

SungHwan Kim

Department of Applied Statistics
Konkuk University

Office: (+82) 02 - 450 - 3658
Email: shkim1213@konkuk.ac.kr

Education

Ph.D. Biostatistics, University of Pittsburgh, Pittsburgh, Apr 2015.

M.S. Statistics, Korea University, Seoul, Feb 2010.

B.A. Education, Korea University, Seoul, Feb 2007.

Experiences

Assistant Professor, Department of Applied Statistics, Konkuk University, Mar 2018 - Present.

Assistant Professor, Department of Statistics, Keimyung University, Mar 2017 - Feb 2018.

Research Grants

[1] "Action recognition models guided by semantic segmentation with the application to monitoring recyclable waste", *National Information Society Agency*, 125M KRW, Jan 2021–Dec 2021 (중소벤처기업부, 총 1억4000만원/12개월).

[2] "Building AI datasets for training predictive models", *National Information Society Agency*, 125M KRW, Sep 2020–Dec 2020 (한국정보화진흥원, 총 1억2500만원/4개월).

[3] "Untact identity verification via facial recognition technique paralleled to vision telecommunication", *Small and Medium Business Administration*, 144M KRW, Sep 2020–Sep 2022 (중소벤처기업부, 총 1억4400만원/24개월).

[4] "Statistical artificial intelligence (AI) algorithms to monitor deleterious and infinitesimal environment materials (particulates and microplastics)", *National Research Foundation of Korea Grant*, 150M KRW, Mar 2020–Feb 2023 (한국연구재단, 총 1억5,000만원/36개월).

[5] "Deep learning based image processing techniques to develop a fine dust metering device", *Small and Medium Business Administration*, 300M KRW, Jan 2020–Dec 2020 (중소벤처기업부, 총 3억원/12개월).

[6] "Techniques to meter particulate matter levels using temporal digital image data", *Research Institute of Industrial Science and Technology (RIST)*, 20M KRW, Dec 2019–Feb 2020 (포스코, 총 2,000만원).

[7] "Integration methods of multi-level omics data via bigdata analytics", *National Research Foundation of Korea Grant*, 90M KRW, Mar 2017–Feb 2020 (한국연구재단, 총 9,000만원/36개월).

[8] "Development of statistical methodologies for integration of multi-omics data", *National Research Foundation of Korea Grant*, 34M KRW, Sep 2016–Aug 2017 (한국연구재단, 총 3,400만원/12개월).

Patents

[1] **Kim S.** and Heo J., "Method for Analyzing Fine Dust and Apparatus for Executing the Method", No. 10-2018-0021631, Feb 2019, *Korean Intellectual Property Office*, Registered.

Conference Proceedings

[1] Han J., Kim G., Lee C., Hwang U., Han Y. and **Kim S.** (2019) Predictive Models of Fire via Deep learning Exploiting Colorific Variation, *ICAIIIC 2019 Conference Proceedings*.

[2] Kim G., **Kim S.**, Kim J. (2019) Fire Detection Using Video Images and Temporal Variations, *ICAIIIC 2019 Conference Proceedings*.

Publications

[1] Jung S., Min J., Jung S., Suh S., Heo S. and **Kim S.*** (2021) Dental Image Data Generation for Instance Segmentation using Generative Adversarial Networks, *in revision*, *Corresponding author.

[2] Han J., Cho K., Lim W. and **Kim S.*** (2021) Reinforcement learning guided by double replay memory with applications to autonomous cars, *in revision*, *Corresponding author.

[3] Park Y., **Kim S.**, Chon K., Lee H., Kim J. and Shin J. (2021) Impacts of heavy rain and floodwater on floating debris entering an artificial lake (Daecheong Reservoir, Korea) During the summer, *Desalination and Water Treatment*.

[4] Suh S., Park Y., Ko K., Yang S., Ahn J., Shin J. and **Kim S.*** (2020) Weighted Mask R-CNN for Improving Adjacent Boundary Segmentation, *Journal of Sensors*, *Corresponding author.

[5] **Kim S.**, Heo S., Yang S., Han J. and Jung S. (2020) Instance Segmentation Guided by Weight Map with an Application to Tooth Boundary Detection, *Quantitative Bio-Science*.

[6] Jung S., Lee E., Lee Y., Ko J., Lee S., Cho J. and **Kim S.*** (2020) Estimation of Particulate Levels Using Deep Dehazing Network and Temporal Prior, *Journal of Sensors*,

*Corresponding author.

[7] Min J., Jung S., Jung S., Yang S. and **Kim S.*** (2020) Grammatical Error Correction Models for Korean Language via Pre-trained Denoising, *Quantitative Bio-Science*, *Corresponding author.

[8] **Kim S.**, Jung S., Yang S., Han J., Lee B., Lee J. and Han S. (2019) Vision-based Deep Q-learning Network Models to Predict Particulate Matter Concentration Levels using Temporal Digital Image Data, *Journal of Sensors* (SCIE).

[9] Lim J., Bang S., Kim J., Park C., Cho J. and **Kim S.*** (2019) Integrative deep learning for identifying differentially expressed (DE) biomarkers, *Computational and Mathematical Methods in Medicine* (SCIE), *Corresponding author.

[10] Han S., Park S., Zhong H., Ryu E., Wang P., Jung S., Lim J., Yoon J.* and **Kim S.*** (2019) Estimation of joint directed acyclic graphs with lasso family for gene networks, *Communications in Statistics - Simulation and Computation* (SCIE).

[11] Lim J., Cho J., Kim J., Kim J. and **Kim S.*** (2019) Deeper Integrative Neural Network Analysis for Multi-level Omics Data, *Quantitative Bio-Science*, *Corresponding author.

[12] Hwang U., Jeong J., Kim J., Cho S. and **Kim S.*** (2019) Computer Vision-based Method to Detect Fire Using Color Variation in Temporal Domain, *Quantitative Bio-Science*, * Corresponding author.

[13] **Kim S.*** and Kim S. (2018) Vision-based Predictive Model on Particulates via Deep Learning, *Journal of Electrical Engineering & Technology* (SCIE) * Corresponding author.

[14] Lee J., Jhong J., Cho Y., **Kim S.**, Koo J. (2018) Penalized log-density estimation using Legendre polynomials, *Communications in Statistics - Simulation and Computation* (SCIE).

[15] Lee J., **Kim S.***, Jhong J. and Koo J. (2018) Variable Selection and Joint Estimation of Mean and Covariance Models with an Application to eQTL Data, *Computational and Mathematical Methods in Medicine* (SCIE), *Corresponding author.

[16] **Kim S.**, Kang D. D., Park Y. and Tseng G. C. (2018) Meta-analytic Principal Component Analysis in Integrative Omics Application, *Bioinformatics* (SCI).

[17] Ma T., Huo Z., Kuo A., Zhu L., Fang Z., Zeng X., Lin C., Liu S., Wang L., Rahman T., Chang L., **Kim S.** et al. (2018) MetaOmics: Analysis Pipeline and Browser-based Software Suite for Transcriptomic Meta-Analysis, *Bioinformatics* (SCI).

[18] **Kim S.**, Jhong J., Lee J., Koo J-Y., Lee B. and Han S. (2017) Node-Structured Integrative Gaussian Graphical Models Guided by Pathway Information, *Computational and Mathematical Methods in Medicine* (SCIE).

[19] **Kim S.**, Oesterreich S., Kim S., Park Y. and Tseng G. C. (2017) Integrative Clustering of Multi-level Omics Data for Disease Subtype Discovery Using Sequential Double Regularization, **Received the Distinguished Student Paper Award for ENAR 2015 and ASA Biometrics Section David P Byar Travel Award at JSM 2015**, *Biostatistics*, 18(1):165-179 (SCIE).

- [20] **Kim S.**, Lin C. and Tseng G. C. (2016) MetaKTSP: A Meta-Analytic Top Scoring Pair Method for Robust Cross-Study Validation of Omics Prediction Analysis, *Bioinformatics*, 32:1966-1973 (SCI).
- [21] **Kim S.**, Lee J., Jhong J. and Koo J-Y. (2017) Meta-analytic Support Vector Machine for Integrating Multiple Omics Data, *BioData Mining*, 10:2 (SCIE).
- [22] Han S.*, **Kim S.***, Seok J., Yoon J. and Zhong H. (2017) Estimation of Directed Subnetworks in Ultra High Dimensional Data for Gene Network Problem, *Statistics and Its Interface* (SCIE), *Joint first authors.
- [23] **Kim S.** (2016) Weighted K-means Support Vector Machine for Cancer Prediction, *SpringerPlus* (section in statistics), 5:1162 (SCIE).
- [24] **Kim S.**, Herazo-Maya J. D., Kang D. D., Juan-Guardela B. M., Tedrow J., Martinez F.J., Sciurba F.C., Tseng G.C. and Kaminski N. (2015) Integrative Phenotyping Framework (iPF): Integrative Clustering of Multiple Omics Data Identifies Novel Lung Disease Subphenotypes, *BMC Genomics*, 16:924 (SCIE).
- [25] Jhong J., Lee J., **Kim S.**, Koo J-Y. (2017) Joint Modeling for Mean Vector and Covariance Estimation with l1-Penalty, *Quantitative Bio-Science*.
- [26] Jung H., **Kim S.**, Lee J., Kim J., Han S. (2017) Differential network analysis for Triple positive and Triple negative breast cancer genes, *Journal of Breast Cancer* (SCIE).
- [27] Zhu L., Ding Y., Chen C., Huo Z., **Kim S.**, Oesterreich S. and Tseng G. C. (2016) MetaDCN: meta-analytic framework for differential coexpression network detection with an application to breast cancer, *Bioinformatics* (SCI).
- [28] Hwang H., **Kim S.** and Kim H. (2016) Reversible Data Hiding Using Sparse Least Square Predictor via the Lasso, *EURASIP Journal on Image and Video Processing*, 2016:42 (SCIE).
- [29] Herazo-Maya J. D.*, Noth I.*, Duncan S. R.*, **Kim S.**, Ma S., Tseng G. C., Feingold E., Juan-Guardela B. M., Richards T. J., Lussier. Y., Huang Y., Vij R., Lindell K. O., Xue J., Gibson K. F., Shapiro S. D., Garcia J. G., and Kaminski N. (2013) Peripheral Blood Mononuclear Cell Gene Expression Profiles Predict Poor Outcome in Idiopathic Pulmonary Fibrosis, *Science Translational Medicine*, 5, 205-136 (SCIE).
- [30] Liu S., Tsai W., Ding Y., Chen R., Fang Z., Huo Z., **Kim S.**, Ma T., Chang T., Priedigkeit M., Lee A., Luo J., Wang H., Chung I., Tseng G., (2015) Comprehensive evaluation of fusion transcript detection algorithms and a meta-caller to combine top performing methods in paired-end RNA-seq data, *Nucleic Acids Research*, 17 (SCI).
- [31] Schwaderer A., Wang H., **Kim S.**, Kline J., Liang D., Brophy P., et al. (2016) Polymorphisms in α -Defensin-Encoding DEFA1A3 Associate with Urinary Tract Infection Risk in Children with Vesicoureteral Reflux, *Journal of the American Society of Nephrology* (SCIE).
- [32] Suryawanshi S., Budiu R. A., Elishaev E., Zhang L., **Kim S.**, Tseng C. G., Mantia-Smaldone G., Ma T., Donnellan N., Lee T., Mansuria S., Edwards R., Huang X. and Vlad A. M. (2014) Complement Pathway Is Frequently Altered in Endometriosis and

Endometriosis-Associated Ovarian Cancer, *Clinical Cancer Research*, 23, 6163-74 (SCIE).

[33] Suh Y., **Kim S.**, Kim S., Park J., Lim H., Park H., Choi H., Ng D., Lee M., and Nam M. (2013) Combined Genome-Wide Linkage and Association Analyses of Fasting Glucose Level in Healthy Twins and Families of Korea, *Journal of Korean Medical Science*, 28, 415-423 (SCI).

Book Chapter

[1] **Kim S.**, Huo Z., Park Y. and Tseng G. C. (2015) MetaOmics: Transcriptomic Meta-Analysis Methods for Biomarker Detection, Pathway Analysis and Other Exploratory Purposes pp. 39-67, Book chapter in Integrating omics data: statistical and computational methods. *Cambridge University Press*.

Honors, Awards & Fellowships

2015 Best Student Presentation Awards, Graduate School of Public Health, University of Pittsburgh, Mar, 2015.

2015 Student of the Year, ASA Pittsburgh Chapter, Apr 2015.

ASA Biometrics Section David P Byar Travel Award, American Statistical Association (ASA), Aug 2015.

Title: "Integrative multi-omics clustering for disease subtype discovery by sparse overlapping group lasso and tight clustering".

Distinguished Student Paper Award for ENAR 2015, ENAR, Mar 2015.

Title: "Integrative multi-omics clustering for disease subtype discovery by sparse overlapping group lasso and tight clustering".

References

Ja-Yong Koo, Ph.D.
Professor of Statistics
Department of Statistics
Korea University
Anam-dong, Seoul, 136-701
South Korea Email: jykoo@korea.ac.kr
Office: +82 (02) 3290 - 2240

George C. Tseng, Sc.D.
Professor of Biostatistics
Department of Biostatistics (primary appointment)
Department of Human Genetics
Department of Computational & Systems Biology

University of Pittsburgh
Pittsburgh, PA 15261
Email: ctseng@pitt.edu
Office: +1 (412) 624-5318

~~~ Last updated: January 15, 2021~~~