# Seojin Bang

#### RESEARCH INTEREST

My research interest is in the interpretability of black-box machine learning models. I develop interpretable machine learning approaches and investigate the relationship between interpretability and robustness in machine learning algorithms. I also develop multimodal/multiview machine learning approaches for combining different data modalities.

#### **EDUCATION**

2020	The School of Computer Science, Carnegie Mellon University	
	PhD in Computational Biology	Advisor: Wei Wu
2013	Seoul National University, Korea MS in Statistics	Advisor: Taesung Park
2010	Sungkyunkwan University, Korea	Advisor. Taesung Lark
2010	BS in Mathetmatical Education · BE in Statistics (double major)	

# PUBLICATIONS/PREPRINTS

[1] Explaining a black-box using deep variational information bottleneck approach. *AAAI*, 2021.

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, 2020.

Bang, Seojin and Yu, Yaoliang and Wu, Wei.

[3] Kw-han: Knowledge infused hierarchical attention network for enhancing interpretability. preprint, under review, 2020.

Jang, Hyeju and Bang, Seojin.

[4] Detecting attackable sentences in argument.

EMNLP, 2020.

Jo, Yohan and Bang, Seojin and Manzoor, Emaad and Hovy, Eduard and Reed, Chris.

[5] Identification of epitope-tcr binding using a generative adversarial network model.

ICML WCB, 2020.

Bang, Seojin and Lee, Heewook.

[6] Dropout prediction over weeks in moocs via interpretable multi-layer representation learning. *AAAI AI4EDU*, 2020.

Jeon, Byungsoo\* and Park, Namyong\* and Bang, Seojin\* (co-first).

[7] Phased-lstm based predictive model for ehrs 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.

[8] A mixture model to detect edges in sparse co-expression graphs.

Preprint arXiv:1804.01185, 2019.

Bar, Haim and Bang, Seojin.

[9] Multiview cluster analysis identifies variable corticosteroid response phenotypes in severe asthma.

American Journal of Respiratory and Critical Care Medicine, (IF 16.49) Highlighted Articles, 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.

[10] Phosphoproteomic analysis of the amygdala response to adolescent glucocorticoid exposure reveals g-protein coupled receptor kinase 2 as a target for reducing motivation for alcohol.

Proteomes, 6(4), 2018.

Bertholomey, Megan L. and Stone, Kathryn and Lam, TuKiet T. and **Bang**, **Seojin** and Wu, Wei and Nairn, Angus C. and Taylor, Jane R. and Torregrossa, Mary M.

[11] Naïve bayes ensemble: A new approach to classifying unlabeled multi-class asthma subjects. In *IEEE International Conference on Bioinformatics and Biomedicine*, 2016.

Bang, Seojin and Wu, Wei.

[12] Joint selection of snps for improving prediction in genome-wide association studies. In *IEEE International Conference on Bioinformatics and Biomedicine. Workshops*, 2012. Bang, Seojin and Kim, Yong-Gang and Park, Taesung.

[13] Ethnic variability in the allelic distribution of pharmacogenes between korean and other populations.

Pharmacogenetics and genomics, 22(12), 2012.

Kim, In-Wha and Im Kim, Kyung and Chang, Hyeu-jin and Yeon, Bora and **Bang**, **Seojin** and Park, Taesung and Kwon, Ji-sun and Kim, Sangsoo and Oh, Jung Mi.

## RESEARCH EXPERIENCE

#### Multimodal/Multiview Machine Learning for Asthma Subtype Identification

2015 - 2020

Research Assistant advised by Dr. Wei Wu

Carnegie Mellon University, Pittsburgh, PA, USA

- · Developed a robust multiple kernel clustering approach.
- · Identified asthma subtypes showing differential responses to corticosteroid 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 [blinded].

### Interpretable Machine Learning

2018 - 2020

Research Intern advised by Dr. Pengtao Xie

AI/ML Solution Team at Petuum, Pittsburgh, PA, USA

- · (Worked as an intern at Petuum between May and Dec 2018. Worked as a student at Carnegie Mellon University from 2019.)
- · Developed a system-agnostic interpretable ML approach using neural networks using information bottleneck principle.
- · Improved an out-of-sample prediction of epitope-TCR binding using an interpretable model.
- · Developed a pytorch based software: VIBI (github.com/SeojinBang/VIBI) and TCR (github.com/SeojinBang/TCR)
- · A paper submitted to [blinded]

#### Machine Learning for Natural Language Processing

2018 - 2020

 $Graduate\ Student$ 

Carnegie Mellon University, Pittsburgh, PA, USA

- · Developed an ontology-based neural network model for patient need detection from an online ovarian cancer discussion forum. A paper submitted to [blinded].
- · Analyzed to detect attackable sentences in arguments for successful persuasion using online discussions from the Change-MyView (CMV) subreddit. A paper submitted to [blinded].

#### Interpretability and Robustness in Machine Learning

2019

- · Investigated the relationship between interpretability and robustness of machine learning and deep learning models.
- · Developing a knowledge guided interpretable ML model using the posterior regularization.

#### Statistical Modeling of Sparse Gene Networks

2013 - 2015

Research Assistant advised by Dr. Haim Bar

University of Connecticut, Storrs, CT, USA

- $\cdot$  Developed a statistical mixture model for better estimating sparse gene network.
- · Helped to develop an R-package: edgefinder
- · A paper under review in Statistics in Medicine.

#### 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.

# PROFESSIONAL SERVICE

2020	Reviewer, NeurIPS, NeurIPS ML4H Workshop, ACM CHIL, ISMB
2019	Reviewer, NeurIPS ML4H Workshop, IEEE Access, IEEE TNNLS
2018	Reviewer, ACM BCB
2018	Admission Committee, Computational Biology Department, Carnegie Mellon University
2012	Program Committee, International Symposium on Statistical Genetics, Korea
2012	Program Committee, Microarray Analysis Workshop: Statistical Analysis using R
	language, Korea
2011	Program Committee, The Spring Conference of the Korean Statistical Society, Korea

#### HONORS AND AWARDS

0010		· · · · · · · · · · · · · · · · · · ·
2018	The Center for Machine Learning and Health Fellowships in Di	igital Health
	full tuition and stipend for 12 months and research-related expenses (total s	\$75,200)
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	
	full tuition for 8 semesters	2006 - 2010
	an additional \$500 grant for a high GPA	2009
	an additional \$500 grant for a high GPA	2008

#### SOFTWARE

Python	VIBI: pytorch implementation of VIBI
	https://github.com/SeojinBang/VIBI
Python	TCR: TCR to epitope binding prediction approach using VIBI
	https://github.com/SeojinBang/TCR
R-package	$\mathbf{MKKC}$ : multiple kernel $k$ -means clustering on a multi-view data
	https://github.com/SeojinBang/MKKC
LaTeX template	TidyCV: simple and tidy LaTeX template for your curriculum vitae
	https://github.com/SeojinBang/TidyCV

# TECHNICAL STRENGTHS

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

# TEACHING EXPERIENCE

gie Mellon University	Teaching Assistant
Quantitative Cell and Molecular Biology Lab	
Computational Methods for Proteogenomics and Metabolomics	
sity of Connecticut	Teaching Assistant
Mathematical Statistics	
Introduction to Mathematical Statistics	
Elementary Concepts of Statistics	
Introduction to Statistics I and II	
Statistical Methods	
National University, Korea	Teaching Assistant
Statistics Laboratory	
Regression and Analysis and Laboratory	
Statistics	
oung Girls' Middle School, Korea	Student Teacher
Middle School Mathematics	
Sungkyunkwan University, Korea Teach	
Alternative Elementary/Middle School Mathematics	
	Quantitative Cell and Molecular Biology Lab Computational Methods for Proteogenomics and Metabolomics sity of Connecticut Mathematical Statistics Introduction to Mathematical Statistics Elementary Concepts of Statistics Introduction to Statistics I and II Statistical Methods National University, Korea Statistics Laboratory Regression and Analysis and Laboratory Statistics Oung Girls' Middle School, Korea Middle School Mathematics Wunkwan University, Korea