## Gyu (Gyuhyeon) Kim

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	<ul> <li>Stanford University, Stanford, CA, USA</li> <li>Building and interpreting machine learning models for predicting transcription activity from transcription factor binding datasets over tissue differentiation</li> </ul>		
	• Characterizing nonconsensus binding of human transcription factors with the ENCODE datasets using machine learning and correlation analyses		
	• Identified and validated long noncoding RNAs impacting skin cancer in a seq and CRISPRi screen (Kim et al., 2025) Advisor: Dr. Paul Khavari Committee: Dr. Anshul Kundaje, Dr. Mark Krasnow, Dr. Aaron Straight	global scale via scRNA-	
Publications	<b>Gyuhyeon Kim</b> , Vivekanandan Ramalingam, Paul A. Khavari. Comprehensive characterization of nonconsensus binding of human transcription factor. <i>Manuscript in preparation</i> .		
	Douglas F. Porter,, <b>Gyuhyeon Kim</b> ,, Paul A. Khavari. Disease-Linked Regulatory DNA Variants and Homeostatic Transcription Factors in Epidermis. <i>Under revision</i> . 2025		
	<b>Gyuhyeon Kim</b> , Zurab Siprashvili,, Paul A. Khavari, Luca Ducoli. In vivo CRISPRi screen reveals lncRNA portfolio crucial for cutaneous squamous cell carcinoma tumor growth. <i>Journal of Investigative Dermatology</i> . 2025		
Leadership & Community Activity	<ul> <li>Vice President for Stanford Biosciences Student Association, Stanford Un</li> <li>Supporting the incoming leadership, organizing the student program pre and community meetings</li> </ul>	v	
	• Working as a student representative in the bioscience faculty meeting  President for Stanford Biosciences Student Association, Stanford Universi	ty 2023 - 2024	
	<ul> <li>Managed overall community activities, the Bioscience Resource Fair, and communications between student officers</li> </ul>	v	
	Social Chair for Stanford Biosciences Student Association, Stanford University of Planned and organized a variety of social and wellness events	ersity 2022 - 2023	
TEACHING & OUTREACH EXPERIENCE	Mentor for Undergraduate Research Program, Stanford University Tutor for YES for CURE Program, Dana-Farber/Harvard Cancer Center	2023 - 2025 2020	
	• Gave virtual lab sessions for an underrepresented minority summer st		
	Tutor for Basic Chemistry, Seoul National University	2014, 2015, 2018	

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

2018

2014, 2015

Organizer for Weekly Public Science Lecture, Seoul National University

Tutor for Math in SAMSUNG Dream Class, SAMSUNG Welfare Foundation

AWARDS,
Honors,
${\bf SCHOLARSHIPS}$

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	2015
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