

# Macroscopes

Decision making in science, industry, and politics, as well as in daily life, requires that we make sense of data sets representing the structure and dynamics of complex systems. Macroscopes provide a "vision of the whole," helping us "synthesize" the related elements and enabling us to detect patterns, trends, and outliers while granting access to myriad details. Rather than make things larger or

smaller, macroscopes let us observe what is at once too great, slow, or complex for the human eye and mind to notice and comprehend.



Microscopes



Telescopes



Macroscopes

## **Plug-and-Play Macroscopes**

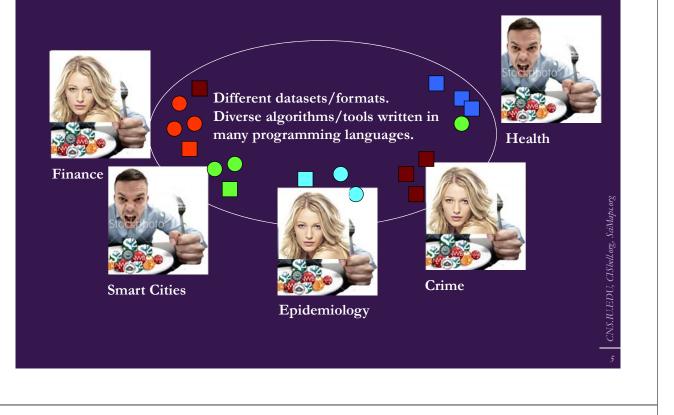
While microscopes and telescopes are **physical instruments**, macroscopes resemble **continuously changing bundles of software plug-ins**. Macroscopes make it easy to select and combine algorithm and tool plug-ins but also interface plug-ins, workflow support, logging, scheduling, and other plug-ins needed for scientifically rigorous yet effective work.

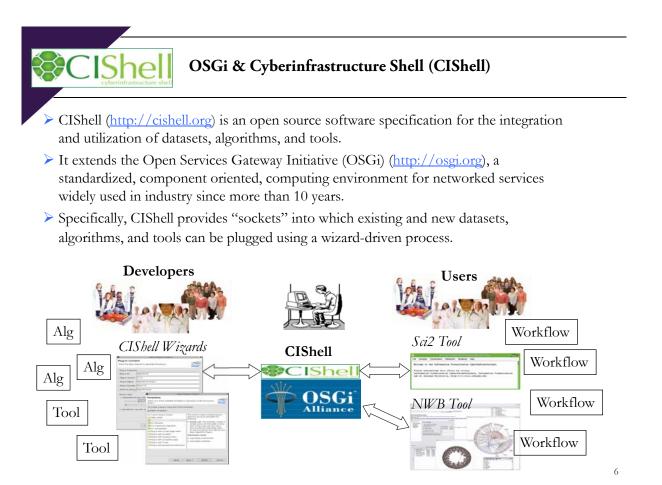
They make it easy to share plug-ins via email, flash drives, or online. To use new plugins, simply copy the files into the plug-in directory, and they appear in the tool

menu ready for use. No restart of the tool is necessary. Sharing

algorithm components, tools, or novel interfaces becomes as easy as sharing images on Flickr or videos on YouTube. Assembling custom tools is as quick as compiling your custom music collection.

# Plug-and-Play Macroscopes





NetworkWorkbench

Network Workbench Tool

http://nwb.cns.iu.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.

The tool provides more 160 plugins that support the preprocessing, analysis, modeling, and visualization of networks.

It has been downloaded more than 110,000 times.



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.

# NetworkWorkbench

### Network Workbench Tool

http://nwb.cns.iu.edu

Velcome to the Network Workberd isualization of small, medium, a the Network Workberd process ward. The primary investgator lessandro Vespignani, Dr. Star he NWB tool was developed by hintan Tank, Joseph Biberski vek Thakre, Ann McCranie, Alessandro Vespignani, and Katy Börner. tetwork Work Bease cite as follows: Weighted automatically Remove Scheduler Remove From List Remove completed automatically Remove Remove From List Remove completed automatically Remove Automatical Science Context (Science Context) Science Completed automatically Remove Linke Science Completed Automatically Remove Linke Sc	e Preprocessing Modeling	Analysis Visualization Scien Network Analysis Toolkit (N	1	Menu		1010 Data Manager	
Node Outdegree      Jagorithm Name    Weighted and Directed      Node Outdegree      Indegree Distribution      Outdegree Distribution      Node Outdegree Distribution      Outdegree Distribution      Outdegree Distribution      Node Outdegree Distribution      Discrete Network Voranics      Textual      Textual      Fextual      PageRank      HITS      Dyad Reciprocity      Arc Reciprocity      Adjacency Transitivity      Node Component Clustering      Stonduler      Remove From List    Remove completed automatically Remove      I Algorithm Name    Date	Velcome to the Network Workb isualization of small, medium, a		f processing, r	nodeling, analysis, ar			ty Desktop \TOOLS \nwb-2009.09.
he NWB tool was developed by bintan Tank, Joseph Biberstin ortunato, Ben Markines, Feix    Discrete Network Dynamics Textual    Outdegree Distribution      K-Nearest Neighbor single Node In-Out Degree Correlations    K-Nearest Neighbor Single Node In-Out Degree Correlations    Data Manager      Vex Thaire, A MoCranie, Alessandro Vespignani, and Katy Borner.    PageRank HITS    PageRank HITS      lease cite as follows: WB Team. (2006). Network Workbench Tool. Indiana University, Nor Idrigan, http://nwb.slis.indiana.edu.    Dyad Reciprocity Arc Reciprocity Adjacency Transitivity    Data Manager      Scheduler    Blondel Community Detection    Extract K-Core Annotate K-Coreness    Extract K-Core      I    Algorithm Name    Date    Time    % Complete							
ortunato, Ben Markines, Feix :    Total and intervention of the provided and the p	The second second second second	and the second se				Data Managan	
etwork Wo    Console    erinfrastructure Shell (http://cishe    PageRank      yberinfrast    Console    einfrastructure Shell (http://cishe    PageRank      HITS    Dyad Reciprocity    Arc Reciprocity      WB Team. (2006). Network Workbench Tool. Indiana University, Nor    Dyad Reciprocity    Arc Reciprocity      and    use relacted    Weak Component Clustering    Strong Component Clustering      Scheduler    Blondel Community Detection    Extract K-Core      Annotate K-Coreness    Annotate K-Coreness    Annotate K-Coreness			Course Made		elations		Janager
WB Team. (2006). Network Workbench Tool. Indiana University, Nor  Arc Reciprocity    Ichigan, http://nwb.slis.indiana.edu.  Adjacency Transitivity    Adjacency Transitivity  Weak Component Clustering    Scheduler  Strong Component Clustering    Remove From List  Remove completed automatically    Remove From List  Remove completed automatically    I  Algorithm Name    I  Algorithm Name	Jabuark Wa	erinfrastructure Shell (http:/	/cishe PageRank			Data N	hanager
Scheduler  Weak Component Clustering    Remove From List  Remove completed automatically    Remove From List  Remove completed automatically    Remove  Blondel Community Detection    Extract K-Core  Annotate K-Coreness    I  Algorithm Name  Date	ichigan, http://nwb.slis.indi		ty, Nor Arc Recipro	ocity			
Image: Strate Complete Statistical y Complete    Image: Strate Complete Statistical y Complete      Image: Strate Complete Strate Complete	Scheduler						
Annotate K-Coreness	Remove From List	e completed automatically R		Blondel Community Detection			
	26			% Complete			

### **Computational Proteomics**

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

Yildriim, 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.

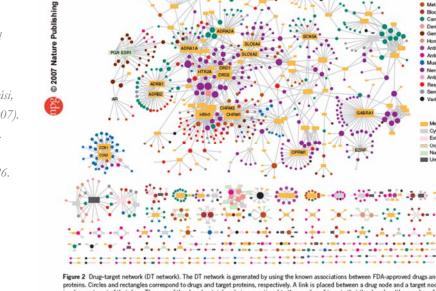
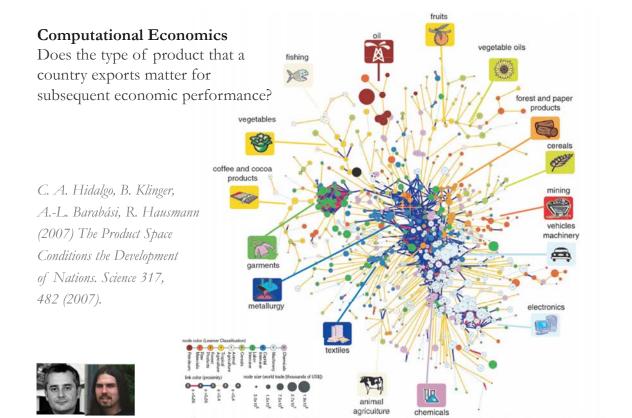




Figure 2 Drug-target network (DT network). The DT network is generated by using the known associations between FDA-approved drugs and their target proteins. Circles and rectangles correspond to drugs and target proteins, respectively. A link is placed between a drug node and a target node if the protein 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 (circle) 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.



### Fig. 1. The product space. (A) Hierarchically clustered proximity (a) 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 Learner. 10



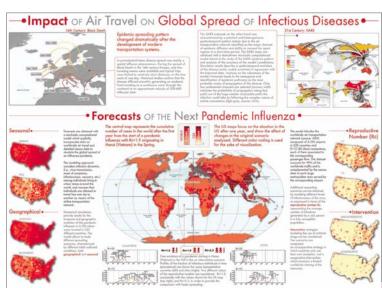
# **Computational Epidemics** Forecasting (and preventing the effects of) the next pandemic.

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).







Science of Science Tool http://sci2.cns.iu.edu

# Sci2 Tool v0.5.1 Alpha (May 4th, 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

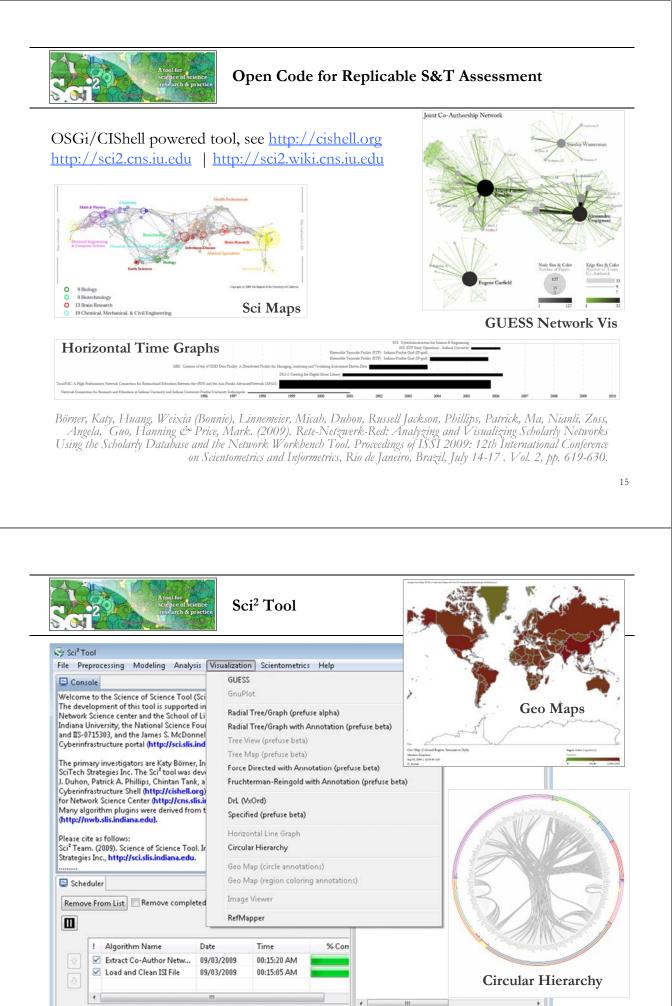
Sci<sup>2</sup> Team. (2009). Science of Science (Sci<sup>2</sup>) Tool. Indiana University and SciTech Strategies, <u>http://sci2.cns.iu.edu</u>

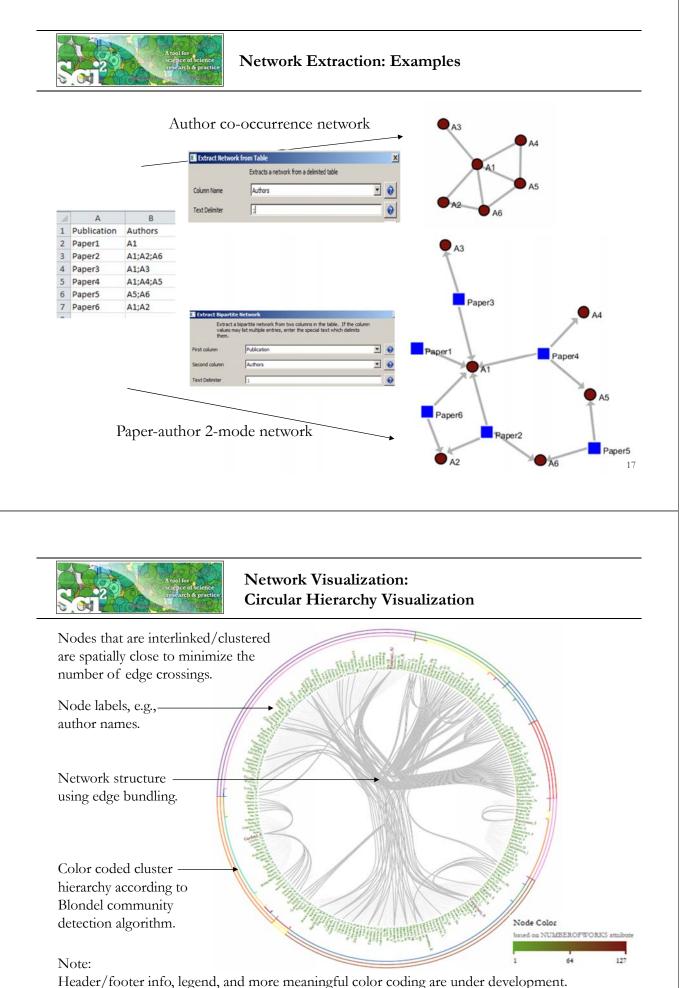


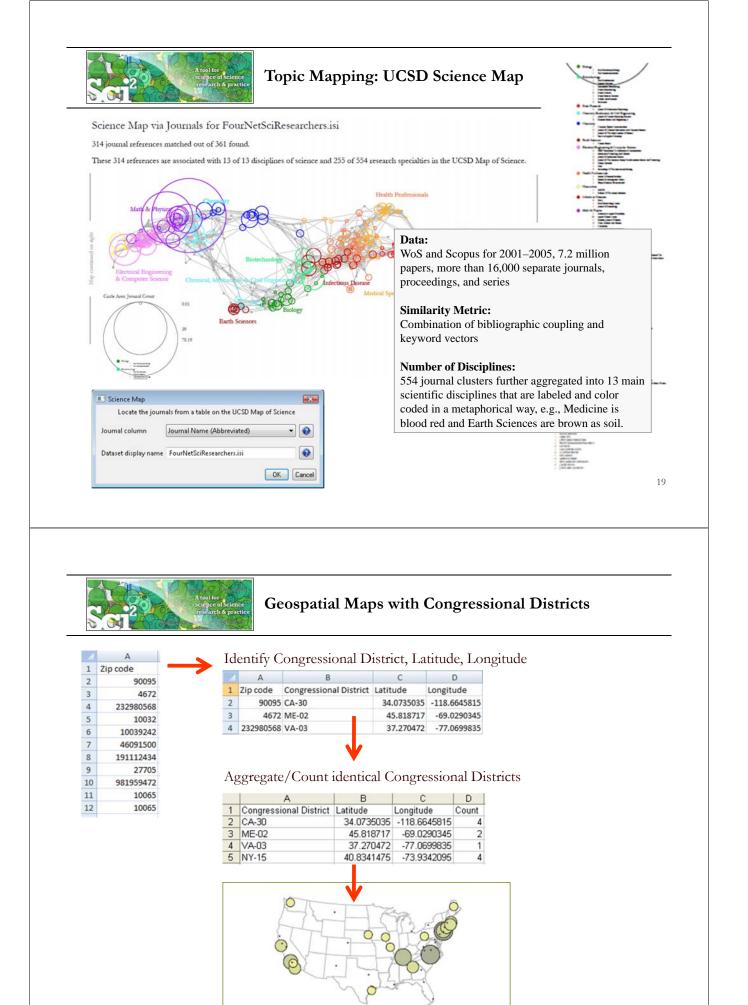


# Type of Analysis vs. Level of Analysis

	Micro/Individual (1-100 records)	Meso/Local (101–10,000 records)	Macro/Global (10,000 < records)
Statistical Analysis/Profiling	Individual person and their expertise profiles	Larger labs, centers, universities, research domains or states	All of NS SA, all of scie
Temporal Analysis (When)	Funding portfolio of one individual	ic bursts	113 Years of P Research
Geospatial Analysis (Where)	Career trajectory of one individual	intellectual le	PNAS
Topical Analysis (What)		research	VxOrd/Topic r NIH funding
Network Analysis (With Whom?)	NSF work of		NIH's
	E a.		

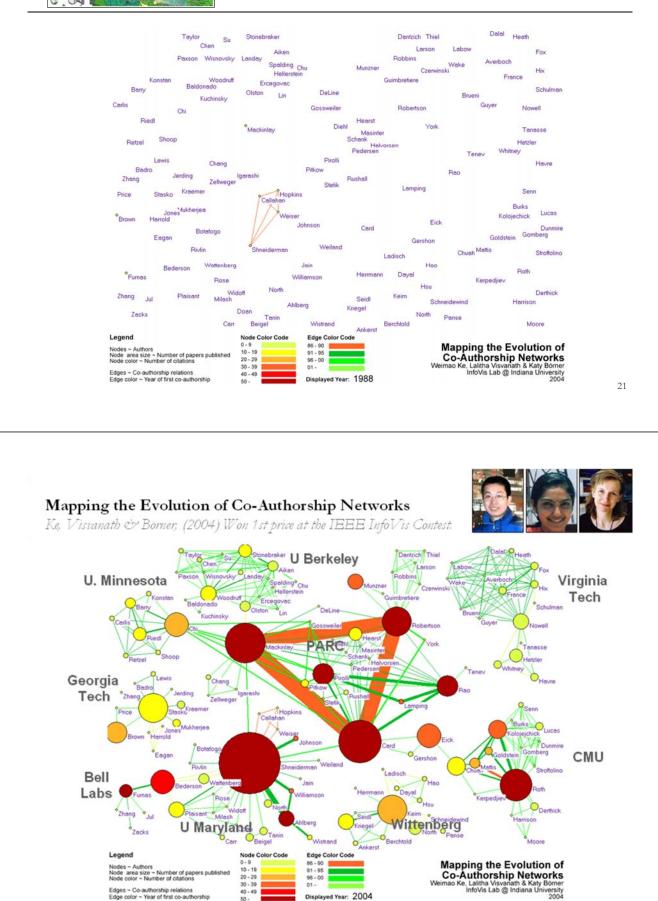


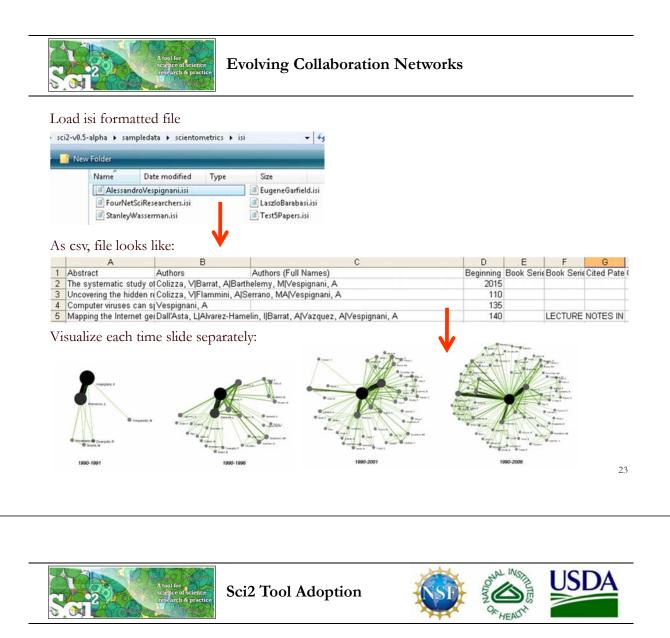






### **Evolving Collaboration Networks**

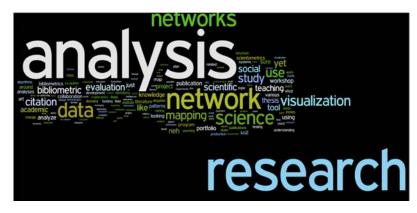


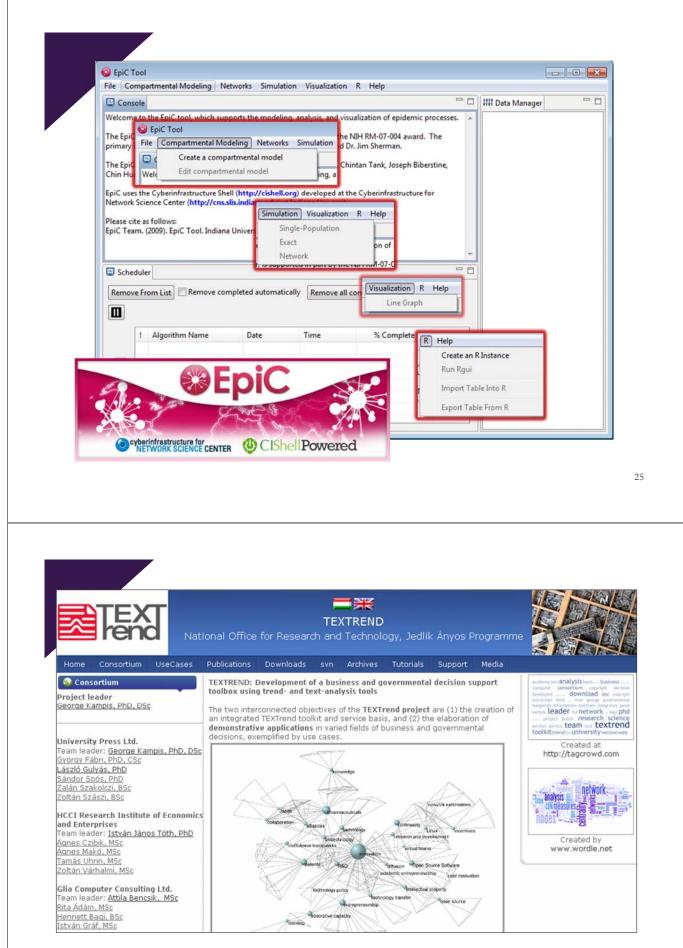


The Sci2 Tool is used by the National Science Foundation, the National Institutes of Health, and the US Department of Agriculture.

"As a new user, I am beginning with very little knowledge of the analyses and modeling techniques that Sci2 enables. I've been able to use my own dataset and follow through some of the workflows to the point of generating the first network and time horizon visuals. That was so exciting I stayed up far past bedtime to get to the visuals."

Dr. Suzanne A. Pierce, Center for International Energy and Environmental Policy Jackson School of Geosciences, The University of Texas at Austin





TEXTrend adds WEKA, UIMA, Wordij, CFinder, and more. See the latest versions of TEXTrend Toolkit modules at *http://textrend.org* 



### CIShell - Integrate New Algorithms

### 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, Sci<sup>2</sup> 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, clientserver, 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...

- <u>CIShell Papers</u>
- <u>CIShell Powered Tools</u>
  Algorithms
- Algorithms
  Plugins (co
- <u>Plugins (coming soon)</u>
  <u>Misc. Tool Documentation</u>
- · CIShell Web Services (coming soon)
- Screenshots

### Getting Started...

- Documentation & Developer Resources
- <u>Download</u>

### Getting Involved...

<u>Contact Us</u>

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

# Shell OSGi/CIShell Adoption

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

Cytoscape (<u>http://cytoscape.org</u>) 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). MAEviz (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.  $\geq$ Taverna Workbench (<u>http://taverna.org.uk</u>) 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 Europe to integrate many different software tools, including over 30,000 web services. TEXTrend (<u>http://textrend.org</u>) 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 (<u>http://www.dynanets.org</u>) Coordinated by Peter M.A. Sloot at the University of Amsterdam, The Netherlands develops algorithms to study evolving networks.  $\geq$ SISOB (http://sisob.lcc.uma.es) An Observatory for Science in Society Based in Social Models. 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.

# Plug-and-Play Macroscopes

Sci2

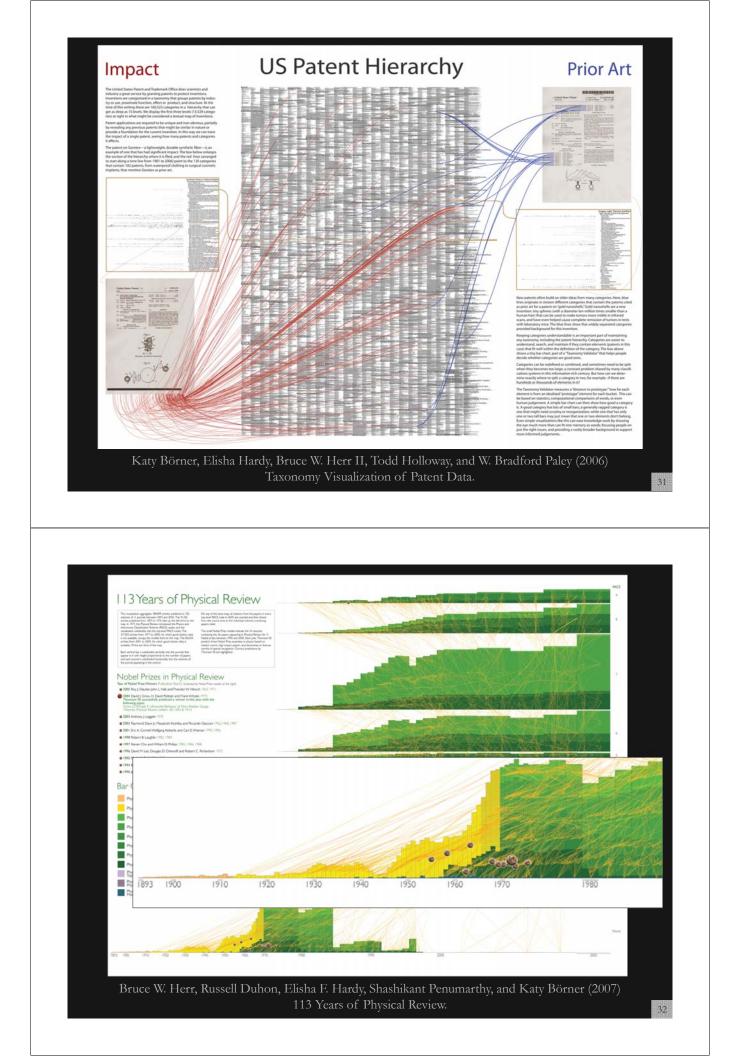
CS

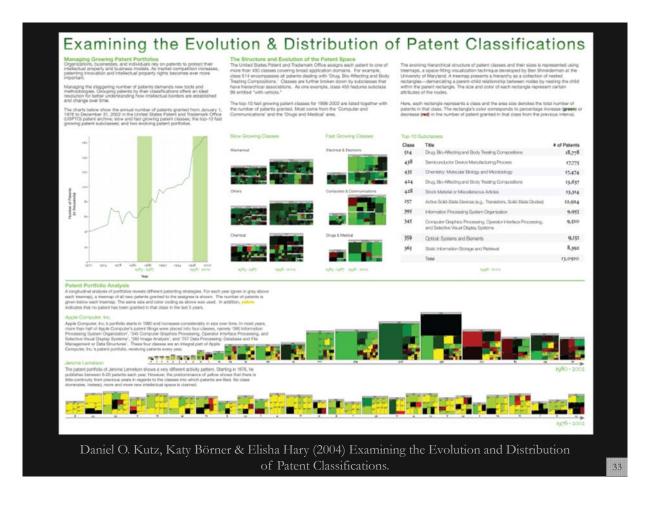
Bio SNA Phys Converters

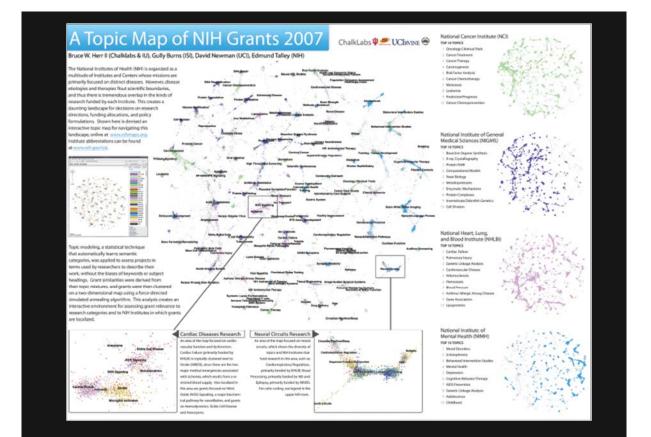
NWB

Maps created using Sci2 are travelling in the "Expedition Zukunft" science train visiting 62 cities in 7 months, 12 coaches, 300 m long. <u>http://www.expedition-zukunft.de</u>

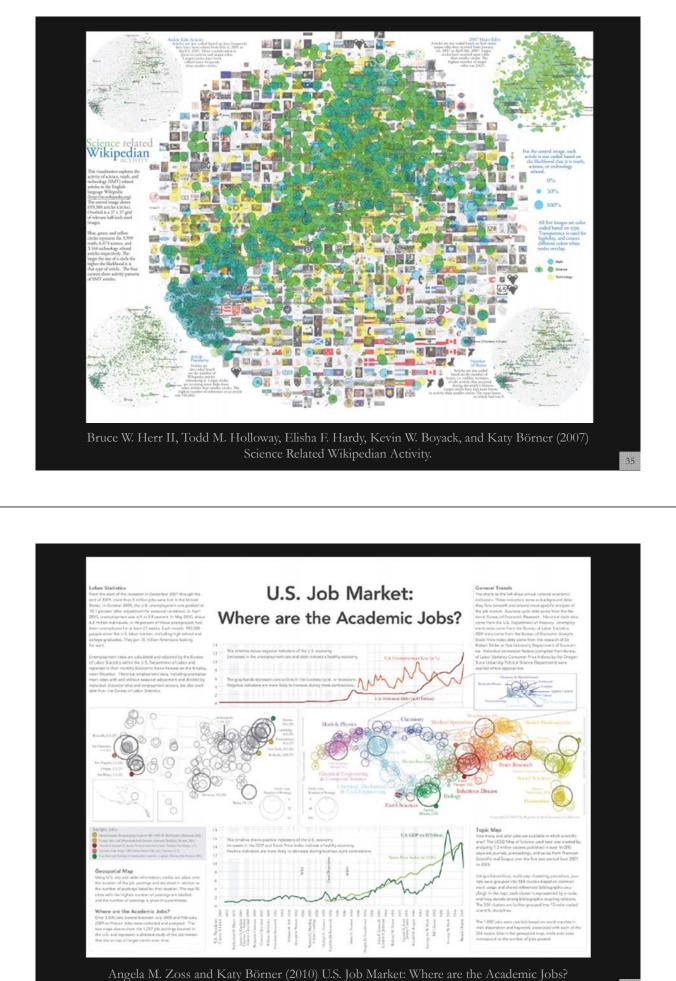
And they are part of the international Mapping Science exhibit: http://scimaps.org

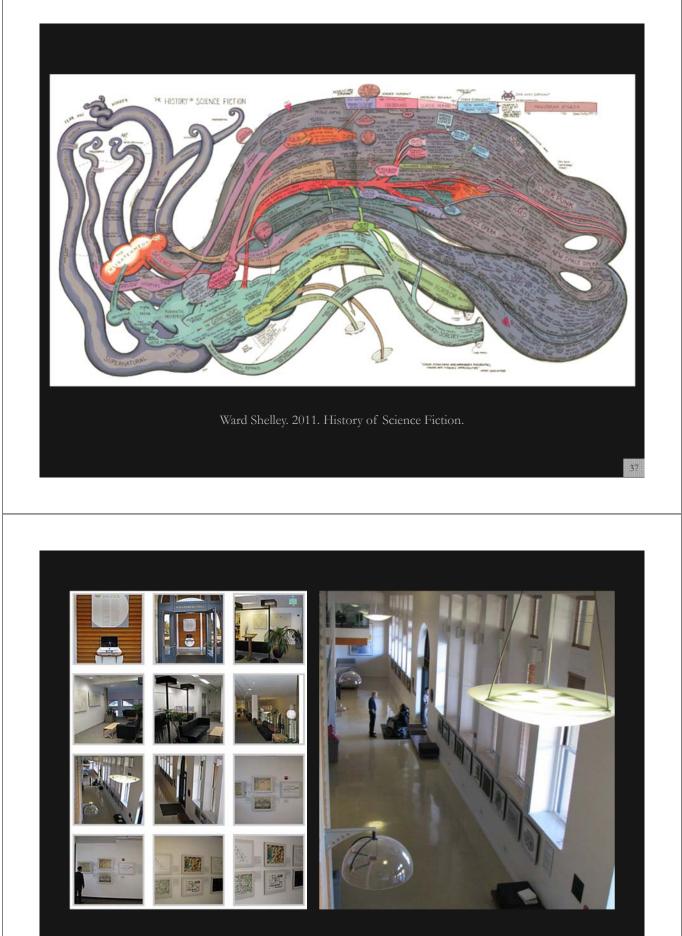






Bruce W. Herr II, Gully Burns, David Newman, Edmund Talley (2007) A Topic Map of NIH Grants.





Debut of 5<sup>th</sup> Iteration of the Mapping Science Exhibit at MEDIA X took place at Wallenberg Hall, Stanford University, <u>http://mediax.stanford.edu</u>, <u>http://scaleindependentthought.typepad.com/photos/scimaps</u>



### References

Börner, Katy, Chen, Chaomei, and Boyack, Kevin. (2003). Visualizing Knowledge Domains. In Blaise Cronin (Ed.), *ARIST*, Medford, NJ: Information Today, Volume 37, Chapter 5, pp. 179-255. http://iyl.slis.indiana.edu/km/pub/2003-borner-arist.pdf

Shiffrin, Richard M. and Börner, Katy (Eds.) (2004). **Mapping Knowledge Domains**. Proceedings of the National Academy of Sciences of the United States of America, 101(Suppl\_1). http://www.pnas.org/content/vol101/suppl\_1/

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

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

Börner, Katy (2010) Atlas of Science. MIT Press. http://scimaps.org/atlas

Scharnhorst, Andrea, Börner, Katy, van den Besselaar, Peter (2011) **Models of Science Dynamics**. Springer Verlag.



Visualizing What We Know Ker Ruse



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

CNS Facebook: http://www.facebook.com/cnscenter Mapping Science Exhibit Facebook: http://www.facebook.com/mappingscience