

MARJAN HOSSEINI

📍 University of Connecticut, Computer Science & Engineering Dept., 369 Fairfield Way, Storrs, CT 06268
✉️ marjan.hosseini@uconn.edu
[Google Scholar](#) [Marjan-Hosseini](#) [marjhosseini](#) [Website](#)

PROFESSIONAL SUMMARY

Ph.D. candidate in Computer Science and Engineering with 10+ years of experience in computational biology, machine learning, and large-scale data analysis. Hands-on background in modeling high-throughput sequencing data (e.g., nanopore), feature selection, and probabilistic/Bayesian ML. Experienced in end-to-end project execution: data preprocessing, model development, evaluation, and clear communication of results to collaborators with diverse technical backgrounds.

CORE SKILLS

Programming & Software Engineering: Python, R, Matlab, Java, Spark, Git/GitHub, Linux (shell scripting), C/C++ (working knowledge).

Machine Learning & Statistics: Deep learning, supervised/unsupervised learning, feature selection, statistical modeling, Bayesian inference, probabilistic graphical models, model evaluation and benchmarking.

Computational Biology: High-throughput sequencing analysis (ONT, PacBio, Illumina), variant calling, haplotype assembly, polyploid genomics, biological signal processing, omics data integration.

ML/AI Frameworks: PyTorch, TensorFlow, scikit-learn, PyMC3, CUDA-accelerated workflows.

Algorithms & Optimization: Graph algorithms, combinatorial, dynamic programming, large-scale data processing.

Collaboration & Leadership: Interdisciplinary teamwork, mentoring students, scientific writing, clear communication with both technical and non-technical collaborators, workshop lecturing and training.

EDUCATION

Ph.D. in COMPUTER SCIENCE AND ENGINEERING — **University of Connecticut**, USA Exp. 2026

Advisor: [Prof. Derek Aguiar](#)

Focus: Computational biology, Bayesian machine learning, deep learning for sequencing data.

Dissertation proposal defended; thesis defense expected in the coming weeks.

M.Sc. in COMPUTER SCIENCE AND ENGINEERING — **Politecnico di Milano**, Italy Jul 2018

Thesis: [“Feature Selection for Microarray Classification Problems”](#)

GPA: **110/110 (Honors – summa cum laude)**

B.Sc. in COMPUTER ENGINEERING (SOFTWARE) — **Shomal University**, Iran Feb 2006

Graduated in Computer Engineering (Software).

SELECTED INDUSTRY & RESEARCH EXPERIENCE

[Department of Computer Science and Engineering](#), University of Connecticut — Storrs, USA Aug 2019 – Present

Graduate Research & Teaching Assistant

- Design and implement machine learning and Bayesian models for high-throughput sequencing and biological time-series data (e.g., nanopore translocation times).

- Build data processing pipelines for large biological datasets, from raw data cleaning to model training and evaluation.

- Collaborate with interdisciplinary teams (bioinformatics, biology, and engineering) to define requirements and interpret results.

- Work on a joint project with deCODE Genetics on non-B DNA structures, including automating preprocessing and large-scale analysis pipelines for very large sequencing datasets.

- Communicate findings through publications, conference presentations, and reports.
- Mentor undergraduate and junior graduate students on research workflows, experimental design, and coding best practices.

DEIB, Politecnico di Milano — Milan, Italy
Research Fellow

- Developed GPU performance prediction models and feature selection methods for large-scale cloud applications (EU H2020 ATMOSPHERE project).
- Implemented machine learning pipelines to evaluate and optimize application performance under resource constraints.
- Built tools for mapping between ontologies and data schemas (SPRINT project), enabling semantic interoperability across complex data sources.

K4Tech, R&D Department — Milan, Italy
R&D Expert in Machine Learning and Signal Processing

- Designed and implemented ML and digital signal/image processing algorithms for smart home appliances and video analytics.
- Worked on applications including face recognition, fire/smoke detection, and virtual makeup using classical and deep learning techniques.

K4Tech, Internship in Machine Learning — Milan, Italy
Machine Learning Intern

- Developed ML algorithms in Matlab/C for computer vision applications, including face detection/recognition and surveillance.
- Built and tested prototypes for real-time video analytics and monitoring systems (e.g., Pantograph Monitoring System).

SELECTED TECHNICAL PROJECTS

MODELING NON-B DNA STRUCTURE IN NANOPORE DATA — UConn, CSE Dept. 2023 – Present

- Developed statistical and deep learning models to detect non-canonical DNA structures from nanopore translocation time signals.
- Performed large-scale data preprocessing and feature engineering from raw sequencing signals.
- Collaborating with deCODE Genetics to scale and automate the analysis pipeline for large sequencing data.

PROBABILISTIC POLYPLOID HAPLOTYPE ASSEMBLY (PHAPCOMPASS) — UConn, CSE Dept. 2022 – 2025

- Developed probabilistic haplotype assembly algorithms for diploid and polyploid genomes that explicitly model and propagate read assignment ambiguity.
- Designed graph-theoretic algorithms to perform statistical inference and uncertainty quantification over an exponential space of possible haplotype phasings.
- Built a simulation workflow for realistic auto- and allopolyploid genomes and DNA-seq data.
- Released four benchmark datasets and an open-source implementation of

BAYESIAN RECONSTRUCTION OF EXCISED mRNA — UConn, CSE Dept. 2021 – 2025

- Built a Bayesian modeling framework for reconstructing excised intron structures and testing differential splicing.
- Implemented inference and evaluation pipelines in Python using probabilistic programming.

FEATURE SELECTION FOR MICROARRAY CLASSIFICATION — Politecnico di Milano 2016 – 2018

- Designed and evaluated distributed feature selection algorithms based on distance correlation for high-dimensional gene expression data.
- Improved classification performance and interpretability of microarray-based models.

CLOUD APPLICATION PERFORMANCE MODELING (ATMOSPHERE) — Politecnico di Milano 2018 – 2019

- Combined Bayesian optimization and ML models to predict performance of GPU-intensive applications under different resource allocations.
- Reduced the number of required production samples while maintaining accurate performance estimates.

LEADERSHIP, TEACHING & SERVICE

Leadership & Mentoring

- Mentored undergraduate and junior graduate students on research projects in computational biology and ML.
- Contributed to organizing and running research group activities and collaborations.

Teaching

- Teaching Assistant, Computer Programming, University of Connecticut (2020).
- Lecturer, Siemens Workshop on Neural Networks and Deep Learning, Milan (2017).
- Teaching Assistant, Computer Networks and Internet Engineering, Shomal University (2005).

Professional Service

- Co-organizer, ASE 2025 Program Workshop (2025).
- Reviewer, Nature Communications (2025) — manuscript on non-B DNA structures.
- Program Committee Member, International Workshop on Semantics and the Web for Transport (2021–2023).
- External Reviewer, SAGE Open Journal (2021).
- Sub-reviewer, RECOMB (Computational Molecular Biology) conference (2019).

Selected Honors

- Predoctoral Fellowship Award, University of Connecticut (2022, 2023, 2025).
- Research Fellowship, Politecnico di Milano (2018).
- Graduated with honors in Computer Science, Politecnico di Milano (2018).

SELECTED PUBLICATIONS

**Deep statistical modelling of nanopore sequencing translocation times reveals latent 2023
non-B DNA structures**
M. Hosseini et al.
Bioinformatics (ISMB proceedings).

Bayesian Reconstruction and Differential Testing of Excised Introns 2025
M. Hosseini, D. McConnell, D. Aguiar.
Bioinformatics (accepted).

**pHapCompass: probabilistic haplotype assembly for diploid and polyploid genomes with 2025
uncertainty quantification**
M. Hosseini et al.
Submitted to *RECOMB* (under review). Code and datasets available at [GitHub](#).

**A Distributed Feature Selection Algorithm Based on Distance Correlation with an Ap- 2018
plication to Microarrays**
A. Brankovic, M. Hosseini, L. Piroddi.
IEEE/ACM Transactions on Computational Biology and Bioinformatics.

LANGUAGES

ENGLISH: Fluent
PERSIAN: Native Speaker

ITALIAN: Intermediate
ARABIC: Elementary