NAM D. NGUYEN

Machine Learning Researcher · Software Engineer

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EXPERIENCE

RESEARCH FELLOW

Daifeng Wang Laboratory, University of Wisconsin-Madison

Sep 2019 - Present

O LISA

- Researched multi-view learning & its application to brain disorder
- Developed Varmole, ECMarker; authored related papers; presented at ISMB 2020 & RSG 2020, resulting in favorable feedback and recommendation for inclusion in the next phase of the project

RESEARCH ASSISTANT

SBU-BNL SEED Grants "Large-Scale Comparative Regulatory Network Analysis in Photosynthetic Organisms"

Daifeng Wang Laboratory, Stony Brook University

May 2017 - Aug 2019

Q USA

- Collaborated with Brookhaven National Lab. to collect & analyze RNA-Seq data of Chlamydomonas reinhardtii
- Researched manifold alignment & its application to multi-view learning
- Developed ManiNetCluster; authored a related paper; presented at ACM-BCB 2018, RSGDREAM 2018 & ICIBM 2019, resulting in favorable feedback and recommendation for inclusion in the next phase of the project
- Solved the trade-off between parametric and nonlinear manifold alignment by designing a deep architecture; implemented in PyTorch

Other Projects

- Led a team in 2 prediction competitions, Zillow Prize & Online Dating Matchmaking (top 22%)
- Led the project "Automated Singing Evaluation" to train the machine to distinguish between good & poor singing

RESEARCHER

Software Engineering Lab., Pohang University of Science & Technology

Mar 2011 - May 2013

♀ S.Korea

SELECTED PUBLICATIONS

(*co-first-author)

- [1] Ting Jin, Nam D Nguyen*, Flaminia Talos, and Daifeng Wang. "ECMarker: Interpretable machine learning model identifies gene expression biomarkers predicting clinical outcomes and reveals molecular mechanisms of human disease in early stages". In: Bioinformatics (Nov. 2020). btaa935. ISSN: 1367-4803.
- [2] Nam D Nguyen, Ian K Blaby, and Daifeng Wang. "ManiNetCluster: a novel manifold learning approach to reveal the functional links between gene networks". In: *BMC genomics* 20.12 (2019), pp. 1–14.
- [3] Nam D Nguyen, Ting Jin, and Daifeng Wang. "Varmole: A biologically drop-connect deep neural network model for prioritizing disease risk variants and genes". In: *Bioinformatics* (Oct. 2020). btaa866. ISSN: 1367-4803.
- Nam D Nguyen and Daifeng Wang. "Multiview learning for understanding functional multiomics". In: *PLOS Computational Biology* 16.4 (2020), e1007677.

HONORS & AWARDS

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NSF Travel Award ICIBM 2019, USA



Best Poster "A Manifold Learning Approach to Reveal Functional Linkage Across Gene Networks"

ACM-BCB 2018, USA



VEF Doctoral Fellowship

The US National Academy of Sciences 2016, USA



Research Scholarship

National Research Foundation of Korea 2011, S.Korea

SKILLS & LANGUAGES

Python PyTorch R MATLAB

Bash AWK Vim Git

Java UML OOAD

Design Patterns

RESEARCH INTERESTS

Machine Learning Network Science

Computational Biology Optimization

EDUCATION

PhD Candidate in Computer Science Stony Brook University

Relevant Coursework: Machine Learning

- · Convex Optimization · Data Science
- · Algorithm

BEng in Computer Science Hanoi University of Science & Technology

2005 - 2010

♀ Vietnam GPA 3.16/4.00

Relevant Coursework: Linear Algebra
Probability & Statistics Numerical
Methods Digital Signal Processing

· Software Design & Development

IBM Certified Solution Designer
Application Developer DB2
Coursera Certified Game Theory

Social Network Analysis