# Chathura Gunasekara

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## Education

2017 Spring (expected)

**PhD, Computational Science & Engineering**; Michigan Technological University (Houghton, MI)

Advisor: [Dr. Hairong Wei](http://www.mtu.edu/forest/about/faculty/wei/)

PhD Thesis title: Bioinformatics Tools and Algorithms Development for Gene Regulatory Network Inference

2010

**BS, Computational Physics**; University of Colombo (Sri Lanka)

Major in Computer Science with minor in Physics and Applied Mathematics

## Professional Experience & Publications

**Graduate Research Assistant : 2013 - Present**

*Curruent Projects*

* TF-miner is a data analysis pipeline based on partial least squares and graphical gaussian models to identify regulatory TFs using microarray gene expression data under stress conditions. [[source]](https://github.com/cjgunase/TF-miner)
* Infer gene regulatory network from DREAM5 Gene expression datasets [[source]](https://github.com/cjgunase/DREAM5_gene_net.git)
* Pairwise analysis of gene expression data to find regulatory TF clusters using novel association methods.[[source]](https://github.com/cjgunase/MIC_based_gene_expr_analysis)
* Currently implementing a web based gene expression data analysis pipeline to identify Transcription Factor(TF) clusters which associates with known biological pathways.[[source]](https://github.com/cjgunase/TF-Cluster), [[web]](http://sys.bio.mtu.edu/cluster/index.php)

*Completed projects*

* Developed an algorithm and web application to search for degenerate motifs in the promoter regions of 50 plant species genomes.[[source]](https://github.com/cjgunase/exactSearch), [[web]](http://sys.bio.mtu.edu/motif/), [[publication]](http://plantmethods.biomedcentral.com/articles/10.1186/s13007-016-0126-6)
* Configured, installed a JBrowse genome browser to visualize RNA-seq and Ribo-seq of wild-type and STTM mutants to parse fastq files.[[web]](https://blossom.ffr.mtu.edu/designindex2.php)
* Collaborated with a lab member to publish an algorithm to infer hierachical gene regulatory network from gene expression data.[[publication]](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4797117/)
* Collaborated with a lab member to develop the Poplar Gene Expression Pipeline web application. [[web]](http://sys.bio.mtu.edu), [[publication]](http://link.springer.com/article/10.1007/s11295-014-0745-x#page-1)
* Contributed to the publication of a lab member by creating a Circos visualization of genomics data from Birch genome.[[source]](https://github.com/cjgunase/myVisualizations/tree/master/circos)

**Software Engineer/Research Engineer : 2010 - 2013**

I Worked on a [research project](http://www.vidusayura.org/?page_id=2) with University of Colombo School of Computing and Sri Lanka Navy.

* Develop algorithms and implemented using Java and web based technologies for a Survailance platfrom to fuse data gathered from multiple transponders situated around Sri Lanken coast line over large local area network. [[Publication]](http://www.icter.org/conference/icter2012/paper/50)
* Maritime Navigation Simulator Project, Low Cost 3D Immersive Telepresence for Survaillance, Planning, Maneuring : 3D-COP 10.5176/2251-1679\_CGAT31. Computer Games, Multimedia & Allied Technology Conference 2012.[[Publication]](http://www.cgames.com.sg/PriorYearsPaper2012.html)
* Maritime Navigation Simulator Project for Simulating Narrow Channel Effect on Surge Motion of a Ship in a Virtual Environment. [[Publication]](http://www.icter.org/conference/icter2012/paper/31)
* Undergraduate Research on Spatialized Real Time Auditory Interface for a Virtual Maritime Application in 2010.[[Publication]](http://www.icter.org/conference/archive2011/index.php/icter/ICTer2010/paper/view/65)

## Technical Experience and Recent Course work

Programming

Software engineering (Object oriented programming, version control, Documentation)

Python (scikit-learn/numpy/scipy/pandas), R, Microsoft Excel

Perl, Java, C++, Database(SQL), Linux/Unix/Shell Scripting

Web Development (Linux/Apache/MySql/PHP) and web server management

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 Foundations: A Case Study Approach by University of Washington on Coursera [[verify]](https://www.coursera.org/account/accomplishments/certificate/UALSGBFJHK7W)

Machine Learning: Regression by University of Washington on Coursera [[verify]](https://www.coursera.org/account/accomplishments/certificate/2AXNSM86GTZS)

Statistics

Statistical Methods (Fall 2013), Regression Analysis (Spring 2013)

Time series analysis and forcasting (Spring 2015)

Algorithms

Advanced Scripting and Programming (Fall 2015)

Algorithmic Toolbox by University of California, San Diego & Higher School of Economics on Coursera.[[verify]](https://www.coursera.org/account/accomplishments/certificate/KSF3Q7M96AWB)

Biology

Bioinformatics Programming Skills (Fall 2013)

Following online Genomic Data Science Specialization Johns Hopkins University

## Professional Development

2016

Presentation at NSFproject/Bioinformatics workshop at Noble Foundation, Ardmore, OK on Gene regulatory network prediction using partial least squares and bayesian network methods.[[presentation]](http://www.slideshare.net/secret/15sqXSMbpOHAOQ)

2016

Poster presentation at the research symposium of Life Science and Technology Institute, Michigan Tech