Gyu (Gyuhyeon) Kim

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EDUCATION	Ph.D. Candidate in Biochemistry Stanford University, Stanford, CA, USA	Sep. 2021 — Present	
	Ph.D. Minor. Candidate in Computer Science Stanford University, Stanford, CA, USA	Sep. 2023 — Present	
	B.S. Chemistry, Minor in Biological Science Seoul National University, Seoul, South Korea	Mar. 2013 — Feb. 2019	
RECENT	Graduate Research	March 2022 — Present	
Research Experience	 Stanford University, Stanford, CA, USA Building and interpreting machine learning models of predicting transcription activity from transcription factor binding data over tissue differentiation 		
	• Characterizing nonconsensus binding of human transcription factors using ENCODE dataset analysis and machine learning		
	• Identified and validated long noncoding RNAs impacting skin cancer in a global scale via scRNA-seq and CRISPRi screen (Kim et al., 2025) Advisor: Dr. Paul Khavari		
	Committee: Dr. Anshul Kundaje, Dr. Mark Krasnow, Dr. Aaron Straight		
	Research Technician Harvard Medical School and Dana-Farber Cancer Institute, Bos • Assessing the association between cell nuclear envelope rupture and D Advisor: Dr. David Pellman		
Publications	Gyuhyeon Kim, Vivekanandan Ramalingam, Paul A. Khavari. Comprehensive characterization of nonconsensus binding of human transcription factor. Manuscript in preparation.		
	Douglas F. Porter,, Gyuhyeon Kim ,, Paul A. Khavari. Disease-Linked Regulatory DNA Variants and Homeostatic Transcription Factors in Epidermis. <i>Revision in Nature Comm.</i> 2025		
	Gyuhyeon Kim , Zurab Siprashvili,, Paul A. Khavari, Luca Ducol reveals lncRNA portfolio crucial for cutaneous squamous cell carcinoma <i>Investigative Dermatology</i> . 2025		
Leadership & Community Activity	Vice President for Stanford Biosciences Student Association, Stanford • Supporting the incoming leadership, organizing the student program and community meetings	n proposals	
	• Working as a student representative in the bioscience faculty meet President for Stanford Biosciences Student Association, Stanford Univ		
	 Managed the overall the Bioscience program community activities, communications between student officers 	· ·	
	Social Chair for Stanford Biosciences Student Association, Stanford U • Planned and organized variety of social events for bioscience programmer.	· ·	
Teaching & Outreach Experience	Mentor for Undergraduate Research Program, Stanford University Tutor for YES for CURE Program, Dana-Farber/Harvard Cancer Cente • Gave virtual lab sessions for an underrepresented minority summe		
	Tutor for Basic Chemistry, Seoul National University	2014,2015,2018	
	Openigan for Wookly Dublic Science Lecture Scoul National University	2019	

Organizer for Weekly Public Science Lecture, Seoul National University

Tutor for Math in SAMSUNG Dream Class, SAMSUNG Welfare Foundation

 $\bullet \ \ \textit{Mentored middle school students from socioeconomic minority backgrounds}$

2018

 $2014,\,2015$

Awards,
Honors,
SCHOLARSHIPS

Doctoral Study Abroad Grant, Korea Foundation for Advanced Studies	2021-2025	
Blavatnik Fellowship, Blavatinik Family Foundation	2023-2024	
MAC3 Graduate Fellowship, MAC3 Impact Philanthropies	2022-2024	
Presidential Award of Korean Chemical Society, Korean Chemical Society	2019	
Honored Graduate of Academic Excellence, Seoul National University	2019	
Kwanjeong Undergraduate Scholarship, Kwanjeong Educational Foundation	2015, 2018	
Dean's List, Seoul National University	2013, 2014, 2015	
Academic Excellence Award, Seoul National University	2015	
Best Tutor for Basic Chemistry, Seoul National University	2015	
1st Place, Science Essay Contest for the Nobel Prize, Korea Ministry of Education		
1st Place, HANCOM Office Software Development Contest, HANCOM Incorporate	e 2015	
National Undergraduate Scholarship, Korea Student Aid Foundation	2013, 2014	

Relevant Skills

Programming

Language: Python (Proficient), R, Bash (Intermediate), C++, Haskell, Agda (Beginner)

Python: ML training with different architectures and model interpretation (Linear model, Deep Neural Network, various tree models, Transformer, LSTM). ML relevant packages (PyTorch, Scikitlearn, SHAP, and other ML tools for tree models). Fine-tuning DNA-LLM. Data analysis with pandas, matplotlib, seaborn. Building pipeline with Snakemake. R: Data analysis and visualization with ggplot Bioinfo: Human genetic sequencing data processing (ChIP, CRISPR screen), Seurat Advanced Coursework Completed: Artificial Intelligence (CS221), Machine Learning (CS229), Continuous Mathematical Methods with an Emphasis on Machine Learning (CS205L), Graph Machine Learning (CS224W), Mining Massive Dataset (CS246), Natural Language Processing with Deep Learning (CS224N), Probabilistic Graph Model (CS228), Introduction to Regression Models and Analysis of Variance (STATS 203)

Experiment

CRISPR-based genetic screen, Immunofluorescence, CRISPR & Lentivirus-mediated cell line generation, Bacterial & mammalian (adherent and suspension) cell culture, RNAi, Transwell migration assay, Colony assay, Biochemistry work on DNA, RNA, and Proteins