

# Akram Vasighi

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## Professional Summary

- PhD, Computer Science with strong background in machine learning, DNN programming & AI modeling.
- AI specialist with 5+ years of experience in implementing & deploying various industrial and academic projects.
- Successful background in leadership in developing large-scale AI models, leading to innovative patent.
- Proficient in Python, R, PyTorch, Scikit-learn, and various data analysis & machine learning frameworks.
- Strong publication records in computational biology & healthcare with valuable cross-functional team experience.
- Delivered courses, workshops, & presentations in top-tier conferences.
- Valued member of scientific journal reviewer community.

## Work Experience

### ML Engineer, App Developer, & Technical Lead, *Freelance*

08/2022 - Present

- Initiated an automating workflow for data integration, cleaning, manipulation, and preprocessing, effectively communicating with stakeholders to understand their needs and streamline complex processes (C#, SQL, ETL ).
- Led a patent project to drive innovative research, involving design & development of a new AI Agent-based tool to help researchers to perform and interpret complex single-cell RNA-seq (scRNAseq) analysis, especially those without programming expertise.
- Ensured testable, maintainable code, subprocess, and refactoring.
- Cross-functional team communication.

### Research Assistant, *School of Computer Science & Windsor Regional Hospital*

09/2024 - Present

- Collaborated with diverse multidisciplinary research teams to identify therapeutically targetable tumour-immune interactions in small cell lung cancer (SCLC), contributing to impactful translational research.
- Communicated findings clearly and effectively with both technical and non-technical audiences, bridging gaps between computational and clinical teams.
- Developed machine learning models to identify predictive biomarkers in SCLC (~95% accuracy), while adapting models based on ongoing feedback and validation results.
- Took the initiative in refining the analysis pipeline, ensuring reproducibility and maintainability for future team use, and contributing to the preparation of publication.

### Post-Doctoral Fellow, Data analyst, *School of Computer Science & Biomedical Science*

07/2024 - 03/2025

- Collaborated with researchers to analyze large-scale experimental genomic data (CRISPR screens), applying machine learning techniques and statistical analysis to detect biologically significant signals, with a focus on non-coding RNA regulation.
- Created an effective analysis pipeline to process, visualize, and analyze data, accelerating research timeline, running on GPU clusters.
- Work closely with researchers to understand and translate their needs into efficient computational solutions.

### Research Assistant, *School of Computer Science, University of Windsor, Canada*

01/2020 - 04/2024

- Demonstrated effective collaboration and mentorship; guiding undergraduate and graduate students to ensure individual and team success; clearly communicating complex research concepts, confidently leading collaborative projects, and applying strong problem-solving and writing skills to contribute to scientific publications and reports, while consistently meeting tight conference/presentation deadlines and research milestones.
- I designed and developed ML pipelines for large-scale data (including data gathering) to identify and characterize intercellular signaling networks as well as cell type annotation from single-cell transcriptomics data (scRNAseq).
- Trained deep learning models, embedding models, optimization, fine-tuning, and benchmarking, addressed the significant challenges of high-dimensionality and sparsity.

<b>University Instructor &amp; Graduate Assistant</b>	<b>01/2020 – 04/2024</b>
<ul style="list-style-type: none"> <li>• Provided supportive mentorship and patient communication, guiding students through complex concepts, facilitating engaging discussions, and collaborating with faculty to enhance student success.</li> <li>• Distributed Systems, Object-oriented programming using Java, Data Structure &amp; Algorithms, Machine Learning, Deep Learning, Computer Networks, AI for Games, Cyber Ethics, C++.</li> </ul>	
<b>Data Analyst, Academic Data Centre, Leddy Library, University of Windsor, Canada</b>	<b>01/2023 -04/2023</b>
<ul style="list-style-type: none"> <li>• Applied strong analytical thinking in integrating NLP techniques for historical text analysis, effectively collaborating with library staff and researchers to facilitate meaningful interpretation and accessibility of archival resources.</li> <li>• Helped instructors and speakers with 3 workshops for Codefest.</li> </ul>	
<b>Help Desk, Leddy Library, University of Windsor, Canada</b>	<b>05/2022 – 12/2022</b>
<ul style="list-style-type: none"> <li>• Provided approachable, patient, and clear communication while assisting students and faculty with technology troubleshooting, facilitating productive collaboration and enhancing user confidence in library IT resources.</li> </ul>	
<b>Python Teacher (Volunteer), Canadian Association for Girls in Science (CAGIS), Canada</b>	<b>01/2022 -04/2022</b>
<ul style="list-style-type: none"> <li>• As a best researcher in the Science department, I have been selected to teach Python to kids (7-12 years) using hands-on Python projects.</li> <li>• I helped to brainstorm, organize, and plan STEM events for members.</li> </ul>	

Education	
<b>Ph.D. in Computer Science</b> , University of Windsor, Canada	<b>01/2020 – 04/2024</b>
<b>M.Sc. in Computer Engineering</b> , University of Tehran, Iran	<b>09/2015 – 01/2017</b>

Technical Skills	
<b>Machine Learning &amp; AI, Data Analysis</b>	
<ul style="list-style-type: none"> <li>• Python   R   Pytorch   TensorFlow   Keras   Streamlit   LLMs (GPT)   API   CI/CD   RAG   AI Agents &amp; Automation</li> </ul>	
<b>Tools &amp; Platforms</b>	
<ul style="list-style-type: none"> <li>• Docker &amp; Containerization   HPC   Bash/Shell Scripting   Data Pipeline &amp; ETL   SQL   LaTeX   MCP   AWS</li> </ul>	
<b>Bioinformatics Tools</b>	
<ul style="list-style-type: none"> <li>• Scanpy   Seurat   Squidpy   DESeq2   SAMtools   BEDTools   IGV   GSEA   Cytoscape   STAR   MaGeCK   CasoFFinder   Crisprscore</li> </ul>	
<b>Leadership &amp; Product Management</b>	
<ul style="list-style-type: none"> <li>• Modularization &amp; Scalable Architecture   Model Optimization &amp; Fine-tuning   Documentation and Version Control (Git)  UI/UX Flow   Cross-functional Team Collaboration</li> </ul>	
<b>Familiar with:</b> Java   front-end technologies (HTML   React   TypeScript   Tailwind CSS   JavaScript   Responsive Design   GCP   Azure ML Studio, Reinforcement Learning techniques (RLHF), RecSys	

Selected Publications & Presentations	
[publications accessible via Google scholar profile ( <a href="https://scholar.google.com/citations?user=mJSJoqIAAAAJ&amp;hl=en">https://scholar.google.com/citations?user=mJSJoqIAAAAJ&amp;hl=en</a> ).]	
<ol style="list-style-type: none"> <li>1. <b>A. Vasighizaker</b>, S. Hora, R. Zeng, L. Rueda, “<a href="#">SEGCECO: Subgraph Embedding of Gene expression matrix for CELL cell COMMunication prediction</a>”, Briefings in Bioinformatics, <b>(2024)</b>.</li> <li>2. <b>Vasighizaker, A.</b>, Danda, S., &amp; Rueda, L. “<a href="#">Discovering cell types using manifold learning and enhanced visualization of single-cell RNA-Seq data</a>”. Scientific Reports, Nature Portfolio, <b>(2022)</b>.</li> <li>3. Nakul Pandya, Raymond Zeng, Biren Dave, <b>Akram Vasighizaker</b>, Swati Kulkarni, Ming Pan, Junaid Yousuf, Luis Rueda, “Identifying Therapeutically Targetable Tumor-Immune Interactions in Small Cell Lung Cancer”, WE-SPARK’s Health Research, Canada, <b>2025</b>.</li> </ol>	

Honors, Awards, & Hobbies	
<ul style="list-style-type: none"> <li>• Member of International Society for Children with Cancer.</li> <li>• Mitacs Research Training Award and NSERC Scholarship in AI.</li> <li>• Traveling, House Decoration, Embroidery, Cooking!</li> </ul>	