Samira Jahangiri

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AI/ML Engineer — Data Scientist — Bioinformatics Engineer

About Me

PhD candidate with 8+ years of interdisciplinary experience in machine learning (transformers, GNNs), large-scale genomic analysis, and computational drug discovery. Proven ability to bridge biotech and AI through projects in protein engineering, NLP for biomedical text, and federated learning. Strong statistical foundation with applied expertise in Python/R and cloud computing, with proven ability to adapt domain-specific knowledge to new problem spaces. For detailed project portfolios: samirajahangiri.github.io

Technical Skills

- **Programming**: Python (PyTorch, TensorFlow, scikit-learn), R (tidyverse, Bioconductor), SQL, Bash, Julia Lang, LaTeX
- **AI/ML**: Transformer models (text embedding), GNNs (3D pose estimation), RNN/CNNs (genomic prediction), federated learning
- **Data Engineering**: AWS (S3, EC2), NGS pipelines (RNA-seq/WGS), Docker, Shiny/RMarkdown
- **Geometric ML**: Matrix algebra (SVD, eigendecomposition), manifold learning, graph embeddings

Professional Experience

Bioinformatics Specialist *Howard Hughes Medical Institute*   *Mar 2022 – Jul 2023*

- Led NGS data analysis (RNA-seq, whole-genome sequencing) for neurological disease research, identifying dysregulated pathways
- Developed ML pipelines with tree-based models for variant prioritization, contributing to 2 publications

Genomics Data Engineer *Shriners Hospital for Children*   *Jul 2021 – Feb 2022*

- Migrated petabyte-scale WGS data between AWS clouds (S3 $\rightarrow$ Glacier), reducing storage costs by 40%
- Automated ETL workflows for pediatric precision medicine research (cerebral palsy, scoliosis)

Data Scientist *USF Omics Hub*   *Aug 2018 – Dec 2021*

- Built R Shiny dashboards for microbiome/RNA-seq analysis (16S, metagenomics) used by Florida DOH
- Created statistical power calculator for microbiome studies (PERMANOVA effect size estimation)

Bioinformatics Consultant *Pars Silico Bioinformatics (Tehran)*   *August 2015 – June 2018*

- Optimized in-house molecular docking algorithm (Cytoscape/DrugBank integration) for pharmaceutical R&D

Research Experience

PhD Candidate in Big Data Analytics *University of Central Florida*   *2023 – Present*

- Developing complex-valued NLP models for biomedical text embedding, capturing semantic relationships via phase angles
- Applied GNNs/LSTMs to 3D pose estimation in autism research (privacy-preserving federated learning)

Graduate Research Assistant *University of South Florida*   *2018 – 2021*

- Developed AI models (RNNs/CNNs) to identify epigenetic drug-resistance markers in malaria

Graduate Research Assistant *University of Tehran*   *2015 – 2018*

- Published first-author study on computational protein-drug interactions (GF-17 antimicrobial peptide).

Education

- **PhD in Big Data Analytics**-Statistics track *University of Central Florida*   *2022 – Present*
- **MS in Public Health**-Biostatistics *University of South Florida*   *2019 – 2021*
- **MS in Biotechnology** *Azad University*   *2014 – 2016*

Publications

- Full list: [Google Scholar](https://scholar.google.com)