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From: dbworld-bounces@cs.wisc.edu on behalf of Ruggero Pensa [rpensa@gmail.com]
Sent: Saturday, June 19, 2010 8:58 AM
To: dbworld@cs.wisc.edu
Subject: [Dbworld] DyNaK 2010: Deadline Extended

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Call for Papers:

Workshop on Dynamic Networks and Knowledge Discovery
(DyNaK 2010)
co-located with ECML PKDD 2010, September 24, 2010, Barcelona, Spain
<http://kdd.di.unito.it/DyNaK2010/>

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Aim and Scope

Modeling and analyzing networks is a major emerging topic in different research areas, such as computational biology, social science, document retrieval, etc. By connecting objects, it is possible to obtain an intuitive and global view of the relationships between components of a complex system.

Nowadays, the scientific communities have access to huge volumes of network- structured data, such as social networks, gene/proteins/metabolic networks, sensor networks, peer-to-peer networks. Most often, these data are not only static, but they are collected at different time points. This dynamic view of the system allows the time component to play a key role in the comprehension of the evolutionary behavior of the network (evolution of the network structure and/or of flows within the system). Time can help to determine the real causal relationships within, for instance, gene activations, link creation, information flow.

Handling such data is a major challenge for current research in machine learning and data mining, and it has led to the development of recent innovative techniques that consider complex/multi-level networks, time-evolving graphs, heterogeneous information (nodes and links), and requires scalable algorithms that are able to manage huge and complex networks.

DyNaK workshop is motivated by the interest of providing a meeting point for scientists with different backgrounds that are interested in the study of large complex networks and the dynamic aspects of such networks. It aims at attracting contributions from both aspects of networks analysis: large real network analysis and modelling, and knowledge discovery within those networks.

Even though each type of real complex networks has some peculiarities related to its specific domain, many aspects of the modeling and mining techniques for such networks are shareable. For instance, gene networks and social networks share a common architecture (scale-free), and involve similar data mining and machine learning methods: module/community extraction, hub single-out, information-flow analysis, missing link detection and link prediction.

Special session on Sentiment Analysis and Opinion Mining

Every day, millions of people write their opinions about any issue in social media, such as social news sites, review sites, and blogs. The distillation of knowledge from this huge amount of unstructured information is a challenging task. Sentiment Analysis and Opinion Mining are two areas related to Natural Language Processing and Text Mining that deal with the identification of opinions and attitudes in natural language texts. In the Opinion Mining session of DyNaK we are interested in research results from academics and practitioners in the task of extracting knowledge from user generated contents, and how time affects to this analysis.

Topics of interest

Contributions to the DyNaK workshop should be focused on this (non exhaustive) list of topics:

Methods:

- * Network inference from raw data
- * Graphical models
- * Graph mining algorithms
- * Graph kernel algorithms
- * Relational learning algorithms
- * Matrix/Tensor methods
- * Information retrieval algorithms
- * Bayesian methods
- * Evolutionary clustering
- * Mining and learning from heterogenous domains
- * Bisociative information discovery
- * Clustering/Co-clustering/Biclustering
- * Pattern mining and clustering with constraints
- * Community detection/Module extraction
- * Analogies between social and biological networks
- * Opinion Extraction and Classification
- * Blogs Analysis and Social Search
- * Temporal Sentiment Analysis
- * Irony and Plagiarism detection in Opinion Mining
- * Recommender Systems

Applications:

- * System biology: regulatory gene networks, protein-protein interaction, miRNA networks, metabolic networks
- * Social networks: folksonomies, digital libraries, information networks, social media, collaborative networks
- * Sensor networks, peer-to-peer networks, Web, agent networks, body sensor networks

Key Dates

- * Paper Submission due: June 28, 2010 (EXTENDED)
- * Notification of Acceptance: July 19, 2010 (EXTENDED)
- * Camera Ready Papers due: July 28, 2010 (EXTENDED)
- * Workshop: September 24, 2010

Organization

- * Ruggero G. Pensa, University of Torino, Italy
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Program Committee

- * Riccardo Bellazzi, University of Pavia, Italy
- * Guillaume Beslon, INSA-Lyon, France

- * Bettina Berendt, Katholieke Universiteit Leuven, Belgium
- * Tanya Berger-Wolf, University of Illinois, USA
- * Karsten M. Borgwardt, MPI, Tübingen, Germany
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- * Tamara G. Kolda, Sandia National Laboratories, USA
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- * Tao Li, Florida International University, USA
- * Pietro Liò, University of Cambridge, UK
- * Huan Liu, Arizona State University, USA
- * Eric Yu-En Lu, University of Cambridge, UK
- * Sara C. Madeira, INESC-ID/IST, Portugal
- * Rosa Meo, University of Torino, Italy
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- * Mirco Nanni, ISTI-CNR, Italy
- * Arlindo Oliveira, INESC-ID, Portugal
- * Andrea Passerini, University of Trento, Italy
- * Lorenza Saitta, University of Piemonte Orientale, Italy
- * Rossano Schifanella, University of Torino, Italy
- * Einoshin Suzuki, Kyushu University, Japan
- * Hanghang Tong, Carnegie Mellon University, USA

Invited Speakers

- * Tanya Berger-Wolf, University of Illinois, USA
- * Stefan Kramer, Technische Universität München, Germany
- * Carlos Rodríguez, Barcelona-Media, Spain

Sponsors

- * BioDigitalValley (<http://www.biodigitalvalley.com/>)
- * Computer Science Department, University of Torino (<http://kdd.di.unito.it/>)
- * SysBioM (http://www.biomolecular_complexsystems.unito.it/)

Paper Submission

We welcome original contributions, either theoretical or empirical, describing ongoing projects or completed work (even partially published). The instructions for authors and the LaTeX packages can be found at <http://www.springer.de/comp/lncs/authors.html>. Paper length should not exceed 12 pages. A selection of the accepted papers will be taken into consideration for a special issue of an international journal.

The instructions for paper submission are now available at <http://kdd.di.unito.it/DyNaK2010/>.

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