KwanHo An

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Education

Masters of Science(M.S.) | McWilliams School of Biomedical Informatics at UTHealth Houston

- · Major: Biomedical Informatics Research Track
- Fall 2023 Spring 2025 | GPA: 4.0/4.0
- Courses: Bioinformatics, Statistics, Health Data Science, Big data analysis, Deep Learning, RDBMS(SQL), NoSQL, Visualization(Tableau)

Bachelor of Science(B.S.) | McNeese State University

- · Major: Medical Laboratory Science
- · Fall 2017 Spring 2020 | Graduated *magna cum laude* (3.86/4.00 GPA)

Technical Skills & Tools

Programming & Data Analysis & visualization

· Python, R, SOL (PostgreSOL, Snowflake), MongoDB, Tableau, Jupyter Notebook, JSON

AI & Bioinformatics

· Linux, Docker, AlphaFold3, ESM-IF, PyTorch, ChimeraX, Protein Data Bank (PBD), ProteinMPNN, FoldX, USCS Genome Browser, GENCODE, bedtools, ClinVar

Lab & Workflow Management

· LIMS, LabVantage, GitHub/GitLab

Experience

Graduate Research Assistant (AI Antibody Engineering) | UTHealth Houston | Jan 2024 - May 2025

- · Performed data cleaning and applied deep learning models (inverse folding, AlphaFold3, structural and functional metrics) to design antibodies targeting influenza H1 and H3 subtypes.
- Engineered monoclonal antibodies for cancer immunotherapy using AI-driven mutation strategies to optimize structural and functional properties through CDRH3 modifications.

Medical Laboratory Scientist II | Natera (Sequencing team) | Aug 2022 - July 2023

- · Performed NGS, MLPA, FraX and Confirmation by using Nova Seq 6000(Illumina) and Capillary Electrophoresis 3730 (Thermo Fisher) instruments.
- · Reviewed sequencing data and ensured QC compliance via LIMS.

Lead, Medical Laboratory Scientist | Natera (Extraction team) | Dec 2020 - Aug 2022

 Oversaw daily lab operations, including troubleshooting and sample case management to identify and resolve stuck cases, trained new hires, and collaborated with supervisors to ensure timely sample processing.

Medical Laboratory Scientist I | Natera (Extraction team) | July 2020 - Dec 2020

 Performed plasma isolation, both automated and manual DNA extraction, and DNA dilutions (TECAN, QIAGEN)

Projects

Bioinformatics Python projects (GitHub: https://github.com/kh7939)

- Developed Python scripts for DNA sequence analysis, including GC content, motif detection, transcription and translation.
- · Built a deep learning model for Acute Lymphoblastic Leukemia (ALL) image classification.

Antibody design using AI (Graduate Research Assistant at UT Health Houston)

 Applied AI-driven methods to design antibodies targeting specific antigens, focusing on CDRH3 optimization to boost binding affinity.

HBB Gene (Sickle Cell Disease) Analysis

• Utilized UCSC Genome Browser, ClinVar, and various genomic data formats (BAM, SAM, GFF3, VCF) to analyze HBB gene variants and assess their clinical implications.

Licensure and Certifications

Board of certificate - MLS (Medical Laboratory Scientist) (ASCP)CM

· ASCP(American Society for Clinical Pathology) certified. KwanHo An, MLS(ASCP)CM