI+Data))

Slice Table by Time

Slice Table by Time

Slice a table into groups of rows by time.

Date/Time Column: **Publication Year**

Date/Time Format: **yyyy**

Slice Into: **Years**

How Many?: **5**

From Time: **1990**

To Time: **2006**

☒ Cumulative?

☐ Align With Calendar

Week Starts On: **Sunday**

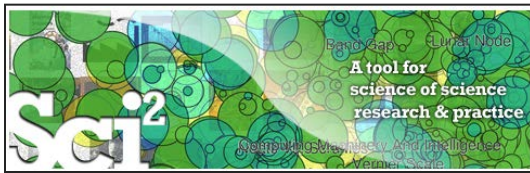
"Slice Into" allows the user to slice the table by days, weeks, months, quarters, years, decades, and centuries. There are two additional parameters for time slicing: cumulative and align with calendar. The former produces tables containing all data from the beginning to the end of each table's time interval, which can be seen in the Data Manager and below:



The latter option aligns the output tables according to calendar intervals:



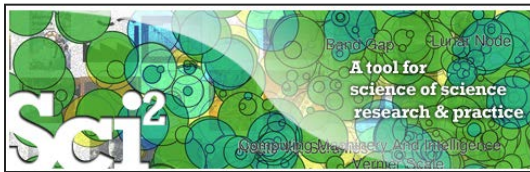
Choosing "Years" under "Slice Into" creates multiple tables beginning from January 1st of the first year. If "Months" is chosen, it will start from the first day of the earliest month in the chosen time interval.



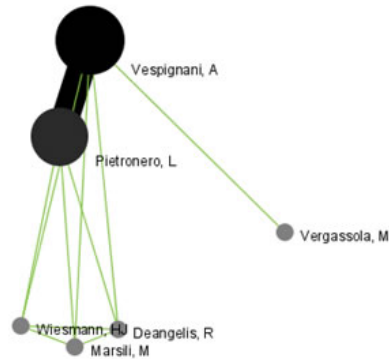
Visualize Each Network, Keep Node Positions

1. To see the evolution of Vespignani's co-authorship network over time, check '*cumulative*'.
2. Extract co-authorship networks one at a time for each sliced time table using '*Data Preparation > Extract Co-Author Network*', making sure to select "ISI" from the pop-up window during the extraction.
3. To view each of the Co-Authorship Networks over time using the same graph layout, begin by clicking on longest slice network (the '*Extracted Co-Authorship Network*' under '*slice from beginning of 1990 to end of 2006 (101 records)*') in the data manager. Visualize it in GUESS using '*Visualization > Networks > GUESS*'.
4. From here, run '*Layout > GEM*' followed by '*Layout > Bin Pack*'. Run '*Script > Run Script ...*' and select '*yoursci2directory/scripts/GUESS/co-author-nw.py*'.
5. In order to save the x, y coordinates of each node and to apply them to the other time slices in GUESS, select '*File > Export Node Positions*' and save the result as '*yoursci2directory/NodePositions.csv*'. Load the remaining three networks in GUESS using the steps described above and for each network visualization, run '*File > Import Node Positions*' and open '*yoursci2directory/NodePositions.csv*'.
6. To match the resulting networks stylistically with the original visualization, run '*Script > Run Script ...*' and select '*yoursci2directory/scripts/GUESS/co-author-nw.py*', followed by '*Layout > Bin Pack*', for each.

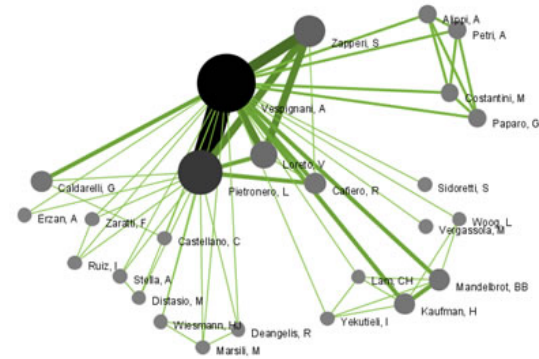
[http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+\(ISI+Data\)](http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+(ISI+Data))



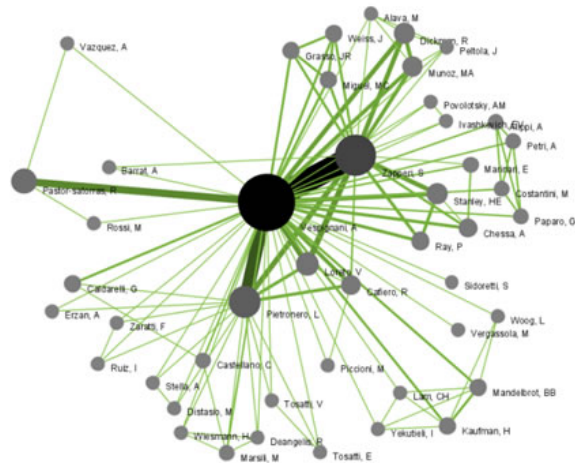
Visualize Each Network, Keep Node Positions



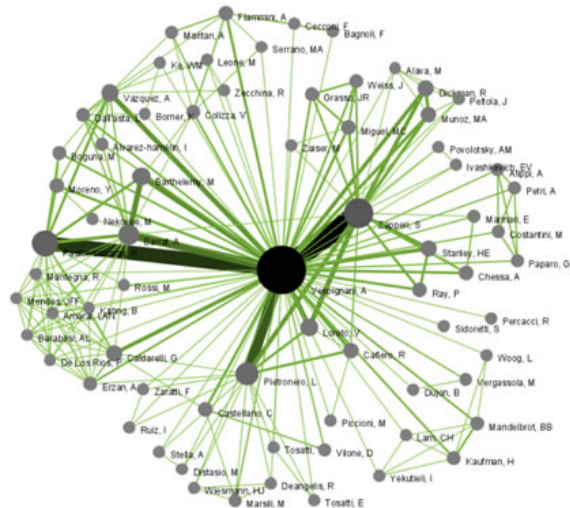
1990-1991



1990-1996

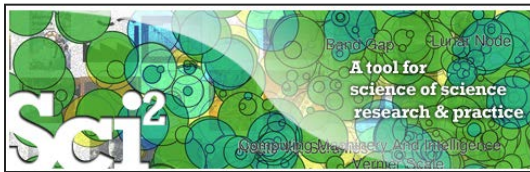


1990-2001



1990-2006

[http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+\(ISI+Data\)](http://sci2.wiki.cns.iu.edu/5.1.2+Time+Slicing+of+Co-Authorship+Networks+(ISI+Data))



Relevant CShell plugin



Added by [Aretha Alencar](#), last edited by [Ted Polley](#) on Jan 12, 2011 ([view change](#))

Description

Slice Table By Time is an algorithm to chop a table up into new tables, based on a date/time column. It takes the column with the date/time data, a string describing the format of that column, the intervals that the data should be sliced into, whether or not the slices are cumulative, whether or not the slices should be aligned with the calendar, and what day the week is considered to start on (which only matters if the slices are aligned with the calendar) as parameters.

The column to use for date/time values should have a single value for each row of data. It is used by the algorithm to choose which slice(s) the row should end up in. In order to determine what date/time is represented by that row, you must provide the algorithm with a descriptive format, in the second parameter. For instance, a four digit year would be represented by yyyy (the default value). See <http://joda-time.sourceforge.net/api-release/org/joda/time/format/DateTimeFormat.html> for details of all the various formatting options.

The next dropdown has the available intervals to slice the table into. These include milliseconds, seconds, minutes, hours, days, weeks, fortnights, months, quarters, years, decades, and centuries. A future version of the algorithm may include the ability to select how many of these intervals should be grouped together at once.

The checkbox that follows determines if the slices will be cumulative. If the slices are not cumulative, every row in the original table is in one and only one resulting slice. However, if the slices are cumulative, every row in the original table is in the slice it is for and every slice for a period after that.

The checkbox that follows determines if the slices will be aligned with the calendar. For instance, if the first row is for June 7th, 2006 and yearly slices are chosen, then the default behavior will be to have the first slice be from June 7th, 2006 to June 6th, 2007. However, if the slices are aligned with the calendar, the first slice will be from January 1st, 2006 to December 31st, 2006. Alignment does not affect the output for intervals of fortnights, quarters, decades, or milliseconds.

If the slices are aligned with the calendar and are weekly, then the day the week starts is used to determine how they are aligned.

Pros & Cons

The output of the slice algorithm is in separate tables, so a longitudinal analysis will require working with each slice separately, which can be awkward. There will likely be future versions of the time slice algorithm that annotate the original table with the slice the rows belong to.

Applications

When doing longitudinal analysis of data, it can be useful to consider it in chunks, such as to calculate how statistics have changed over time. Alternatively, only a particular time period might be of interest, and this algorithm can extract it from data for a larger time range.

Implementation Details

This algorithm uses the Joda Time library extensively, which provides significantly improved capabilities compared to the default Java algorithms for dates and times.

<http://cishell.wiki.cns.iu.edu/Slice+Table+by+Time>



Workshop Overview

1:15 Marcoscope Design and Usage & CShell Powered Tools: NWB & Sci2

1:45 Sci2 Tool Basics

➤ Download and run the tool.

2:00 Sci2 Sample Workflow: Padgett's Florentine Families - Prepare, load, analyze, and visualize family and business networks from 15th century Florence.

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➤ Load and clean a dataset as text file; process raw data into networks.

➤ Find basic statistics and run various algorithms over the network.

➤ Visualize as either a circular hierarchy or network

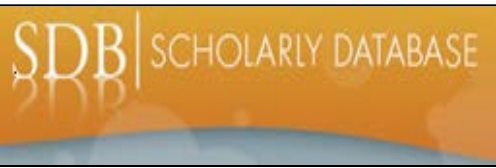
3:30 Break

4:00 Sci2 Demo I: Geospatial maps with congressional districts

4:30 Sci2 Demo II: Evolving collaboration networks

4:45 Marcoscopes: Outlook and Discussion


5:00 Adjourn



Scholarly Database at Indiana University

<http://sdb.wiki.cns.iu.edu>

Supports federated search of 25 million publication, patent, grant records.
Results can be downloaded as data dump and (evolving) co-author, paper-citation networks.



SCHOLARLY DATABASE

Cyberinfrastructure for Network Science Center, SLIS, Indiana University, Bloomington

IU User

IU Users must login using the Central Authentication Service (CAS), the standard IU authentication system. Please click the button below to proceed to the IU login page.

[Go to IU Login](#)

Non-IU User

Email

Password

[Login](#)

Not Registered Yet?

[Register as an IU User](#)

[Register as a Non IU User](#)

In the News

Whitfield, John. 2008. *Group Theory*. Nature, 455, 9: 720-723.


Please Cite As

La Rove, Gavin, Ambre, Sumeet, Burgoon, John, Ke, Weimao and Börner, Katy. (2007) The Scholarly Database and Its Utility for Scientometrics Research. In Proceedings of the 11th International Conference on Scientometrics and Informetrics, Madrid, Spain, June 25-27, 2007, pp. 457-462.
<http://ella.slis.indiana.edu/~katy/paper/07-issi-sdb.pdf>



Acknowledgements

The Scholarly Database is funded by the School of Library and Information Science and the Cyberinfrastructure for Network Science center at Indiana University, the National Science Foundation under Grants No. IIS-0238261 and IIS-0513650, and a James S. McDonnell Foundation grant in area Studying Complex Systems.


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Search

Creators:

Title:

Abstract:

Full Text:

First Year:

Last Year:

☒ **Medline (1898 - 2008)**

☒ **NIH (1961 - 2002)**

☒ **NSF (1985 - 2004)**

☒ **USPTO (1976 - 2007)**

[Search](#)

If multiple terms are entered in a field, they are automatically combined using 'OR'. So, 'breast cancer' matches any record with 'breast' or 'cancer' in that field.

You can put AND between terms to combine with 'AND'. Thus 'breast AND cancer' would only match records that contain both terms.

Double quotation can be used to match compound terms, e.g., "breast cancer" retrieves records with the phrase 'breast cancer', and not records where 'breast' and 'cancer' are both present, but not the exact phrase.

The importance of a particular term in a query can be increased by putting a ^ and a number after the term. For instance, 'breast cancer^10' would increase the importance of matching the term 'cancer' by ten compared to matching the term 'breast'.


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Scholarly Database :: Results - Mozilla Firefox

File Edit View History Bookmarks Tools Help

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Browse Results

Your search returned 13,231 results in 0.295 seconds.

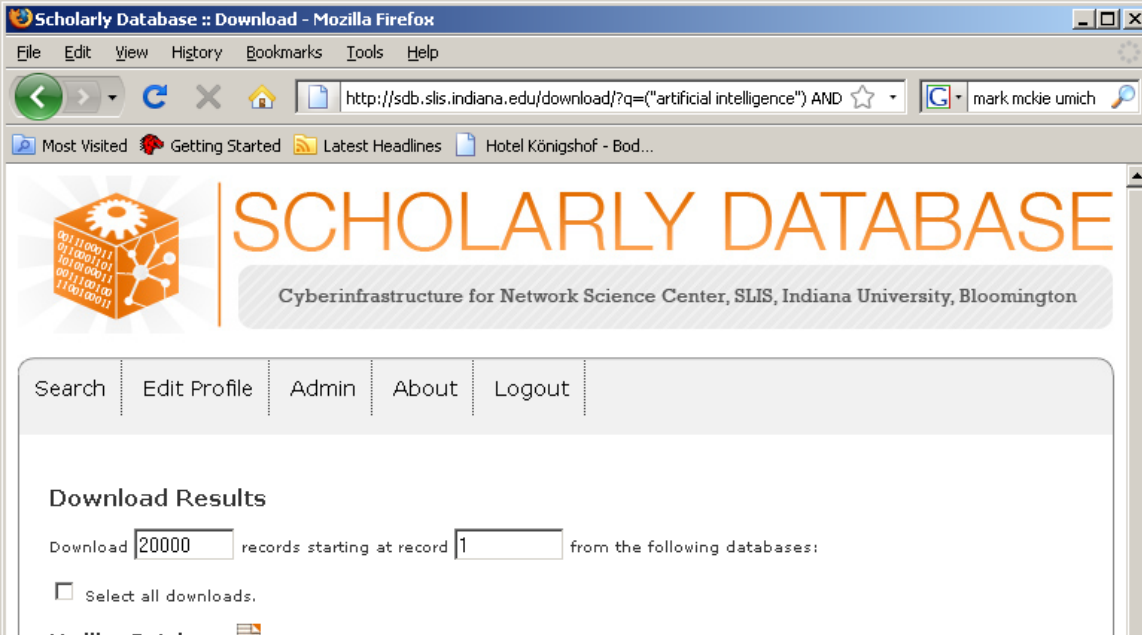
[📄 Download](#)

Total results per database: NIH: 2,103, Medline: 10,235, USPTO: 279, NSF: 614.

Results 1 through 20.

[Next>>](#)

Source	Authors/Creators	Year	Title	Score (out of 5.71)
Medline	LaCombe	1987	Artificial intelligence.	5.71
Medline		1989	Artificial intelligence: expert systems.	5.71
Medline	Schmitt	1990	[Artificial intelligence in dentistry]	5.71
Medline	Adlassnig and Adlassnig	2002	Artificial-intelligence-augmented systems.	5.60
Medline	Touretzky	1980	Artificial intelligence.	4.86
Medline	Goldenberg	1980	Artificial intelligence.	4.86

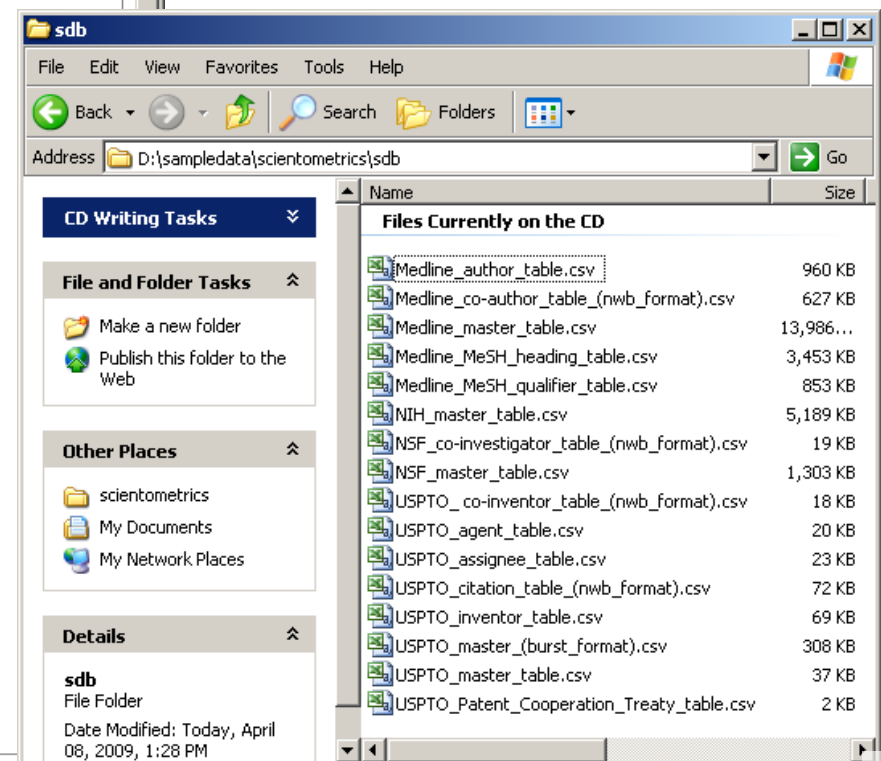


Since March 2009:

Users can download networks:

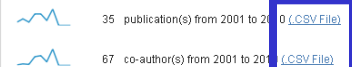
- Co-author
- Co-investigator
- Co-inventor
- Patent citation

and tables for
burst analysis in NWB.



This information is based solely on publications which have been loaded into the VIVO system. This may only be a small sample of the person's total work.

General Statistics

Co-Author Network ([GraphML File](#))

Legend



Interact

Hover over any name to see the number of joint publications and co-authors with Borner, Katy. Click on a name to see details on the right.

Thresholding

Only people that co-authored more than 1 paper(s) with Borner, Katy are shown.

Sorted into communities: Co-authors are placed near one another if they frequently collaborate with each other and each other's co-authors in the graph.

[Change to log scale](#) [Refresh](#) [Sort alphabetically](#) [Save as image](#)

Tables

Publications per year (.CSV File)	Co-authors (.CSV File)
Year	Author Publications with Borner, Katy
2001	2
2002	4
2003	2
2004	7
2005	7
2006	3
2007	10

Download Data

General Statistics

- 36 publication(s) from 2001 to 2010 ([.CSV File](#))
- 80 co-author(s) from 2001 to 2010 ([.CSV File](#))

Co-Author Network

([GraphML File](#))

Save as Image (.PNG file)

Tables

- Publications per year ([.CSV File](#))
- Co-authors ([.CSV File](#))

http://vivo-netsci.cns.iu.edu/vivo/visualization?uri=http%3A%2F%2Fvivo-trunk.indiana.edu%2Findividual%2FPerson74&vis=person_level&render_mode=standalone

36 publication(s) from 2001 to 2010 ([.CSV File](#))

Year	Publications
2001	2
2002	4
2003	2
2004	7
2005	7
2006	3
2007	10
2010	1

80 co-author(s) from 2001 to 2010 ([.CSV File](#))

Year	Count	Co-Author(s)
2001	1	Chen C.
2002	3	Chen C.; McMahon T.; Feng Y.
2003	2	Chen C.; Boyack K.W.
2004	17	Sengupta A.; Penumarthi S.; Thakur S.; Sooriamurthi R.; Maru J.T.; Shiffrin R.M.; Mane K.; Moor K.A.;

Co-author network ([GraphML File](#))

```

1 <?xml version="1.0" encoding="UTF-8"?>
2 <graphml xmlns="http://graphml.graphdrawing.org/xmlns"
3   xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
4   xsi:schemaLocation="http://graphml.graphdrawing.org/xmlns
5     http://graphml.graphdrawing.org/xmlns/1.0/graphml.xsd">
6 <key id="label" for="node" attr.name="label" attr.type="string" />
7 <key id="number_of_authored_works" for="node" attr.name="number_of_authored_works" attr.type="int" />
8 <key id="num_unknown_publication" for="node" attr.name="num_unknown_publication" attr.type="int" />
9 <key id="num_latest_publication" for="node" attr.name="num_latest_publication" attr.type="int" />
10 <key id="latest_publication" for="node" attr.name="latest_publication" attr.type="int" />
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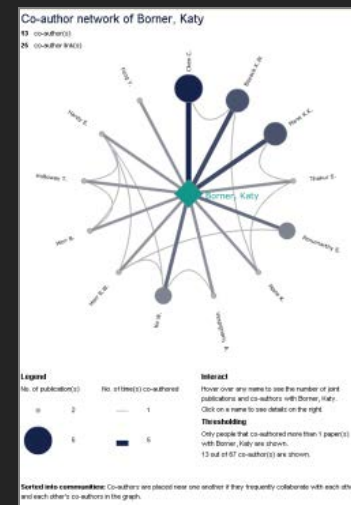
```

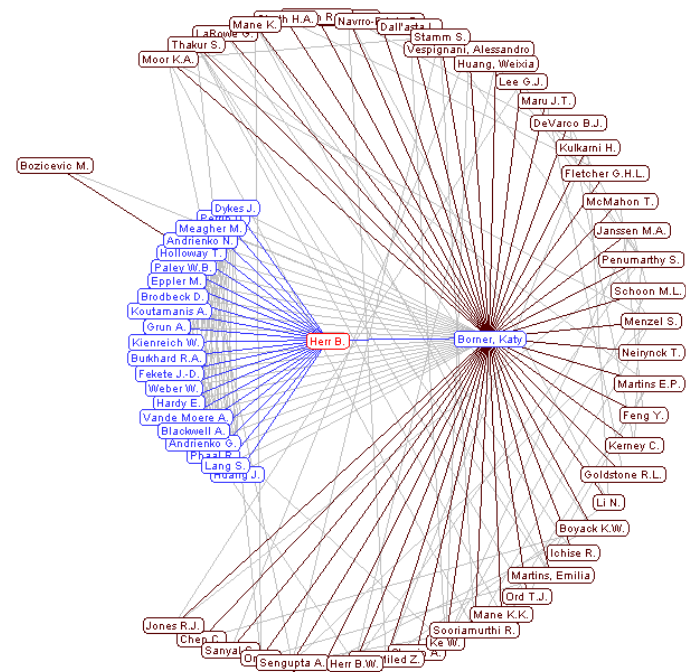
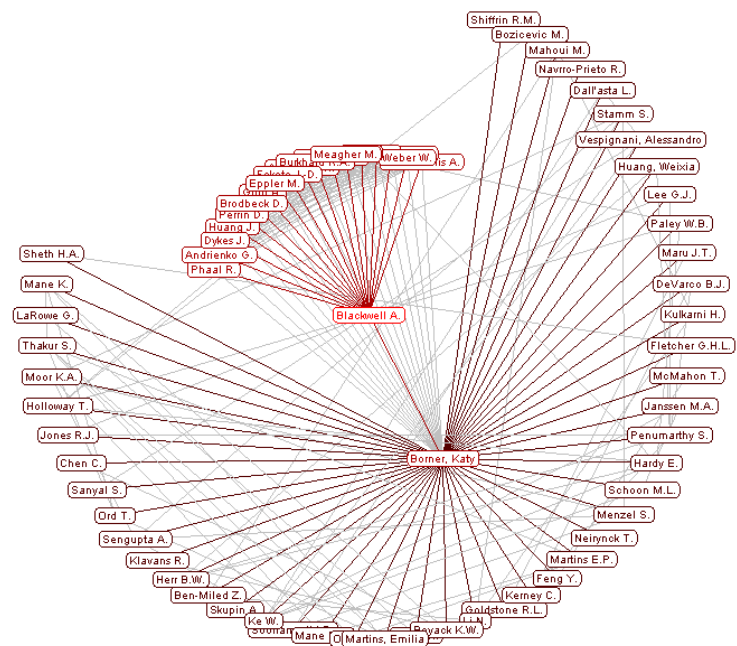
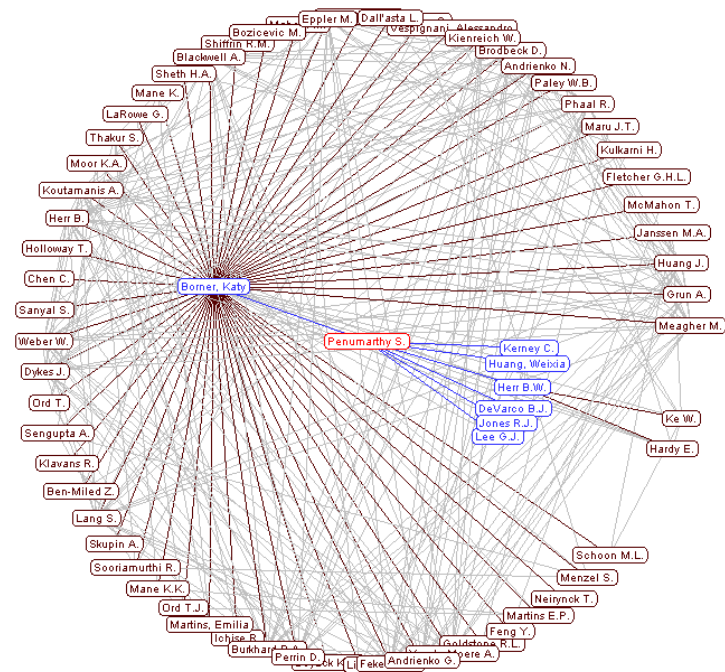
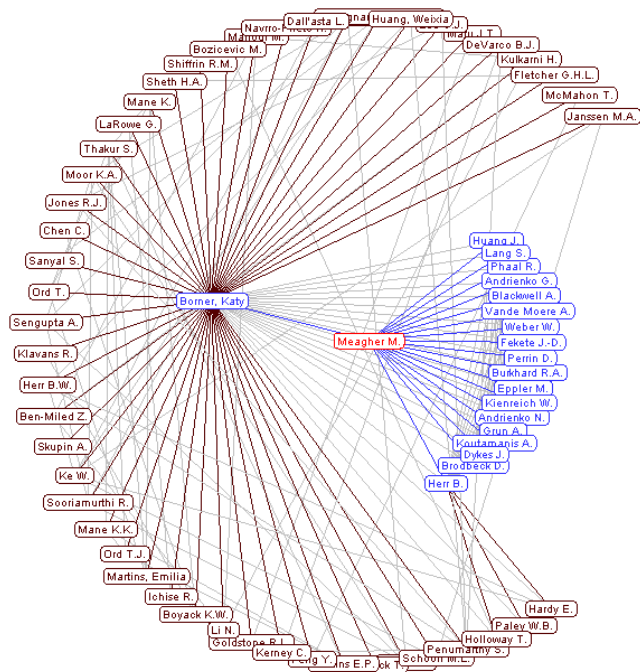
Save as Image (.PNG file)

Publications per year ([.CSV File](#)), see top file.

Co-authors ([.CSV File](#))

Co-Author	Count
Andrienko G.	1
Andrienko N.	1
Ben-Miled Z.	1
Blackwell A.	1
Boyack K.W.	4
Bozicevic M.	1
Brodbeck D.	1
Burkhard R.A.	1
Chen C.	5





About the Cyberinfrastructure Shell

The Cyberinfrastructure Shell (CIShell) is an open source, community-driven platform for the integration and utilization of datasets, algorithms, tools, and computing resources. Algorithm integration support is built in for Java and most other programming languages. Being Java based, it will run on almost all platforms. The software and specification is released under an Apache 2.0 License.

CIShell is the basis of [Network Workbench](#), [TexTrend](#), [Sci2](#) and the upcoming [EpiC](#) tool.

CIShell supports remote execution of algorithms. A standard web service definition is in development that will allow pools of algorithms to transparently be used in a peer-to-peer, client-server, or web front-end fashion.

CIShell Features

A framework for easy integration of new and existing algorithms written in any programming language

Using CIShell, an algorithm writer can fully concentrate on creating their own algorithm in whatever language they are comfortable with. Simple tools are provided to then take their algorithm and

Learn More...

- [CIShell Papers](#)
- [CIShell Powered Tools](#)
- [Algorithms](#)
- [Plugins \(coming soon\)](#)
- [Misc. Tool Documentation](#)
- CIShell Web Services (coming soon)
- [Screenshots](#)

Getting Started...

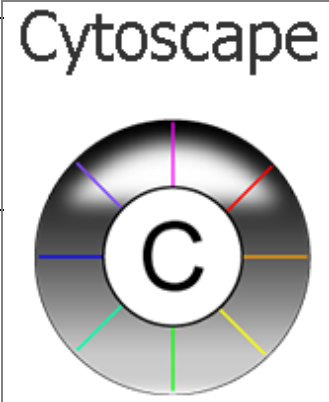
- [Documentation & Developer Resources](#)
- [Download](#)

Getting Involved...

- [Contact Us](#)

CIShell Developer Guide is at <http://cishell.wiki.cns.iu.edu>

Additional Sci2 Plugins are at <http://sci2.wiki.cns.iu.edu/3.2+Additional+Plugins>

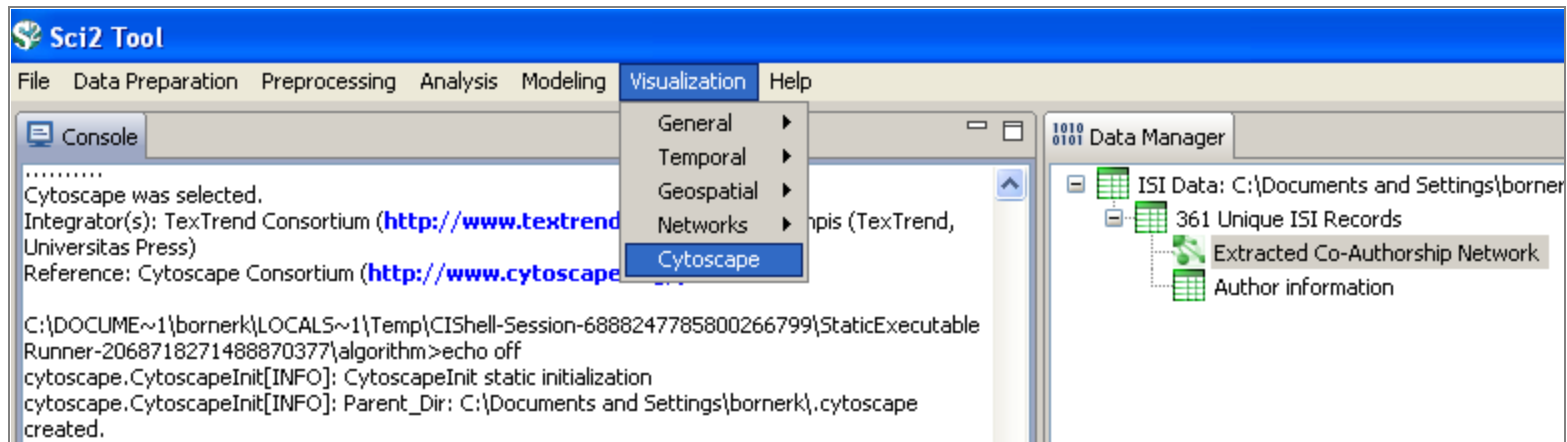


Adding more layout algorithms and network visualization interactivity via Cytoscape <http://www.cytoscape.org>.

Simply add *org.textrend.visualization.cytoscape_0.0.3.jar* into your /plugin directory.

Restart Sci2 Tool.

Cytoscape now shows in the Visualization Menu.



Select a network in Data Manager, run Cytoscape and the tool will start with this network loaded.

Search:

Control Panel

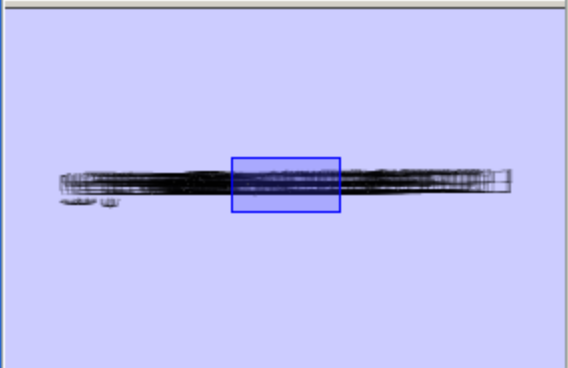
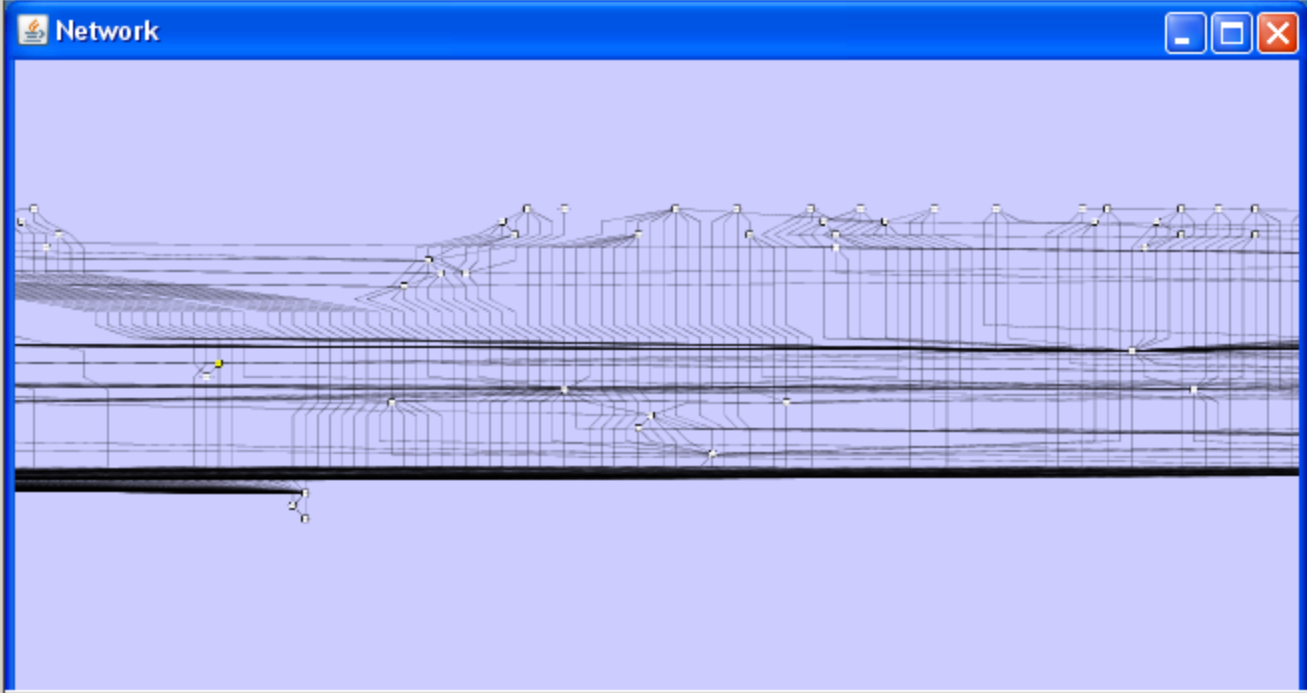
Network

VizMapper™

Editor

Filters

Network	Nodes	Edges
● Network	247(1)	891(0)



Data Panel

ID

Munoz, Ma

FileEditViewSelectLayoutPluginsHelp

Control Panel

NetworkVizMapper™

Network

Network

Rotate

Scale

Align and Distribute

Settings...

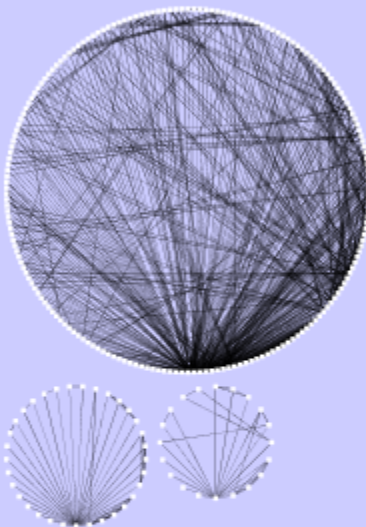
yFiles

Cytoscape Layouts

JGraph Layouts

Search:

Network



Data Panel

ID

Node Attribute Browser

Edge Attribute Browser

Network Attribute Browser

Welcome to Cytoscape 2.6.2

Right-click + drag to ZOOM

Middle-click + drag to PAN

A number of other projects recently adopted OSGi and/or CIShell:

- Cytoscape* (<http://cytoscape.org>) Led by Trey Ideker at the University of California, San Diego is an open source bioinformatics software platform for visualizing molecular interaction networks and integrating these interactions with gene expression profiles and other state data (Shannon et al., 2002).
- *Taverna Workbench* (<http://taverna.org.uk>) Developed by the myGrid team (<http://mygrid.org.uk>) led by Carol Goble at the University of Manchester, U.K. is a free software tool for designing and executing workflows (Hull et al., 2006). Taverna allows users to integrate many different software tools, including over 30,000 web services.
- *MAEviż* (<https://wiki.ncsa.uiuc.edu/display/MAE/Home>) Managed by Jong Lee at NCSA is an open-source, extensible software platform which supports seismic risk assessment based on the Mid-America Earthquake (MAE) Center research.
- *TEXTrend* (<http://textrend.org>) Led by George Kampis at Eötvös Loránd University, Budapest, Hungary supports natural language processing (NLP), classification/mining, and graph algorithms for the analysis of business and governmental text corpuses with an inherently temporal component.
- *DynaNets* (<http://www.dynanets.org>) Coordinated by Peter M.A. Sloot at the University of Amsterdam, The Netherlands develops algorithms to study evolving networks.

As the functionality of OSGi-based software frameworks improves and the number and diversity of dataset and algorithm plugins increases, the capabilities of custom tools will expand.



Consortium

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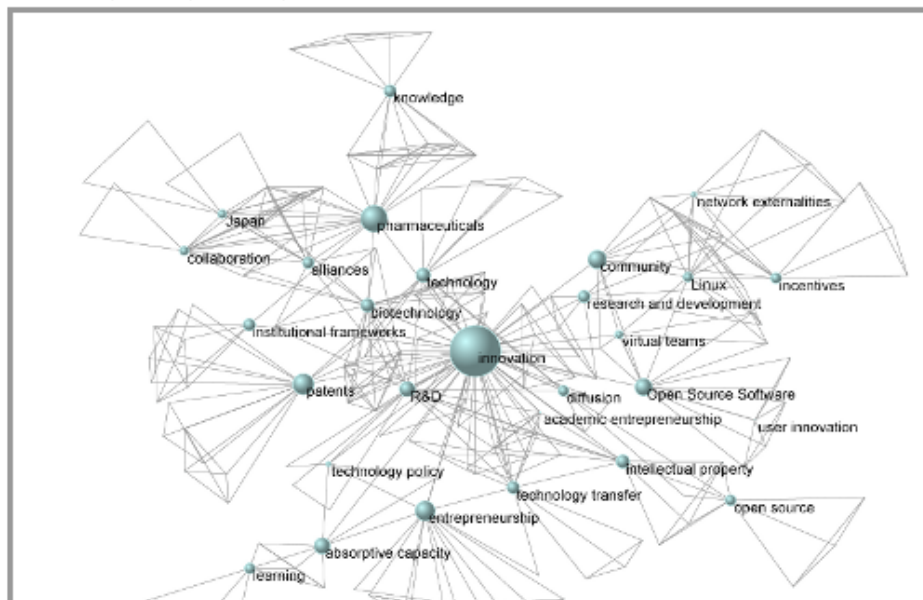
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TEXTrend: Development of a business and governmental decision support toolbox using trend- and text-analysis tools

The two interconnected objectives of the **TEXTrend project** are (1) the creation of an integrated TEXTrend toolkit and service basis, and (2) the elaboration of **demonstrative applications** in varied fields of business and governmental decisions, exemplified by use cases.



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toolkit trend tv university version web

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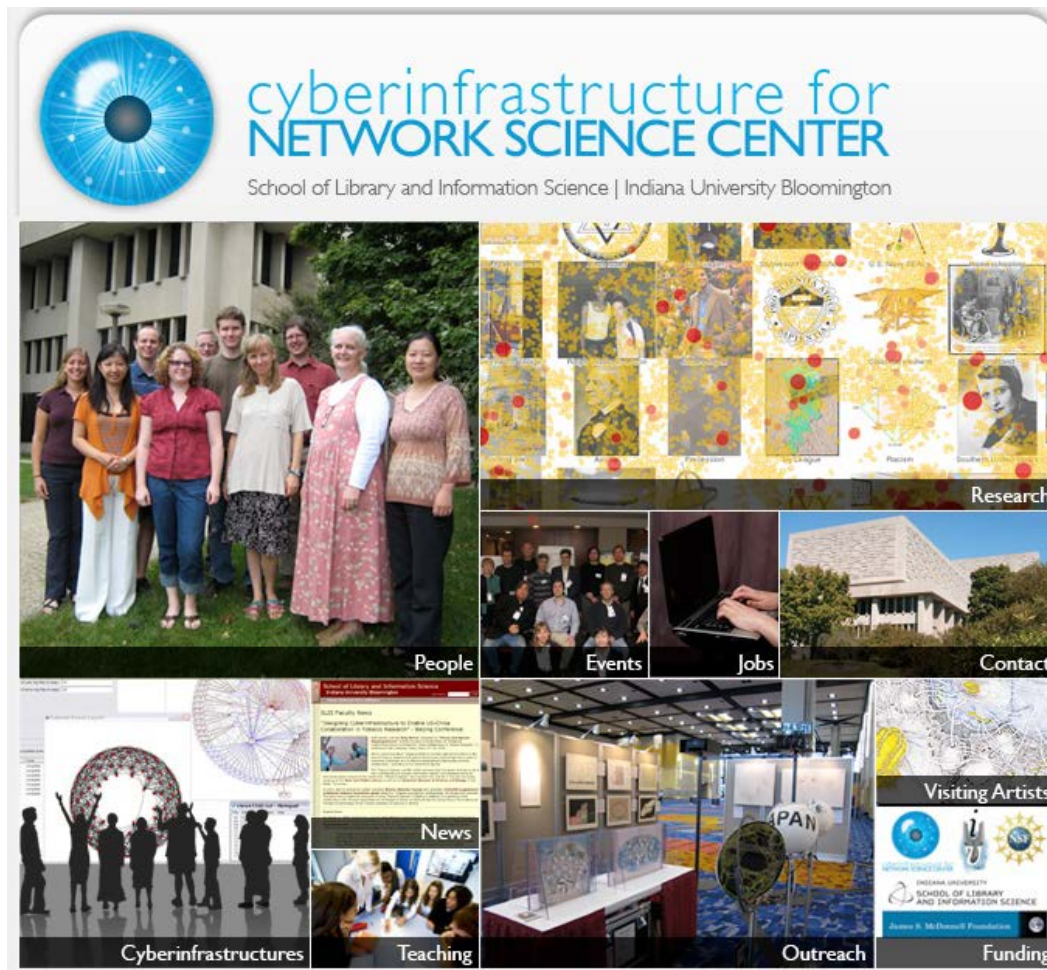


Created by
www.wordle.net

TEXTrend adds R bridge, WEKA, Wordij, CFinder, and more.

See the latest versions of TEXTrend Toolkit modules at

http://textrend.org/index.php?option=com_content&view=article&id=47&Itemid=53



All papers, maps, tools, talks, press are linked from <http://cns.iu.edu>

CNS Facebook: <http://www.facebook.com/pages/Cyberinfrastructure-for-Network-Science-Center/144339535612571>

Mapping Science Exhibit Facebook: <http://www.facebook.com/mappingscience>