

Seojin Bang

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PROFILE

Broad research experience in machine learning, deep learning and software development. Graduate researcher in CS with math/stat backgrounds.

EDUCATION

Aug 2020 (expected)	The School of Computer Science, Carnegie Mellon University PhD candidate in Computational Biology	Advisor: Wei Wu
Feb 2013	Seoul National University, Korea MS in Statistics	Advisor: Taesung Park
Aug 2010	Sungkyunkwan University, Korea BS in Mathematical Education · BE in Statistics	

RESEARCH EXPERIENCE

Interpretability and Robustness in Machine Learning Jul – Aug 2019
Visiting Graduate Student advised by Dr. Adrian Weller *ML Group at University of Cambridge, UK*

- Investigated the relationship between interpretability and of machine learning and deep learning models.
- Developing a knowledge guided interpretable ML model using the posterior regularization. (in progress)

Interpretable Machine Learning May – Dec 2018
Research Intern advised by Dr. Pengtao Xie *AI/ML Solution Team at Petuum, Pittsburgh PA*

- Developed a system-agnostic interpretable ML approach using neural networks using information bottleneck principle.
- Developed a pytorch based software: **VIBI** (github.com/SeojinBang/VIBI)
- A paper submitted to *ICLR*

Multimodal/Multiview Machine Learning 2015 – 2019
Research Assistant advised by Dr. Wei Wu *Carnegie Mellon University, Pittsburgh PA*

- Developed a multiple kernel k-means clustering approach that is robust against adversarial features and identified variable asthma subtypes by combining multiview clinical data.
- Developed an R-package: **MKKC** (github.com/SeojinBang/MKKC)
- Three papers published in *AJRCCM*, *IEEE BIBM* and *Proteomes*, and a paper submitted to *AAAI*.

Other Collaborative Experience with Graduate Colleagues 2017 – 2019
Graduate Student *Carnegie Mellon University, Pittsburgh PA*

- Developed an ontology-based neural network model for patient need detection. A paper in progress.
- Developed a phased-LSTM based predictive model for EHRs with complex missing patterns. Released a preprint.

Statistical Modeling of Sparse Gene Networks 2013 – 2015
Research Assistant advised by Dr. Haim Bar *University of Connecticut, Storrs, CT*

- Developed a statistical mixture model for better estimating sparse gene network.
- Helped to develop an R-package: **edgefinder**
- A paper under review in *Biostatistics*.

Statistical Approaches for Biomedical Problems 2012 – 2013
Research Assistant and Scientist advised by Dr. Taesung Park *Seoul National University, Korea*

- Developed a joint feature selection method using the elastic-net regularization to high-dimensional data.
- Deployed a time-dependent survival model to identify subtypes of intraductal papillary mucinous neoplasm.
- Two papers published in *IEEE BIBM* and *Pharmacogenetics and genomics*.

OTHER EXPERIENCE

Professional Experience

2019	Reviewer of <i>NeurIPS ML4H Workshop</i> , <i>IEEE Access</i> , and <i>IEEE TNNLS</i>
2018	Reviewer of <i>ACM BCB</i>
2018	Admission Committee at Computational Biology Department, Carnegie Mellon University
2012	Program Committee of two workshops held at Seoul National University and a conference of the Korean Statistical Society
2012 – 2018	Teaching Assistant at Carnegie Mellon University, University of Connecticut and Seoul National University

Leadership Experience

2017 – 2018	Led the communication team in Korean Graduate Student Association to establish and maintain relationship with ~30 companies seeking to advertise events/job openings.
2016 – 2019	Led the squash team of 35 CMU students. Initiated and organized weekly training. Applied and get a financial support from KGSA.

TECHNICAL STRENGTHS

Computer Languages	Python, R, C/C++, MATLAB, Bash, HTML
Library & Others	Pytorch, Tensorflow, Caffe, Keras, Git, LaTeX

HONORS AND AWARDS

2018	The Center for Machine Learning and Health Fellowships in Digital Health
2013	The Korean Statistical Society Paper Awards (3st Place)
2012	The Korean Statistical Society Poster Awards (1st Place)
2006 – 2010	National Science and Engineering Undergraduate Scholarship

SELECTED PAPERS

- [1] **Explaining a black-box using deep variational information bottleneck approach.**
preprint *arXiv:1902.06918*, (submitted to *ICLR*), 2019.
Bang, Seojin and Xie, Pengtao and Lee, Heewook and Wu, Wei and Xing, Eric.
- [2] **Robust multiple kernel k-means clustering using min-max optimization.**
preprint *arXiv:1803.02458*, (submitted to *AAAI*), 2019.
Bang, Seojin and Yu, Yaoliang and Wu, Wei.
- [3] **Phased-lstm based predictive model for ehers with complex missing patterns.**
preprint, <https://www.cs.cmu.edu/~epxing/Class/10708-17/project-reports/project8.pdf>, 2019.
Bang, Seojin and Yang, Yang and Wang, Yuchuan.
- [4] **Multiview cluster analysis identifies variable corticosteroid response phenotypes in severe asthma.**
American Journal of Respiratory and Critical Care Medicine, 2019.
Wu*, Wei and Bang* (co-first), Seojin and Bleecker, Eugene and Castro, Mario and Denlinger, Loren and Erzurum, Serpil and Fahy, John and Fitzpatrick, Anne and Gaston, Ben and Hastie, Annette and Israel, Elliot and Jarjour, Nizar and Kerr, Sheena and Levy, Bruce Meyers, Deborah and Moore, Wendy and Peters, Michael and Phipatanakul, Wanda and Sorkness, Ronald and Wenzel, Sally.