Kishan K C

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SUMMARY

- I develop machine-learning algorithms to model, predict and explain biological data.
- I design, implement, and train machine-learning models, usually in scikit-learn (python) and PyTorch.
- I have a strong background in probabilistic machine learning, deep learning, graph neural networks, network biology, and interpretability.

EDUCATION

Rochester Institute of Technology, Rochester, New York

August 2016 - Present

PhD in Computing and Information Sciences. GPA 4/4

Relevant coursework: Statistical Machine Learning; Deep Learning;

Big data; Statistical Analysis

Institute of Engineering, Tribhuvan University, Kathmandu, Nepal

January 2010 – October 2014

B.E. in Computer Engineering. Passed with Distinction

Relevant Courses: Artificial Intelligence; Data Mining; Big Data

Technologies; Computer Programming; Data Structures and Algorithms;

Probability and Statistics

RESEARCH EXPERIENCE

Graduate Research Assistant, Lab of Use-Inspired Computational Intelligence

Rochester Institute of Technology, Rochester, NY

August 2016 – Present

- Developed higher-order graph convolutional networks (github.com/kckishan/HOGCN-LP) on PyTorch to aggregate information from multi-hop neighborhood for link prediction
- Developed interpretable and sparse gated sequence encoder (github.com/kckishan/InterpretablePIP) on PyTorch tailored to biological sequences
- Designed representation learning framework (github.com/kckishan/GNE) on TensorFlow to integrate biological networks with additional node information for link prediction

INDUSTRY EXPERIENCE

Data Engineer, Research & Development, Project: Medical Intelligence

Verisk Information Technologies, Kathmandu, Nepal

May 2015 – June 2016

- Developed machine learning model on scikit-learn to predict the execution time of SQL queries for optimization
- Collaborated with a team of 10 to develop a framework for norm processing and received the team
 of the quarter award
- Awarded two individual awards for outstanding contributions to the projects and workplace

SKILLS

Data Science tools NumPy, Pandas, SciPy, Matplotlib, NetworkX, Jupyter

Deep Learning Frameworks PyTorch, TensorFlow

Programming Languages Python, R

Machine Learning libraries Scikit-learn (python), Caret (R)

Databases SQL

SELECTED PUBLICATIONS

- 3. **K C K.**, Li R., Cui F., Haake A. "Predicting biomedical interactions with higher-order graph convolutional networks." *IEEE/ACM Transactions on Computational Biology and Bioinformatics*. Feb 2021.
- 2. **K C K.**, Cui F., Haake A., Li R. "Interpretable Structured Learning with Sparse Gated Sequence Encoder for Protein-Protein Interaction Prediction, *25th International Conference on Pattern Recognition*, Jan 2021.
- 1. **K C K.**, Li R., Cui F., Yu Q., Haake A. "GNE: A deep learning framework for gene network inference by aggregating biological information" *The Asia Pacific Bioinformatics Conference*. Jan 2019.

HONORS, AND AWARDS

| RIT PhD Merit Scholarship | 2016-Present |
|---|--------------|
| RIT Graduate Showcase Oral Presentation Award | 2019 |
| Team of the Quarter | 2017 |
| Data Science Certification Grant | 2016 |
| The employee of the Year | 2016 |
| Rookie of the Year | 2015 |