Chathura Gunasekara

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Education

2013 May -

PhD, Computational Science & Engineering; Michigan Technological University

2017 Spring

(Houghton, MI)

(expected) PhD Thesis title: Bioinformatics Tools and Algorithms Development for Gene Reg-

ulatory Network Inference

2006-2010 BS, Computational Physics; University of Colombo (Sri Lanka)

Major in Computer Science with minor in Physics and Applied Mathematics

Projects, Work Experience & Publications

Graduate Research Assistant: 2013 - Present

Currently I am Working under **Dr. Hairong Wei**, conducting research to infer gene regulatory networks and indentify regulatory transcription factors (TFs) which control known biological pathways in Arabidopsis thaliana under stress conditions using gene expression data.

- TF-miner
- My early attempts for this research was presented at NSF Project/Bioinformatics Workshop @ Noble Foundation, Ardmore, Oklahoma.
- Pairwise analysis of Pathway Transcription Factor gene expression data to find regulatory TF clusters.Git
- Currently implementing a web based gene expression data analysis pipeline to identify Transcription Factor(TF) clusters which associates with known biological pathways using novel pair-wise gene association methods such as Maximum Information Coefficient(MIC), Distance Correlation.

Completed projects

- Developed, implemented and published an algorithm and web application to search for degenerate motifs in the promoter regions of 50 plant species genomes. Our suffix tree based search algorithm can search for 100 degerate motifs in 35000 genes under 4 minutes!.
- I configured, installed and developed Perl scripts for parsing the FASTQ files using open sourse
 tools for a genome browser to visualize RNA-seq and Ribo-seq of wild-type and STTM mutants.
 This browser provides information on global translation status and miRNA-directed translation
 inhibition, which is currently lacking in plants. This web portal make them available to the research
 community.
- Co-authored an algorithm to infer hierarchical gene regulatory network from gene expression data.
- Implemented and Co-authored Poplar Gene Expression Pipeline web application.
- Contributor, to Birch Genome publication building a Circos Visualization

Software Engineer/Research Engineer: 2010 - 2013

Worked on a **collaborative research project** with University of Colombo School of Computing and Sri Lanka Navy. Following are the list of publications I authored/contributed:

- Develop algorithms and to implement using Java and web based technologies a Survailance platfrom to fuse data from multiple transponders such as AIS, RADAR sensors around Sri Lanken coast line. First-Author
- Maritime Navigation Simulator Project, First-Author CGAT 2012, Thailand, Low Cost 3D Immersive Telepresence for Survaillance, Planning, Maneuring: 3D-COP 10.5176/2251-1679_CGAT31.
 Computer Games, Multimedia & Allied Technology Conference 2012.
- Maritime Navigation Simulator Project, Co-Author ICter 2012 for Simulating Narrow Channel Effect on Surge Motion of a Ship in a Virtual Environment
- Undergraduate Research Conference paper, Spatialized Real Time Auditory Interface for a Virtual Maritime Application in 2010.

■ Technical Experience and Recent Course work

Data Science Applied Predictive Modelling Fall 2014

Introduction to Data Science Fall 2014

Data Mining Spring 2014

Data mining for geo spatial applications Fall 2015

Machine Learning - Regression Coursera Verified Certification (online)

Statistics Statistical Methods Fall 2013

Regression Analysis Spring 2013

Time series analysis and forcasting Spring 2015

Computer Advanced Scripting and Programming Fall 2015

Science Algorithmic Toolbox Coursera Verified Certification (online)

Bioinformatics Programming Skills Fall 2013

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