# Beethika Tripathi Indian Institute of Technology Madras

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# Education

Program	Institution	%/CGPA	Year of Completion
M.S. (Computer Science and Engg.)	Indian Institute of Technology Madras	8.6	2018
M.Tech (Computer Science and Engg.)	Birla Institute of Technology, Mesra	8.75	2015
B.Tech (Computer Science and Engg.)	SRMCEM, UPTU, Lucknow	78.5	2013
XII (ISC)	Saint Dominic Savio College, Lucknow	83.7	2009
X (ICSE)	Saint Dominic Savio College, Lucknow	88.0	2007

## **Technical Skills**

• Languages: Proficient - Java, C, Python. Intermediate - HTML, CSS, AWK

- o Tools: MATLAB, LATEX, Visual Studio, Eclipse, Code Blocks, Weka, Gephi, SQL Plus
- Databases: Oracle 10g, DB2, MySQL

## **Course Work**

• Data Science: Machine Learning, Natural Language Processing, Data Mining, Searching & Indexing in Large Dataset

- Computer Science: Advanced Data Structures and Algorithms, Operating Systems, Database, Networks
- Mathematics: Linear Algebra and Random Processes, Probability & Statistical Analysis

# **Publications**

- B.Tripathi and Amritanjali, "*Finding sorting traces of reversals in the presence of hurdles.*" Proceedings of the 9th International Conference on Bioinformatics and Biomedical Engineering 2015. (iCBBE 2015), pp. 85–90.
- B.Tripathi and Amritanjali, "An improved algorithm for reducing the solution space for sorting by reversals." International Journal of Basic and Applied Biology, pp. 23-27.

# Key Projects

#### **Neural representation learning for biological networks** *Guide: Dr Balaramn Ravindran and Dr Karthik Raman*

- Leveraging the power of complex network analysis to bypass exhaustive characterizing of genes through biological intractable experiments.
- Neural representation learning techniques employed to encode complex relations in a continuous vector space.
- For reliable inferring in case of noisy networks, developing a principled approach for integrating heterogeneous sources of information to learn shared representations across multiplex networks.

### Disease module identification

Guide: Dr Balaraman Ravindran, Dr Karthik Raman and Dr Himanshu Sinha

• Employed graph theoretic and machine learning techniques to exploit the basic assumption of "guilt by association", where colocalized genes are assumed to be functionally correlated hence responsible for a disease.

### Analysis of NLP techniques for protein family classification

Faculty: Dr Sutanu Chakraborti, Natural Language Processing, Team Size:3

- Continuous space representation models such as word2vec, performs better than traditional statistical models such as statistical language model that rely on discrete features.
- Achieved F-score of 0.98 for protein family classification in comparison to discrete bag of words model (F-score 0.82).

### Jun 2016 - Present M.S. Project, IIT Madras

# Sept - Nov 2016

DREAM Challenge

### Course Project, IIT Madras

Jun 2016 - Dec 2016

#### Spell-checker Faculty: Dr Sutanu Chakraborti, Natural Language Processing, Team Size:3

• Predict corrections to erroneous words at 3 levels namely standalone words, phrases and sentences, based on a noisy channel model, phonetics and cross entropy based N-gram model to account for the context of the word.

# Semantically matching documents using term co-occurrence graph

Faculty: Dr Sayan Ranu, Searching & Indexing in Large Dataset, Team Size:2

- Quantified document matching using the hidden semantic meaning instead of standard lexical features using point mutual information (PMI), TF-IDF and cosine similarity.
- Created a term co-occurrence graph using 8.21GB of clean Wikipedia text and tested on Flipkart review corpus.

### Efficient solution to the problem of sorting genomes by reversals Guide: Dr Amritanjali

- Comparing genomes of two different species to reconstruct evolutionary history by finding out sequence of the possible genome rearrangement (reversal) operations that could transform the shared gene order into another genome.
- Developed heuristics for grouping into equivalence classes to reduce the solution space, as it increases exponentially with the size of shared genes. Resulting in 10 times speedup.

# Emotion based music player

Guide: Dr Radhey Shyam, Team Size:2

- o Image of the person localized to find the region of interest. Important features extracted using principle component analysis. Support Vector Machine used for classification into four emotions namely happy, anger, disgust and neutral.
- Detected emotion used as input to music player to play the desired playlist.

# **Scholastic Achievements**

- Scholar of Grace Hopper Conference India 2016.
- Won  $2^{nd}$  prize in Data Challenge organized by Visa at GHCI 2016.
- Secured **1**<sup>st</sup> rank worldwide in the sub-challenge of disease module identification **DREAM Challenge** 2016.
- Obtained certificate of merit, outstanding performance in all 4 years B-Tech.

# **Positions of Responsibility**

- Faculty Scholarship Application Reviewer at Grace Hopper Conference 2017
- Mentored interns at IIT Madras.(May June, 2017)
  - Assisted in deciding problem for B.Tech project.
  - Designed project framework.
- Teaching Assistant for Machine Learning, Deep Learning & Introduction to Programming courses at IIT Madras.
  - Prepared tutorial questions and programming assignments.
  - Evaluated guizzes and conducted vivas.

# Co-Curricular and Extra-Curricular Activities

- Coding Competitions: Current InterviewBit global ranking 4194/102724 (99.95 percentile) as on 6 September 2017.
- $\circ$  Presented posters in various workshops at IIT Madras, University of Hyderabad & TCS Research. pprox 4
- Participated in Android Application Development Workshop, Tarang 2012, SRMCEM Lucknow.
- Volunteer in cultural events at annual fest Abhivyakti 2012, SRMCEM Lucknow.
- Dancing: Trained in Bharatnatyam for 5 years from Prayag Sangeet Samiti, Allahabad.

Course Project, IIT Madras

#### Jul 2014 - Apr 2015 M.Tech Project, BIT Mesra

# Jan - May 2015

B. Tech Project, SRMCEM Lucknow

Course Project, IIT Madras

Aug - Sept 2016

Jul - Nov 2015