Haeun (Hannah) Hwangbo

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◆ haeunhwangbo.github.io

HIGHLIGHTS

- Ph.D. candidate in computational biology and bioinformatics at UNC-Chapel Hill (Expected to graduate in May 2024)
- Interested in clinical and (reverse-)translational cancer research with 7-year research experience
- Experienced in analyzing clinical trials and multi-omics data in preclinical and clinical samples

WORK EXPERIENCE

Bristol Myers Squibb, Informatics and Predictive Sciences

Computational Systems Biology Intern

San Diego, CA Jun 2023 – Aug 2023

- Predicted cell viability from perturbation transcriptomics by building a machine learning model; This approach is now integrated into compound selection strategy
- Utilized public resources to construct the model and validated on internal datasets
- Drove improvements in data processing and deposition from external collaborations

RESEARCH EXPERIENCE

The University of North Carolina at Chapel Hill, Adam Palmer Lab

Graduate Research Assistant

Chapel Hill, NC Aug 2019 - Present

- Developed computational method to identify individual drug activity in clinical trials of cancer combination therapy; Identified combinations that will benefit most from patient selection; Analyzed 92 FDA-approved drug combinations in the past 25 years
- Modeled drug-drug interactions in clinical trials of cancer combination therapy
- Performed RNA-seq trajectory analysis to discover biomarkers for aging in intestinal stem cells (collaboration with Oxford university)

Korea Advanced Institute of Science and Technology, Omics Laboratory

Research Scientist

Daejeon, Korea Mar 2018 — Jun 2019

- Conceived and co-led project on discovering biomarkers for cancers with homologous recombination deficiency; Analyzed transcript isoforms and used machine learning to predict clinical outcomes
- Developed vectorized algorithm for predicting vulnerabilities in breast cancer; Significantly improved the simulation speed; Used Connectivity Map to validate simulation results
- Constructed RNA-seq pipeline for government-funded project ("Integrated Multi-genomics-based Precision Medicine in Colon Cancer."); Performed pilot analysis on single-cell ATAC-seq

Seoul National University, Laboratory of Evolutionary Bioinformatics

Undergraduate Research Assistant

Seoul, Korea Jun 2016 – Jun 2017

- Designed and implemented autoencoder for dimensionality reduction of prokaryotic genome signatures; Clustered taxa based on low-dimensional projections
- Constructed pipeline predicting taxa of query sequences using tetranucleotide signatures

TECHNICAL SKILLS

- **Programming:** Python (pandas, numpy, scikit-learn, matplotib, seaborn, lifelines), R (bioconductor, ggplot2), bash, Linux CLI, C, MATLAB, Mathematica, SQL
- Data Science/Cloud Computing: Git, Docker, GNU make, snakemake, nextflow, HPC systems, AWS, web scraping
- **Bioinformatics/Omics analysis**: variant calling, peak calling, copy number variation, differential expression, differential splicing, trajectory analysis, methylation, CRISPR-KO, dose-response analysis
- Data processing: high-throughput sequencing, WGS, WES, RNA-seq, ATAC-seq, scATAC-seq, proteomics
- Public Databases: TCGA, GTEx, CCLE, Connectivity Map, GDSC, CTRPv2, COSMIC, DepMap
- Machine Learning/Statistics: regression analysis, generalized linear model, elastic net, xgboost, random forest, clustering, autoencoder, dimensionality reduction, classification, survival analysis, design & analysis of clinical trials

EDUCATION

The University of North Carolina at Chapel Hill

Ph.D. in Bioinformatics and Computational Biology

Big Data to Knowledge Certificate

Seoul National University

LEADERSHIP EXPERIENCE

B.S. in Biological Sciences (summa cum laude); Minor in Computer Science and Engineering

Bioinformatics and Computational Biology Steering Committee, Student Representative

Advocated student perspectives and contributed to curriculum development

Chapel Hill, NC Expected May 2024

> Seoul, Korea Feb 2018

Chapel Hill, NC

Aug 2022 - Present

Participated in search committees

UNC Computational Biosciences Club, Leadership, International Student Liaison

Assisted international students to navigate through career development

Coordinated with departments to facilitate internships for international students

UNC How to Learn to Code, Course Instructor

Created curricula and taught introductory Python to biomedical researchers

Chapel Hill, NC

Chapel Hill, NC Nov 2021 - Present

July 2022

PUBLICATIONS

- 1. Hwangbo, H., Patterson, S., Dai, A., Plana, D., & Palmer, A. C. (2023). Additivity predicts the efficacy of most approved combination therapies for advanced cancer. Nature Cancer, 1-12.
- (Preprint) Chen, J., Merrick, K.A., ..., Hwangbo, H., ..., & Yaffe, M. B. (2023) An RNA Damage Response Network Mediates the Lethality of 5-FU in Clinically Relevant Tumor Types. bioRxiv.
- 3. Palmer, A. C., Izar, B., Hwangbo, H., & Sorger, P. K. (2022). Predictable Clinical Benefits without Evidence of Synergy in Trials of Combination Therapies with Immune-Checkpoint Inhibitors. Clinical Cancer Research, 28(2), 368-377.
- 4. Kang, H. G., **Hwangbo, H.**, Kim, M. J., Kim, S., Lee, E. J., Park, M. J., Kim, J., Kim, B., Cho, E., Chang, S., Lee, J., & Choi, J. K. (2021). Aberrant transcript usage is associated with homologous recombination deficiency and predicts therapeutic responses. Cancer Research, 82(1), 142-154.
- 5. Jang, K., Park, M. J., Park, J. S<u>., **Hwangbo, H.,**</u> Sung, M. K., Kim, S., Jung, J., Lee, J.W., Ahn, S.H., Chang, S. & Choi, J.K. (2020). Computational inference of cancer-specific vulnerabilities in clinical samples. Genome biology, 21(1), 1-24.

HONORS & AWARDS

- Best Talk Award (2023, UNC Genetics Department)
- Korean Government Scholarship for Study Overseas (2019-2021, ~50 scholars selected nationwide)
- Study Abroad Applicant Scholarship (2018, Korea Foundation for Advanced Studies)
- Outstanding Peer Tutor 2nd Prize (2017, Seoul National University)
- DUO-Korea Fellowship (2015, ASEM-DUO Fellowship Programme, supported study abroad in United Kingdom)
- Dean's List (2014, 2015, Seoul National University)
- National Science & Technology Scholarship (2013-2016, 4-year full scholarship)