

Soo Min Lee

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EDUCATION

Korea Advanced Institute of Science and Technology (KAIST)
M.S., Major in Bio and Brain Engineering

Daejeon, Korea
August 2020 – August 2022

Handong Global University (HGU)
B.S., Double major in *Computer Science* and *Life Science*
Summa cum laude, Early graduation

Pohang, Korea
Feb 2017 – August 2020

Allendale Columbia School
Studied abroad for high school in the United States

Rochester, NY, U.S.A.
Jun 2016

RESEARCH EXPERIENCE

Korea Advanced Institute of Science and Technology (KAIST)
Researcher, SBiE laboratory (prof. Kwanghyun Cho)

Daejeon, Korea
July 2020 – Feb 2023

1. ‘Deep learning untangles the resistance mechanism of p53 reactivator in lung cancer cells’
(*Master Thesis & Journal publication & KOR/US patent*)

- **Problem:** While the most common TP53 mutations in lung cancer make p53 activator treatments essential, drug resistance significantly limits their effectiveness.
- **Contributions:** Developed a novel computational approach using Generative adversarial networks and Graph neural networks to identify drug resistance mechanisms and therapeutic targets. Conducted *in vitro* validation, confirming the therapeutic potential of the identified gene targets.
- **Results:** Identified a key pathway and biomarker associated with p53 resistance, leading to a Korean/US patent application and publication in *Cell iScience*.

2. ‘A systems biological study to investigate the fundamental principle and overcoming method of adaptive resistance to targeted anti-cancer therapy’

- **Problem:** Overcoming the limitations of colon cancer treatment requires understanding adaptive resistance to targeted therapies, like MEK inhibitors, and developing strategies to combat it.
- **Contributions:** Performed drug concentration and timepoint screenings, established MEK inhibitor-resistant colon cancer cell lines, analyzed gene expression changes in key pathways, and produced RNA sequencing data.
- **Results:** Discovered significant resistance-related genes, supporting model construction and contributing to securing \$300K in funding from the *National Research Foundation of Korea*.

Handong Global University (HGU)
Researcher, Biodata laboratory (prof. Taejin Ahn)

Pohang, Korea
Jan 2019 – Jun 2020

1. ‘Identification of Lifestyle Factors correlated with Gastroesophageal Reflux Disease’ (*Bachelor Thesis*)

- **Problem:** The increasing global incidence of Gastroesophageal Reflux Disease (GERD) is associated with unhealthy lifestyles, making it crucial to identify contributing lifestyle factors.
- **Contributions:** Collaborated with a physician from Seoul National University Hospital (SNUH) to analyze GERD risk factors using logistic regression, hierarchical clustering, and subgroup analysis on a large patient dataset.
- **Results:** Identified lifestyle factors contributing to GERD risk, which were presented in my bachelor’s thesis.

2. ‘Diabetic Status Monitoring and Prediction using Clinical and Genomic Data’

- **Problem:** Diabetes is a chronic disease that requires continuous monitoring, which can be difficult for patients to manage on their own, highlighting the need for robust predictive tools for prevention and management.
- **Contributions:** Developed a blood glucose level forecasting model using deep neural networks. Analyzed clinical data from SNUH and genomics data from the GWAS catalog.
- **Results:** Built an effective blood glucose prediction model with accurate results, leading to a scholarship award through the Capstone project submission.

Seoul National University (SNU)
Research intern, Bioinformatics & Biostatistics laboratory (prof. Taesung Park)

Seoul, Korea
Jan 2020 – Feb 2020

1. ‘Bio-synergy: Metabolites Data Analysis’

- **Problem:** Metabolites are crucial for predicting type 2 diabetic status, as diabetes is a metabolic disease; however, accurate predictions depend on effective data preprocessing and appropriate model selection.

- **Contributions:** Preprocessed LC/MS-based metabolite data from the Korea Basic Science Institute using Combat and quantile normalization to reduce non-biological variation. Improved a diabetes prediction model by applying elastic net regression.
- **Results:** Enhanced prediction accuracy and gained experience in handling metabolite data.

INDUSTRY EXPERIENCE

Inocras (Headquarter in San Diego, CA)	Daejeon, Korea
<i>Computational Genomics Center – Pipeline Development Team</i>	<i>March 2023 – Present</i>
<ul style="list-style-type: none"> ▪ Developed algorithms for actionable findings, including targetable mutations, on-label drug recommendations, and clinical trial matching based on patients' genetic variations. This involved integrating, refining, and optimizing data from The Clinical Knowledgebase (CKB) into our somatic pipeline. Collaborated with oncologists to develop a Disease Checklist for precise on-label drug recommendations. ▪ Improved Structural Variation (SV) detection in somatic and germline pipelines by optimizing SV calling methods and post-processing, resulting in more accurate variant identification (e.g., detecting missing fusion variants, filtering false positives, and reducing short inversions, which decreased SV calls by up to 78%) and speeding up the process. ▪ Contributed to an AI project by developing an AI-driven deep mutation caller for small variants, enhancing the accuracy and efficiency of mutation detection. 	

PUBLICATIONS & PATENTS

Publications	
<ul style="list-style-type: none"> ▪ <u>Lee SM</u>, Han Y, Cho KH. Deep learning untangles the resistance mechanism of p53 reactivator in lung cancer cells. <i>iScience</i>. 2023 Nov 1;26(12):108377. doi: 10.1016/j.isci.2023.108377. PMID: 38034356; PMCID: PMC10682260. 	
Patents	
<ul style="list-style-type: none"> ▪ USES OF THBS1 INHIBITOR FOR OVERCOMING DRUG RESISTANCE OF CANCER <ul style="list-style-type: none"> • Application: 10-2023-0038078, 2023 March 23 (Republic of Korea), PCT/KR2023/003905, 2023 March 23 (PCT), 18/840,605, 2024 August 22 (USA) 	

AWARDS & OTHER EXPERIENCES

Awards	
▪ Top honor in Life Science from Handong Global University (1%)	2020
▪ Academic Excellence Scholarship	2018, 2019, 2020
▪ Software Mileage Scholarship	2020
▪ Excellence prize at Software Festival held by Handong University; Convergence Research Contest & SW Problem-Solving Idea Contest.	2019
Other Experiences	
▪ Compassion Mate Translation Volunteer	Sep 2016 – Dec 2020
▪ First-year Student Mentoring	Feb 2019 – Dec 2019
▪ Machine Learning with Real World Data Camp, C++ Camp	Jul 2019
▪ Student Government – Scribe, Director, Treasurer	Feb 2017 – Jun 2019
▪ Translation Society – Co-Chair	Feb 2018 – Dec 2018
▪ National Undergraduate Symposium on Biology – Topic: Autoimmune Disease	Jun 2018 – Sep 2018
▪ World Friends ICT Volunteers Program in Kenya (National Information Society Agency)	Jul 2017 – Aug 2017

SKILLS & OTHERS

Languages: Native fluency in Korean, fluent in English
Programming Languages: Python, R, Java, C, C++
Laboratory Skills: Cell culture, DNA/RNA extraction and purification, recombinant DNA technology, PCR, qRT-PCR, Western blotting, gel electrophoresis, transfection, viral transduction
Coursework: Applied Genomics, Biodata Analysis, Molecular Biology, Machine Learning, Graph Machine Learning and Mining, Data Mining, Genome Bioinformatics
Test Scores: TOEIC (990/990), TOEFL (111/120, My Best Score)
Qualifications: Big Data Analyst (Issued by Ministry of Science and ICT / Statistics Korea)
Interests: Machine Learning, Multiomics, Bioinformatics, Personalized Medicine, Drug Discovery