

Mirza Ahmadi

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SKILLS SUMMARY

- **Languages & Tools:** Python, R, Bash, Linux, High-Performance Computing (HPC), Git, HTML & CSS
- **Certifications:** [Machine Learning \(ML\) + Deep Learning \(DL\) - Compute Ontario](#) | [AI + Python - Harvard CS50](#) | [DL + Reinforcement Learning - CIFAR](#) | [Linear Algebra + Calculus - Coursera](#)

PERSONAL PROJECTS

- [Enhancing EEG Classification with GANs \(BrainHack 2024, SickKids\)](#) | [Demo](#) | *Python* Dec 2024
- Collected and cleaned EEG data from individuals performing two cognitive exercises, creating a labeled dataset
 - Trained a **Random Forest** model with **95%** accuracy using **Pandas**, **NumPy**, and **Scikit-Learn**
 - Generated 200 000 synthetic datapoints with a **GAN** to test if synthetic data can serve as a proxy for real data
 - Visualized results using Confusion matrices, ROC-AUC curves, loss plots, and learning curves with **Matplotlib**
- [Evaluating Algorithms for Gene Sequence Classification](#) | *R (Tidyverse)* Nov 2024
- Compared the performance of **Random Forest** and **Linear Regression** models in gene sequence classification
 - Visualized sequence length distribution, k-mer frequency proportions, and feature importance
- [Road Sign Categorizer](#) | *Python* May 2024
- Implemented a **convolutional neural network** to classify road signs into 43 categories with **90%** accuracy
 - Leveraged the libraries **TensorFlow**, **Scikit-learn** and **OS** for model development and data preprocessing
- [Venom Type Classifier](#) | [Demo](#) | *Python* Mar 2024
- Developed a **Support Vector Machine** model to predict venom type from protein data, achieving **94%** accuracy
 - Utilized **Scikit-learn**, **Matplotlib**, and **Pandas** for model building and visualization of PCA data

EXPERIENCE

- M. Sc. Research Project - University of Guelph** Guelph, ON | Jan 2024 - Present
- Developing an **ETL** pipeline to derive genome annotation insights, informing disease, variability, and evolution
 - Integrating **Python**, **R**, bioinformatics tools, and **ML** for data processing, sequence classification, and modeling
 - Using Compute Canada systems and **SLURM** to manage **HPC** jobs for genomic data processing and storage
 - Deploying the pipeline on a cloud platform to enhance efficiency and data quality for genomic researchers
- Evolutionary Biology Researcher - University of Guelph** Guelph, ON | May 2022 - Present
- Designed and executed an independent research report on snake venom evolution across 127 species
 - Analyzed proteomic and phylogenetic data in **Excel** and **R** to compile phylogenies and create graphical analyses
 - Authoring a research paper for publication to enhance our understanding of venom and complex trait evolution
- Teaching Assistant (1st year Biology Course) - University of Guelph** Guelph, ON | Sept 2022 - Dec 2024
- Led weekly seminars for 120 students and provided teaching support during two weekly lectures
 - Developed seminar material aligned with lecture topics, including genetics, evolution, and organismal biology
 - Evaluated and provided feedback on seminar worksheets to enhance understanding and learning outcomes

EDUCATION

- M. Sc., Bioinformatics & AI (Research-Based) - University of Guelph** Guelph, ON | Jan 2024 - Present
- Course: Machine Learning, Math for Machine Learning, Genomic Methods, Bioinformatics Software Tools
- B. Sc., Zoology Major with Minor in French - University of Guelph** Guelph, ON | Sept 2019 - April 2023