

Computational Epidemiology

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ABSTRACT

As recent pandemics such as SARS and the Swine Flu outbreak have shown, diseases spread very fast in today's interconnected world, making public health an important research area. Some of the basic questions are: How can an outbreak be contained before it becomes an epidemic, and what disease surveillance strategies should be implemented? These problems have been studied traditionally using differential equation methods, which are amenable to analysis and closed form solutions. However, these models are based on complete mixing assumptions, which do not hold for realistic populations, thereby limiting their utility.

In this tutorial, we focus on an approach based on diffusion processes on complex networks. This captures more realistic populations, but leads to novel mathematical and computational challenges. The structure of the underlying networks has a significant impact on the dynamical properties, motivating the need for improved network models, and efficient algorithms for computing network and dynamical properties that scale to large networks. We provide an overview of the state of the art in computational epidemiology, which is a multi-disciplinary research area, that overlaps different areas in computer science, including data mining, machine learning, high performance computing and theoretical computer science, as well as mathematics, economics and statistics. Specifically, we will discuss mathematical and computational models, problems of inference, forecasting and state assessment, and epidemic containment.

Target audience and prerequisites

The tutorial is suitable for both novice and expert researchers working in the areas of social and information networks, link prediction and public health. Problems of big data, including large scale optimization and novel statistical techniques are inherent aspects of this area. Public health is an area of great societal importance, with novel problems arising out of the use of network models. Recent advances in network modeling and agent based simulation tools, combined with

the use of new datasets, such as social media data, have led to new challenges and opportunities. The tutorial will introduce this emerging area, and will identify some of the key research topics. It will be at a level that is accessible to most KDD attendees who work in these areas.

Instructors

Madhav V. Marathe is a Professor of Computer Science and Director, Network Dynamics and Simulation Science Laboratory, Virginia Bioinformatics Institute, Virginia Tech. He obtained his Bachelor of Technology in 1989 in Computer Science and Engineering from the Indian Institute of Technology, Madras, and his PhD in 1994 in Computer Science from the University at Albany. He is the recipient of the Distinguished Copyright award for TRANSIMS software, LANL's achievement award, and a recipient of the University at Albany's Distinguished Alumni Award. His research interests are in high-performance computing, modeling & simulation of socio-technical systems, service oriented architectures, computer and communication networks, theoretical computer science, social networks and graph theory, computational epidemiology and computational economics. He is a fellow of the IEEE and ACM.

Anil Kumar S. Vullikanti is an Associate Professor in the Dept. of Computer Science and the Virginia Bioinformatics Institute at Virginia Tech. He received his undergraduate degree from the Indian Institute of Technology, Kanpur, and his Ph.D. from the Indian Institute of Science, Bangalore. He was a post-doctoral researcher at the Max-Planck Institute for Informatics, and a Technical Staff Member at the Los Alamos National Laboratory. His research interests are in the broad areas of approximation and randomized algorithms, dynamical systems, computation epidemiology, wireless networks, social networks, data mining and the modeling, simulation and analysis of socio-technical systems.

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