International, Interdisciplinary Plug-and-Play Macroscopes

Katy Börner

Cyberinfrastructure for Network Science Center, Director Information Visualization Laboratory, Director School of Library and Information Science Indiana University, Bloomington, IN <u>katv@indiana.edu</u>





With special thanks to the members of the Cyberinfrastructure for Network Science Center, the Sci2 team, NWB team, EpiC team, and OSGi/CIShell tool developers in Europe.

From OpenSHAPA to Open Data Sharing Workshop NSF, Stafford Place II, 4121 Wilson Boulevard, 5th Floor, Arlington, VA, 22230

September 15, 2011



By George Shaw, Phillip Decamp, and Deb Roy, MIT

The map shows the experimental setup used to record the language development of Deb Roy's son at home. Approximately **10 hours of high fidelity audio and video were recorded on a daily basis from birth to age three.** The resulting corpus contains over **100,000 hours of multitrack recordings** and constitutes the most comprehensive record of a child's development made to date.

To analyze the massive data set, new tools had to be developed **to rapidly annotate, transcribe, analyze, and visualize salient patterns of behavior and interaction**. Results of this analysis comprise **human movement traces, word birth places, and social networks**, see map. For most children, language development is steady, progressive, and to a casual observer effortless. But for some children—those with developmental delays due to biological or environmental causes— language is a major developmental hurdle. Understanding the **regularities in home environments** is essential to understanding mechanisms of language acquisition, causes of delay, and ultimately appropriate intervention procedures.





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 of PNAS	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

Needs-Driven Workflow Design using a modular data acquisition/analysis/ modeling/ visualization pipeline as well as modular visualization layers.



Börner, Katy (2010) Atlas of Science. MIT Press. 5

COMMUNICATIONS

Börner, Katy. (March 2011). Plug-and-Play Macroscopes. *Communications of the ACM*, 54(3), 60-69.

Video and paper are at <u>http://www.scivee.tv/node/27704</u>



Designing "Dream Tools"

Many of the best micro-, tele-, and macroscopes are designed by **scientists keen to observe and comprehend what no one has seen or understood before.** Galileo Galilei (1564–1642) recognized the potential of a spyglass for the study of the heavens, ground and polished his own lenses, and used the improved optical instruments to make discoveries like the moons of Jupiter, providing quantitative evidence for the Copernican theory.

Today, scientists repurpose, extend, and invent new hardware and software to

create **"macroscopes"** that may solve both local and global challenges.

Plug-and-play macroscopes **empower** me, my students, colleagues, and 100,000 others that downloaded them.



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. Analysis, navigation, and management of these continuously evolving data sets require a new kind of data-analysis and visualization tool we call a macroscope (from the Greek macros, or "great," and skopein, or "to observe") inspired by de Rosnay's futurist science writings.

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



Macroscopes cont.

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.



9





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.



Timeline Visualization: Example

Project Timeline



2	Sci ² Tool A tool for science of science research & practice Email Address Password Login
Forgot your pas To recover your Not registered Register now	ssword? account password, please visit our <u>password recovery page</u> . yet?
Tutorials Scott Weingart, J Biberstine (2010 Science, Indiana	Katy Börner (2010) Science of Science Research and Tools (12 Tutorials). Reporting Branch, Office of Extramural Research/Office of the Director, National Institutes of Health, Bethesda, MD. • Tutorial #01: Science of Science Research • Tutorial #02: Network Science / Information Visualization • Tutorial #03: CIShell Powered Tools: Network Workbench and Science of Science Tool • Tutorial #03: Geospatial Analysis — Burst Detection • Tutorial #05: Geospatial Analysis and Mapping • Tutorial #06: Topical Analysis and Visualization • Tutorial #07: Tree Analysis and Visualization • Tutorial #08: Network Analysis and Visualization • Tutorial #09: Large Network Analysis and Visualization • Tutorial #10: Using the Scholarly Database at IU • Tutorial #11: YIVO National Researcher Networking • Tutorial #12: Future Developments
	Geetha Senthal (2010). <u>Multidisciplinary Nature of Work With Reference to PIs and ICs Within a Portfolio</u> . PA Group at NIH. NIH Office of Extramural Research and Katy Börner (2010) <u>Network Visualizations Using SPIRES Data and the Sci2 Tool</u> . Office of Extramural Research at NIH.



Preprocessing

Extract Top N% Records Extract Top N Records Normalize Text Slice Table by Line

Extract Top Nodes Extract Nodes Above or Below Value Delete Isolates

Extract top Edges Extract Edges Above or Below Value Remove Self Loops Trim by Degree MST-Pathfinder Network Scaling Fast Pathfinder Network Scaling

Snowball Sampling (in nodes) Node Sampling Edge Sampling

Symmetrize Dichotomize Multipartite Joining

Geocoder

Extract ZIP Code

Sci² Tool: Algorithms

Modeling Random Graph Watts-Strogatz Small World Barabási-Albert Scale-Free TARL

Analysis Network Analysis Toolkit (NAT) Unweighted & Undirected Node Degree Degree Distribution

> K-Nearest Neighbor (Java) Watts-Strogatz Clustering Coefficient Watts Strogatz Clustering Coefficient over K

Diameter Average Shortest Path Shortest Path Distribution Node Betweenness Centrality

Weak Component Clustering Global Connected Components

Extract K-Core Annotate K-Coreness

HITS

Weighted & Undirected

Clustering Coefficient Nearest Neighbor Degree Strength vs Degree Degree & Strength Average Weight vs End-point Degree Strength Distribution Weight Distribution Randomize Weights

Blondel Community Detection

HITS Unweighted & Directed Node Indegree Node Outdegree Indegree Distribution

Outdegree Distribution

K-Nearest Neighbor Single Node in-Out Degree Correlations

Dyad Reciprocity Arc Reciprocity Adjacency Transitivity

Weak Component Clustering Strong Component Clustering



Sci² Tool: Algorithms cont.

Extract K-Core Annotate K-Coreness HITS PageRank Weighted & Directed HITS

Weighted PageRank

Textual Burst Detection

Soon:

Visualization

GnuPlot GUESS Image Viewer

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 Bar Graph Circular Hierarchy Geo Map (Circle Annotation Style) Geo Map (Colored-Region Annotation Style) Science Map (Circle Annotation)

Scientometrics

Remove ISI Duplicate Records Remove Rows with Multitudinous Fields Detect Duplicate Nodes Update Network by Merging Nodes

Extract Directed Network Extract Paper Citation Network Extract Author Paper Network

Extract Co-Occurrence Network

Extract Word Co-Occurrence Network Extract Co-Author Network Extract Reference Co-Occurrence (Bibliographic Coupling) Network

Extract Document Co-Citation Network

19

Database support for ISI and NSF data.







Network Visualization: Example







The Sci² Tool is used by NSF, NIH, USDA, and private foundations.

Upcoming Tutorial: Sci2: A Tool of Science of Science Research and Practice

Instructor:	Dr. Katy Börner, Indiana University
Time/Date:	8:30a-11:30a on Oct 17, 2011
Place:	Room II-555 in NSF's Stafford Place II Conference Center,
	4121 Wilson Boulevard, Arlington, Virginia 22230, USA
Audience:	This tutorial is designed for researchers, practitioners, program
	staff from federal agencies interested to use advanced data
	mining algorithms and visualizations in their work and daily
	decision making.
Cost:	Free. Registration by Oct 10, 2012 required.
Register:	Please use <u>http://www.surveymonkey.com/s/MVC8LWW</u> to
-	register by Oct 10, 2012. NSF will issue visitor badges.



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.

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 (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.

fruits

25



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. 26



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).





- cpr	
File	ompartmental Modeling Networks Simulation Visualization R Help
Welco	nsole iffit Data Manager
weico	© EpiC Tool
The E	File Compartmental Modeling Networks Simulation d Dr. Jim Sherman.
The E	Create a compartmental model
Chin	WelqEdit compartmental modeling, a
EpiC u	ses the Cyberinfrastructure Shell (http://cishell.org) developed at the Cyberinfrastructure for
Netwo	Simulation Visualization R Help
EpiC 1	eam. (2009). EpiC Tool. Indiana Univer Single-Population
	Exact pn of
E Se	eduler
	Visualization R Help
Kem	Line Graph
ш	
	! Algorithm Name Date Time % Complete R Help
	Create an R Instance
5h	EPIC Import Table Into R
	Export Table From R
200	
G	cyberinfrestructure for
	Chall OSCi/CIShell Adoption
CI	Shell OSGi/CIShell Adoption
CI	of other projects recently adopted OSCi and /or CIShelly
Imber	of other projects recently adopted OSGi and/or CIShell:
CI umber Cytosca	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: pe (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegon n source bioinformatics software platform for visualizing molecular interaction
CI Imber Cytosca in open	of other projects recently adopted OSGi and/or CIShell: <i>be (<u>http://cytoscape.org</u>) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction the sand integrating these interactions with gene expression profiles and other state</i>
Imber Cytosca in openetwo Shann	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: pe (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction :ks and integrating these interactions with gene expression profiles and other state ton et al., 2002).
Cytosca Cytosca in openetwo Shann MAE	of other projects recently adopted OSGi and/or CIShell: <i>be (http://cytoscape.org)</i> Led by Trey Ideker at the University of California, San Diegon n source bioinformatics software platform for visualizing molecular interaction the sand integrating these interactions with gene expression profiles and other state ion et al., 2002). <i>iz (https://wiki.ncsa.ninc.edu/display/MAE/Home</i>) Managed by Jong Lee at NCSA is
Imber Cytosca in open Shant MAE Open-S	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegon source bioinformatics software platform for visualizing molecular interaction ks and integrating these interactions with gene expression profiles and other state on et al., 2002). iz (https://wiki.ncsa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based
Cl Cytosca in open etwo Shann MAE open-s the M	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegon source bioinformatics software platform for visualizing molecular interaction cks and integrating these interactions with gene expression profiles and other state ion et al., 2002). viz (https://wiki.ncsa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research.
Clumber Cytosca in openetwo Shann MAE open-s he Mi Favern	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegon source bioinformatics software platform for visualizing molecular interaction et al., 2002). iz (https://wiki.ncsa.ninc.edn/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. workbench (http://labora.org.uk) Developed by the myGrid team
Clamber Cytosca in ope intwo Shant WAE open-s ine Mi Cavern http://	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction ks and integrating these interactions with gene expression profiles and other state ton et al., 2002). iz (https://wiki.ncsa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://taverna.org.uk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free research for desiring and executing workflows (Hull at al. 2006). Towards allows of the set of the set of the university of Manchester, U.K. is a free research.
Imber Cytosca in open- Shant MAE Open-s he Mi Tavern http:// softwa	of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diego n source bioinformatics software platform for visualizing molecular interaction ks and integrating these interactions with gene expression profiles and other state ion et al., 2002). iz (https://wiki.ncsa.uiuc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://taverna.org.uk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows us prate many different software tools including over 30 000 web services
Imber Cytosca In open-s In open-s Shant MAE Shant MAE Shant Shant Cavern http:// softwa co inter	of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diego n source bioinformatics software platform for visualizing molecular interaction the and integrating these interactions with gene expression profiles and other state ion et al., 2002). iz (https://miki.ncsa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://laverna.org.nk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. irond (http://lowing.including over 30,000 web services.
Clamber Cytosca in open-s netwo Shann MAE Spen-s he Mi Cavern http:// coftw/ o inter TEXT	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://evtoscape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction et al., 2002). viz (https://wiki.ncsa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://taverna.org.uk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows us grate many different software tools, including over 30,000 web services. rend (http://textrend.org) Led by George Kampis at Eötvös Loránd University, Budz
Clamber Cytosca in openes betwo Shann MAE Shann Shan Sha	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://eutoscape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction et al., 2002). big (https://wiki.nesa.uiuc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://taverna.org.uk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. rend (http://textrend.org) Led by George Kampis at Eötvös Loránd University, Buda ry supports natural language processing (NLP), classification/mining, and graph nums for the analysis of business and governmental text corouses with an inherently
Imber Cytosca in ope Shan MAE Spen-s he Mi Copen-s in ope Softwa co inter TEXT Hunga Ilgorit cempo	Specify OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: Description be (<u>http://eutoscape.org</u>) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction the sand integrating these interactions with gene expression profiles and other state on et al., 2002). big (<u>http://wiki.ncsa.uiuc.edu/display/MAE/Home</u>) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (<u>http://Iarerna.org.uk</u>) Developed by the myGrid team (mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. big many different software tools, including over 30,000 web services. big many different software tools, including over 30,000 web services. big many different software tools, including over 30,000 web services. big many different software tools, including over 30,000 web services. big many different software tools, including over 30,000 web services. big many side processing (NLP), classification/mining, and graph nums for the analysis of business and governmental text corpuses with an inherently ral component.
Imber Cytosca In open- Shann MAE Spen-s he Mi Favern http:// softwa so inter TEXT Hunga Ilgorit cempo DynaN	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (<u>http://cytoscape.org</u>) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction the state interactions with gene expression profiles and other state on et al., 2002). big (<u>http://wiki.nesa.ninc.edu/display/MAE/Home</u>) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (<u>http://taverna.org.uk</u>) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. birnd (<u>http://textrend.org</u>) Led by George Kampis at Eötvös Loránd University, Budary supports natural language processing (NLP), classification/mining, and graph mus for the analysis of business and governmental text corpuses with an inherently ral component.
Imber Cytosca In open-s Detwo Shant MAE Open-s Che Mi Cavern http:// softwa co inte TEXT Hunga ilgorit cempo DynaN Amste	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction these and integrating these interactions with gene expression profiles and other state on et al., 2002). iz (https://miki.nesa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. tworkbencb (http://taverna.org.uk) Developed by the myGrid team (mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows ugrate many different software tools, including over 30,000 web services. trend (http://textrend.org) Led by George Kampis at Eötvös Loránd University, Buda ry supports natural language processing (NLP), classification/mining, and graph mus for the analysis of business and governmental text corpuses with an inherently ral component. Lets (http://numv.dynanets.org) Coordinated by Peter M.A. Sloot at the University of rdam, The Netherlands develops algorithms to study evolving networks.
umber Cytosca in open- netwo Shann MAE Shann MAE Shann MAE Shann Shann MAE Shann Shann MAE Shann MAE SISOE	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://citascape.org) Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction et al., 2002). iz (http://wiki.nesa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is on ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. a Workbench (http://taverna.org.uk) Developed by the myGrid team /mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. irend (http://textrend.org) Led by George Kampis at Eötvös Loránd University, Budz ty supports natural language processing (NLP), classification/mining, and graph mus for the analysis of business and governmental text corpuses with an inherently ral component. Vets (http://new.dynanets.org) Coordinated by Peter M.A. Sloot at the University of rdam, The Netherlands develops algorithms to study evolving networks.
Cytosca in open- betwo Shann MAE Softwa co inte TEXT Hunga ligorit co inte TEXT Hunga Softwa co inte TEXT Hunga higorit co inte SISOE he fur	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: be (http://cytoscape.org) Led by Trey Ideker at the University of California, San Dieger n source bioinformatics software platform for visualizing molecular interaction tcs and integrating these interactions with gene expression profiles and other state ion et al., 2002). iz (http://miki.nesa.ninc.edu/display/MAE/Home) Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. n/mygrid.org.uk) led by Carol Goble at the University of Manchester, U.K. is a freed re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. irend (http://textrend.org) Led by George Kampis at Eötvös Loránd University, Budary supports natural language processing (NLP), classification/mining, and graph nms for the analysis of business and governmental text corpuses with an inherently ral component. ets (http://num.dynanets.org) Coordinated by Peter M.A. Sloot at the University of rdam, The Netherlands develops algorithms to study evolving networks. (http://isab.lec.uma.es) An Observatory for Science in Society Based in Social Mode to clonality of OSGi-based software frameworks improves and the number and
umber Cytosca in ope Shan MAE Spen-s he Mi Copen-s in the Copen-s in the Copen-s in the Copen-s in the Copen-s in ope Shan MAE Shan MAE Softwa co inte Copen-s in the Copen-s in the in the	OSGi/CIShell Adoption of other projects recently adopted OSGi and/or CIShell: <i>pe (http://eviscape.org)</i> Led by Trey Ideker at the University of California, San Diegen source bioinformatics software platform for visualizing molecular interaction the and integrating these interactions with gene expression profiles and other state to en et al., 2002). <i>iz (http://wiki.nesa.nine.edu/display/MAE/Home)</i> Managed by Jong Lee at NCSA is ource, extensible software platform which supports seismic risk assessment based d-America Earthquake (MAE) Center research. <i>wWorkbench (http://tarema.org.uk)</i> Developed by the myGrid team /mygrid.org.uk) Ied by Carol Goble at the University of Manchester, U.K. is a free re tool for designing and executing workflows (Hull et al., 2006). Taverna allows u grate many different software tools, including over 30,000 web services. <i>rend (http://lextmd.org)</i> Led by George Kampis at Eötvös Loránd University, Buda ry supports natural language processing (NLP), classification/mining, and graph mus for the analysis of business and governmental text corpuses with an inherently camponent. <i>ets (http://imm.dynanets.org)</i> Coordinated by Peter M.A. Sloot at the University of rdam, The Netherlands develops algorithms to study evolving networks. (<i>http://imm.dynanets.org)</i> An Observatory for Science in Society Based in Social Mod ctionality of OSGi-based software frameworks improves and the number and f dataset and algorithm plugins increases, the cambilities of custom tools will experimental for the analysis increases.



scientific collaboration and sharing of data, software, and educational materials on the web. <u>http://hubzero.org</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://ivl.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.





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

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