AIL NAHTANHOL

Data scientist interested in development and implementation of new deep learning algorithms

PROFESSIONAL EXPERIENCE

2023

Data Analyst Associate

Baylor College of Medicine

- Houston, TX
- · Developed automated pipeline for processing of NGS data with Docker, Python, and R
- Applied scRNA/snRNA-seq pipelines for mechanistic analysis of transcriptomic data
- Collaborated with software engineers and clinical geneticists to leverage LLMs for precision diagnosis of rare Mendelian genetic disorders
- · Supervised bioinformatics team to work with collaborators in using algorithms and data to drive hypothesis-driven research conclusions
- · Worked with Dr. Jeong and graduate students to develop mentor-mentee matching tool using LLMs with RAG and RLHF

2022 2020

Graduate Research Assistant

UTHSC Houston Graduate School of Biomedical Sciences

- O Houston, TX
- Developed a convolutional neural network with self-attention architecture using TensorFlow/Keras libraries to identifying high confidence transcription factor targets for HTLV-1.
- · Performed pseudotime analysis on scRNAseq data to infer which transcription factor programs drive COVID-19 infection severity with Seurat and DriVAER

2020 2019

Research Assistant I

MD Anderson Cancer Center

OHouston, TX

Processed, prepared, and stained patient samples for Nanostring and Cytometry

EDUCATION

2022 2020

M.S. in Quantitative Sciences

UTHSC Houston Graduate School of Biomedical Sciences

• Houston, TX

- Thesis: DeepHTLV: a deep learning model for detecting and elucidating human T-cell leukemia virus type 1 integration sites
- UTHSC Houston McGovern School of Medicine

OHouston, TX

2018 2015 2015

2011

B.S. in Biology, B.A. in Chemistry

Emory University

Atlanta, GA

Thesis: Investigating heterogeneity in the dynamics of virus and immune response

AWARDS

2022

Best Quantitative Sciences Program Seminar Presentation

UTHSC Houston Graduate School of Biomedical Sciences

O Houston, TX

2022

First Prize Poster Presentation

UTHSC Houston Graduate School of Biomedical Sciences

O Houston, TX

2016

Rheumatology Research Foundation Research Award

UTHSC Houston McGovern School of Medicine

O Houston, TX

CONTACT INFO

C) jjia1

Me

LANGUAGES

Python

Shell/Bash

Git

Node.js

PACKAGES

Tensorflow

PyTorch

Cuda

LangChain

PEFT

LoRA

Transformers

PUBLICATIONS

Deep learning for detecting and elucidating human T-cell leukemia virus type 1 integration sites (Cell Patterns 2023)

H.Xu* and J.Jia*, H.Jeong,Z.Z

Delineating COVID-19 immunological features using single-cell RNA sequencing (The Innovation 2022)

W.Liu, J.Jia, Y.Dai, W.Chen, G.Pei.Q.Yan. Z.Zhao

Investigating Cellular Trajectories in the Severity of COVID-19 and Their **Transcriptional Programs Using Machine Learning** Approaches (Genes 2021)

H.Jeong* and J.Jia*, Y.Dai, L.Simon, Z.Zhao

What Controls the Acute Viral Infection Following Yellow Fever Vaccination? (Bulletin of Mathematical Biology 2016)

J.Moore, H.Ahmed, **J.Jia**, R.Akondy, R.Ahmed, R.Antia

CERTIFICATES

Enhancing Data Science Outcomes With Efficient Workflow

NVIDIA

2024

Last updated on 2024-06-11.