# **RESEARCH INTERESTS**

Computational Biology; Network Representation Learning; Deep Learning

## **CURRENT POSITION**

2017 Rochester Institute of Technology, Rochester, NY

Graduate Research Assistant, Golisano College of Computing and Information Science

Advisor: Prof. Anne Haake, Prof. Rui Li

### **EDUCATION**

Rochester Institute of Technology, Rochester, NY

2016 -Doctor of Philosophy, Computing and Information Science

Tribhuwan University, Pulchowk Engineering Campus, Lalitpur, Nepal

2011 - 2014**Bachelor of Engineering (B.E.)**, Computer Engineering (GPA: 4.0)

Institute of Engineering: Pulchowk Campus, Tribhuwan University

Thesis: Agricultural Data Integration and Analysis - Analyzing environmental factors

to recommend appropriate crops for a place.

### PROFESSIONAL EXPERIENCE

2016 -Rochester Institute of Technology, Research Assistant

ABI Innovation - Novel Methodology for Leveraging Metabolic Simulation to Improve

Regulatory Reconstruction.

Advisor: Prof. Anne Haake, Prof. Rui Li

2015 - 2016Verisk Information Technologies, Data Engineer

Analyzed US Healthcare (medical and pharmacy claims) to provide insights about

people's health.

2014 - 2015Yomari Pvt. Ltd., Software Trainee

Developed ETL framework to load retail data into data warehouse.

2013 E&T Nepal Pvt. Ltd., Intern

Developed web application that provides user friendly interface to simulate CFD results

in 3D models and share among users.

RIT Ph D. Merit Scholarship

## **HONORS & AWARDS**

2017

2017	KIT THE SCHOOL SHOP
2016	Verisk Way to Go Award (Best Employee, Verisk Information Technologies)
2015	Rookie of the Year (Verisk Information Technologies)
2014	The College Fellowship Scholarship (6 times during <b>B.E.</b> )
2014	Academic Excellence Award (8 times during <b>B.E.</b> )

Academic Excellence Award (8 times during **B.E.**)

## RECENT ACADEMIC PROJECTS

2017 Learning Hidden Representation of Gene Expression Data.

Analyzed gene expression data to see if they provide clues about the regulatory relationship between genes. Implemented autoencoder architecture to learn hidden representation of expression data. learned representation is visualized with tSNE.

Independent Study (Deep Learning), Spring 2017

2017 Ensemble SVM for Reconstruction of Gene Regulatory Network.

Developed ensemble feature selection approach based on support vector machines to select the subset of transcription factors (genes) that have huge impact in the expression

level of target genes. Implemented multiprocessing to speed up the execution.

Foundations of CyberInfrastructure Foundations, Spring 2017

2016 Predicting the outcome of soccer matches.

Investigated various features related to soccer match to design a framework to predict

the outcome of soccer matches. *Introduction to Big Data, Fall 2016* 

### TALKS AND POSTERS

2017 Reconstruction of Gene Regulatory Network using Ensemble SVM with Recursive

Feature Elimination

[Poster] GCCIS Research Showcase, Rochester Institute of Technology

## **TECHNICAL SKILLS**

Programming Languages Python, R, Java, C/C++, MATLAB.

Markup Languages & Web HTML/5, CSS, Javascript, Java Spark Framework, PHP, Shiny.

Databases & Query Languages SQL, MySQL, PL/SQL, Oracle.

Other GitHub, SVN, Scrum Agile Development.