

Hannah Kim

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Research Interest

VIRAL EVOLUTION | ALGORITHM DEVELOPMENT | DISEASE DYNAMICS

Education

Temple University

PHD BIOINFORMATICS

• Advisor: Dr. Sergei L Kosakovsky Pond

Philadelphia, PA

08/2019 - 05/2024

Carnegie Mellon University

MS COMPUTATIONAL BIOLOGY

Pittsburgh, PA

08/2015 - 12/2016

Carnegie Mellon University

BS CHEMISTRY

Pittsburgh, PA

08/2010 - 05/2013

Relevant Experience

- 2021-2022 **Bioinformatics Engineer**, Lifetime Omics
- 2017-2019 **Bioinformatics Analyst / Software Engineer**, Children's Hospital of Philadelphia
- 2017 **Research Programmer**, Computational Biology Department, Carnegie Mellon University
- 2016 **Graduate Researcher**, Computational Biology Department, Carnegie Mellon University
- 2016 **Course Developer**, Computational Biology Department, Carnegie Mellon University
- 2013-2014 **Post-Baccalaureate Researcher**, Department of Biological Sciences, Carnegie Mellon University
- 2012-2013 **Undergraduate Student Researcher**, Department of Chemistry, Carnegie Mellon University
- 2011 **Student Intern, Summer Research Institute**, Department of Biological Sciences, Carnegie Mellon University

Publications

PUBLISHED

- Huzar, J., **Kim, H.**, Kumar, S., Miura, S. (2022). MOCA for integrated analysis of gene expression and genetic variation in single cells. *Frontiers in Genetics*. doi:10.3389/fgene.2022.831040
- Ding, Y., **Kim, H.**, Madden, K., Loftus, J., Chen, G., Allen, D., Zhang, R., Xu, J., Chen, C., Xu, Y., Tasian, S., Tan, K. (2021). Network Analysis Reveals Synergistic Genetic Dependencies for Rational Combination Therapy in Philadelphia Chromosome-like Acute Lymphoblastic Leukemia. *Clinical Cancer Research*. doi:10.1158/1078-0432.CCR-21-0553
- Tarca, A. L., Pataki, B. Á., Romero, R., Sirota, M., Guan, Y., Kutum, R., Gomez-Lopez, N., Done, B., Bhatti, G., Yu, T., Andreoletti, G., Chaiworapongsa, T., **The DREAM Preterm Birth Prediction Challenge Consortium**, Hassan, S. S., Hsu, C., Aghaepour, N., Stolovitzky, G., Csabai, I., Costello, J. C. (2021). Crowdsourcing assessment of maternal blood multi-omics for predicting gestational age and preterm birth. *Cell Reports Medicine*, 2(6). doi:10.1016/j.xcrm.2021.100323
- Ichikawa, Y., Bruno, V. M., Woolford, C. A., **Kim, H.**, Do, E., Brewer, G., Mitchell, A. P. (2021). Environmentally contingent control of *Candida albicans* cell wall integrity by transcriptional regulator Cup9. *Genetics*. doi: 10.1093/genetics/iyab075
- Tao, Y., Rajaraman, A., Cui, X., Cui, Z., Chen, H., Zhao, Y., Eaton, J., **Kim, H.**, Ma, J., Schwartz, R. (2021). Assessing the Contribution of Tumor Mutational Phenotypes to Cancer Progression Risk. *PLOS Computational Biology*, 17(3). doi:10.1371/journal.pcbi.1008777

- He, B., Gao, P., Ding, Y., Chen, C., Chen, G., Chen, C., **Kim, H.**, Tasian, S. K., Hunger, S. P., Tan, K. (2020). Diverse noncoding mutations contribute to deregulation of cis-regulatory landscape in pediatric cancers. *Science Advances*, 6(30). doi:10.1126/sciadv.aba3064
- Lin, C., Jain, S., **Kim, H.**, Bar-Joseph, Z. (2017). Using neural networks for reducing the dimensions of single-cell RNA-Seq data. *Nucleic Acids Research*, 45(17). doi:10.1093/nar/gkx681

PREPRINT

- Tao, Y., Rajaraman, A., Cui, X., Cui, Z., Eaton, J., **Kim, H.**, Ma, J., Schwartz, R. (2019). Improving personalized prediction of cancer prognoses with clonal evolution models. *bioRxiv*. doi:10.1101/761510

Presentations

* *presenting author*

CONTRIBUTED PRESENTATIONS

- Kim, H.***, Shank, S., Kosakovsky Pond, S. L. (2022). PProperty Informed Models of Evolution (PRIME). Poster: The 31st KSEA Northeast Regional Conference, Hybrid
- Hu, Y., Chen, C., Ding, Y.*, **Kim, H.**, Tan, K. (2019). Synergistic Control Genes in Cancer Gene Networks as Targets for Combination Therapy. Poster: Children's Hospital of Philadelphia Research Poster day and Scientific Symposium, Philadelphia, PA.

Awards, Fellowships, & Grants

2022	KSEA Excellent Poster Award , The 31st KSEA Northeast Regional Conference	\$ 100
2022	CST Three-Minute Thesis Competition 2nd Place Award , Temple University	\$ 250
2015	Departmental Merit Fellowship , Carnegie Mellon University	\$ 3000
2013	Mellon College of Science Research Honors , Carnegie Mellon University	
2012	Summer Undergraduate Research Fellowship , Carnegie Mellon University	\$ 3500

Teaching Experience

* *awarded Research Assistantship unless otherwise noted*

- F2020 **BIOL-3111/5111 Genomics in Medicine**, Teaching Assistant
- S2020 **BIOL-1012 General Biology II**, Teaching Assistant
- F2019 **BIOL-2112 Introduction to Cellular and Molecular Biology**, Teaching Assistant

Doctoral Coursework

- F2020 **BIOL-5128 Genomics and Infectious Disease Dynamics**
- F2020 **BIOL-8210 Seminar: "Ecoevo discuss"**
- F2020 **STAT-8109 Applied Statistics and Data Science**
- S2020 **BIOL-5241 Genomics and Evolutionary Biology of Parasites**
- S2020 **CIS-5517 Data-Intensive and Cloud Computing**
- S2020 **CIS-5523 Knowledge Discovery and Data Mining**
- F2019 **BIOL-5111 Genomics in Medicine**
- F2019 **BIOL-5466 Topics in Bioinformatics**
- F2019 **BIOL-8210 Seminar Biol 8210 at Center for Computational Genetics and Genomics**

Outreach & Professional Development

PROFESSIONAL MEMBERSHIPS

College of Science and Technology-Graduate Student Organization (CST-GSO), DEI Representative, *12/2021-*
College of Science and Technology Diversity, Equity, and Inclusion Committee, Student Representative, *11/2021-*
AnitaB.org, Member, *09/2021-*
Biology Graduate Student Society (BGSS), Vice President, *09/2021-*
Society for Molecular Biology and Evolution (SMBE), Member, *01/2021-*
Korean-American Scientists and Engineers Association (KSEA), Member, *02/2020-*
Philadelphia Korean Scholars Association (PKSA), Member, *06/2019-*

VOLUNTEERING

George Washington Carver Science Fair, Science Fair Judge, *03/2022*

CERTIFICATES

Matrix Algebra for Engineers, Coursera *2022*
Viruses and How to Beat Them: Cells, Immunity, Vaccines, IsraelX *2022*

Coding Languages

PYTHON , R , MATLAB , BASH , JAVASCRIPT , GOLANG