

Your Name: Li Zhang

Your E-mail address: lizhang@cse.buffalo.edu

Your Dissertation Proposal Title:

VIZSTRUCT: VISUAL EXPLORATION FOR

GENE EXPRESSION PROFILING

Your Committee:

Chair: Dr. Aidong Zhang

Members: Dr. Jian Pei, Dr. Murali Ramanathan

Your Dissertation Proposal Abstract:

In many applications, understanding data is of primary importance. Visualization may provide more insightful information than traditional numerical methods. Nevertheless, this is often quite difficult due to the complexity of the data and the limitation of human perspective. DNA microarray technology provides a broad snapshot of the state of the cell by measuring the expression levels of thousands of genes simultaneously. It has already had a significant impact on the field of bioinformatics and has proposed a unique challenge: information in gene expression matrices is special in that the sample space and gene space are of very different dimensionality and it can be studied in either sample space or gene space.

To meet this challenge, this proposal presents a theoretically solid visualization framework to explore gene expression datasets. Motivated by radial coordinate visualizations such as RadViz, we consider the two dimensional visualization problem as a problem of approximation of discrete-time signals. In this proposal, we propose a new paradigm of information visualization through frequency domain representation of signals. We introduce Fourier harmonic projections (FHPs) to map the high dimensional gene expression data onto a two dimensional space to create an intuitive scatterplot. FHPs have desired properties preserving crucial characteristics during the mapping. In this proposal, we present a detailed discussion and rigorous mathematical proofs of these properties. Furthermore, we design some possible enhancements and user interactions. Based on the unique nature of gene expression data, two novel and related processes are proposed, explorative visualization and confirmative visualization.