# Hannah Kim

Temple University, 1925 N 12th St, Room 643, Philadelphia, PA 19122

■ hannah.kim0007@temple.edu | ★ https://hannahkimincompbio.github.io | ■ hannahkimincompbio

Research Interest \_\_\_\_ COMPUTATIONAL BIOLOGY | ALGORITHM DEVELOPMENT | EVOLUTIONARY GENOMICS | VIRAL EVOLUTION Education \_\_\_\_ **Temple University** Philadelphia, PA PhD BIOINFORMATICS 08/2019 - 05/2024 Advisor: Dr. Sergei L Kosakovsky Pond **Carnegie Mellon University** Pittsburgh, PA MS COMPUTATIONAL BIOLOGY 08/2015 - 12/2016 **Carnegie Mellon University** Pittsburgh, PA 08/2010 - 05/2013 **BS CHEMISTRY** Relevant Experience \_ 2021-2022 Bioinformatics Engineer, Lifetime Omics Bioinformatics Analyst / Software Engineer, Children's Hospital of Philadelphia 2017-2019 Research Programmer, Computational Biology Department, Carnegie Mellon University 2017 **Graduate Researcher**, Computational Biology Department, Carnegie Mellon University 2016 2016 Course Developer, Computational Biology Department, Carnegie Mellon University Post-Baccalaureate Researcher, Department of Biological Sciences, Carnegie Mellon University 2013-2014 **Undergraduate Student Researcher**, Department of Chemistry, Carnegie Mellon University 2012-2013 Student Intern, Summer Research Institute, Department of Biological Sciences, Carnegie Mellon University

#### **PUBLISHED**

Publications \_\_

- 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*, 13:831040. 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 Chromosomelike Acute Lymphoblastic Leukemia. *Clinical Cancer Research*, 27(18). 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., Aghaeepour, N., Stolovitzky, G., Csabai, I., Costello, J. C. (2021). Crowdsourcing assessment of maternal blood multiomics 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*, 218 (3). 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 \_

#### **CONTRIBUTED PRESENTATIONS**

- Kim, H.\*, Shank, S., Kosakovsky Pond, S. L. (2022). PRoperty 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	Selected Attendee Support, Scientists and Engineers Early Career Development (SEED)	\$ 200+
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

#### Doctoral Coursework

F2020 BIOL-5128 Genomics and Infectious Disease	D:
F/11/11 BIGH-51/X GENOMICS AND INTECTIONS DISEASE	IIIVnamics

F2020 BIOL-8210 Seminar: "Ecoevo discuss"

## Teaching Experience \_\_\_\_\_

<sup>\*</sup> presenting author

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

<sup>\*</sup> 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

Outreach & Professional Development
LEADERSHIPS
Temple University Korean Graduate Student Association (TKGSA), Founding Member & Treasurer, 03/2022-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-Biology Graduate Student Society (BGSS), Vice President, 09/2021-
Professional Memberships
Korean-American Innovative Technology Engineers and Entrepreneurs (KITEE), 04/2022-Society for the Study of Evolution (SSE), 04/2022-AnitaB.org, 09/2021-Society for Molecular Biology and Evolution (SMBE), 01/2021-Korean-American Scientists and Engineers Association (KSEA), 02/2020-Philadelphia Korean Scholars Association (PKSA), 06/2019-
COMMUNITY INVOLVEMENT
Frontiers in Oncology (1x/yr), Ad Hoc Reviewer, 2022  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