

Hannah Kim

Temple University, 1801 N Broad St, Philadelphia, PA 19122

✉ hannah.kim0007 [at] temple.edu | 🏠 <https://hannahkimincompbio.github.io> | 🌐 hannahkimincompbio

Research Interest

DISEASE DYNAMICS | VIRAL EVOLUTION | ALGORITHM DEVELOPMENT

Education

Temple University

PHD BIOINFORMATICS

• Advisor: Dr. Sergei L Kosakovsky Pond

Philadelphia, PA

2019 - present

Carnegie Mellon University

MS COMPUTATIONAL BIOLOGY

Pittsburgh, PA

2015 - 2016

Carnegie Mellon University

BS CHEMISTRY

Pittsburgh, PA

2010 - 2013

Relevant Experience

- 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

- 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., **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

ACCEPTED

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:10.1101/2021.01.06.425608

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

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

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

F2021	BIOL-3111/5111 Genomics in Medicine , Teaching Assistant
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