

ZIYANG “Claude” HU

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Charlottesville, VA – 22901, USA

SKILLS

Programming Languages: Python, R, C, Rust, Java, Bash/Shell, SQL

Machine Learning: PyTorch Lightning, Transformers, PyTorch, TensorFlow, scikit-learn

Natural Language Processing: SentenceTransformers, LangChain, FastEmbed, CoreNLP, Stanza

Data Analysis and Management: NumPy, Pandas, SQLite, Hugging Face Datasets, Qdrant

Visualization and Design: ggplot2, matplotlib, Inkscape, LaTeX

EDUCATION

DUKE UNIVERSITY, School of Medicine, Durham, NC

Master of Biostatistics, May 2023

Relevant coursework: Software Tools for Data Science, Statistical Programming for Big Data, Probabilistic Machine Learning

EMORY UNIVERSITY, College of Arts and Sciences, Atlanta, GA

Bachelor of Science, May 2021

Double Major: Computer Science, Neuroscience and Behavioral Biology

Relevant coursework: Analysis of Algorithms, Machine Learning, Numerical Analysis, Big/Small Data and Visualization

EXPERIENCE

UNIVERSITY OF VIRGINIA, Charlottesville, VA

2023-Present

Sheffield Lab, Department of Genome Sciences

Scientific Programmer

- Developing Python/Rust packages for genomic interval data analysis and machine learning applications.
- Investigating text-file retrieval for ChIP-seq genomic interval output in bed narrowPeak format.
- Optimized genomic interval data storage and reduced database reindex runtime by 40%.
- Built a file retrieval pipeline combining semantic search with genomic interval representation learning.
- Curated ad-hoc datasets with biomedical ontologies.
- Fine-tuned open-source language models for information retrieval.

DUKE UNIVERSITY, Durham, NC

2022-2023

Department of Biostatistics & Bioinformatics

Graduate Research Assistant

- Developed Python/R functions to automate preprocessing of unstructured clinical data.
- Participated in systematic review by extracting and summarizing key information from relevant scientific literature.
- Performed medical concept extraction and entity mapping from electronic health records with NLP tools.
- Retrieved and preprocessed raw data from the Duke Clinical Research Data Mart (CRDM).
- Trained predictive models and analyzed model fairness across demographic groups.

EMORY UNIVERSITY, Atlanta, GA

2020-2021

Department of Sociology

Research Assistant

- Contributed to the development of a computational social science data analysis toolkit.
- Designed a pipeline using the annotation of Stanford CoreNLP for efficient relationship extraction.
- Preprocessed raw data for an interdisciplinary project to analyze statements from HIV patients.
- Utilized open-source Python libraries to perform sentiment analysis on interviews.

PUBLICATIONS

Franzosi, R., Dong, W., **Hu, Z.**, Dai, W., Cha, M., Piloto, R., & Wang, G. (2024). Automatic information extraction of the narrative elements who, what, when, and where. [Manuscript submitted for publication]. *Social Science Computer Review*.

Yang, R., Tong, J., Wang, H., Huang, H., **Hu, Z.**, Li, P., ... & Hong, C. (2025). Enabling inclusive systematic reviews: incorporating preprint articles with large language model-driven evaluations. *Journal of the American Medical Informatics Association*, ocaf137.

LeRoy, N. J., Campbell Jr, D. R., Stadick, S., Khoroshevskyi, O., Park, S. H., **Hu, Z.**, & Sheffield, N. C. (2025). Fast, memory-efficient genomic interval tokenizers for modern machine learning. *arXiv preprint arXiv:2511.01555*.

PRESENTATIONS

Huang, H., Tong, J., **Hu, Z.**, Li, Y., Pencina, M., Chen, Y., & Hong, C. . Enabling Inclusive Systematic Reviews: Incorporating Preprint Articles based on Semantic Learning and Large Language Model. Poster presentation at the ENAR Spring Meeting, Baltimore, MD, USA. March 10-13, 2024.

Xue, B., Khoroshevskyi, O., Stolarczyk, M., Mosquera, J. V., Campbell, D., **Hu, Z.**, Tambe, S., LeRoy, N., Gharavi, E., Duzlevski, O., & Sheffield, N. C. . BEDbase: A web application and API for genomic region sets. Poster presentation at the Biological Data Science Conference, Cold Spring Harbor, NY, USA, November 13-16, 2024.