# Lin (Bill) Qi

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#### **PROFILE**

PhD-level AI specialist with over 8 years of experience spanning advanced ML research and the development of production-grade AI systems. Expertise in architecting Retrieval-Augmented Generation (RAG), multi-agent systems, and fine-tuning computer vision (CV) and large language models (LLMs). Track record of leading high-impact projects, supported by a strong publication history (contributed to 11 papers, 7 as lead/co-lead author).

# **EXPERIENCE**

CGI | Montreal, QC

### **Data Scientist, Generative AI**

June 2024 - Present

Architecting and deploying enterprise-grade generative AI solutions for a major public sector client, demonstrating expertise in tackling advanced RAG, agentic use cases, and model fine-tuning.

- Enterprise Conversational AI: Led the architecture and development of an enterprise-scale conversational AI assistant for compensation advisers. Achieved 91% benchmarked response accuracy by fine-tuning LLaMa 3.1 models with LoRA and serving quantized models using SGLang.
- Advanced RAG & Search: Engineered a sophisticated hierarchical vector search system, significantly outperforming traditional methods on complex policy documents. The system featured custom fine-tuned embedding and re-ranking models to enhance retrieval precision.
- Multi-Agent Systems: Developed a novel multi-agent OCR pipeline using OpenAl Agents and Prompt Flow, combining Azure Document Intelligence with GPT-4o. Achieved >99% accuracy in extracting structured data from scanned PDF documents.
- Infrastructure & MLOps: Automated the provisioning of secure, zero-trust generative AI architectures using **Terraform**. Built robust **CI/CD pipelines** to accelerate the testing and deployment of AI solutions in a cloud-native environment (Azure AI Foundry).

# McGill University, Department of Human Genetics | Montreal, QC PhD Candidate & Researcher Sep

September 2018 - February 2024

- Engineered scalable Python and TensorFlow data pipelines on HPC clusters (SLURM) to process terabyte-scale genomic and biological data, structuring it into graph-based representations for deep learning analysis.
- Developed a novel Graph Convolutional Neural Network to integrate complex, multi-modal genomic and clinical data for predictive modeling.
- Contributed to **11 peer-reviewed papers** (**7 as lead/co-lead author**) in top-tier journals, including *Biological Psychiatry*, and introduced novel methods for ML model selection.

#### Machine Learning Developer

July 2017 - September 2018

- Developed a classification model for engineer assignment for 1,000+ support engineers. Backend written in Python, deployed as a **Kubernetes** microservice pod.
- Prototyped a question-answering system using the RoBERTa model for answer span-classification and information retrieval methods on product documentation.
- Frontend Javascript (AngularJS) development for a ticket management application.

### **TECHNICAL SKILLS**

- Generative AI & LLMs: RAG (Retrieval-Augmented Generation), Fine-Tuning (LoRA), Agentic Workflows, LangChain, Prompt Flow, OpenAI Agents, Vector Databases (Pinecone, Azure AI Search, MongoDB vCore), SGLang, vLLM
- Al & Machine Learning: TensorFlow, PyTorch, Scikit-Learn
- Cloud & MLOps: Docker, Kubernetes, Terraform, CI/CD, Azure (Al Foundry, Prompt Flow, Al Search, Document Intelligence), AWS (EC2, Lambda, S3), Databricks, MLflow, Wandb
- Backend & Databases: Python, JavaScript, FastAPI, Flask, REST APIs, Azure Cosmos DB, PostgreSQL, Git, Linux/Unix
- Data Processing: Pandas, Numpy, Jupyter

#### **EDUCATION**

- PhD in Human Genetics (Statistical & Machine Learning focus) | McGill University, Montreal, QC
- Al in Healthcare Nanodegree | Udacity, Online
- Bachelor of Science, Microbiology & Immunology | McGill University, Montreal, QC

### **AWARDS & PUBLICATIONS**

# **Kaggle Competition Medals:**

- **Ubiquant Market Prediction** (Rank 48/2893) [Link]: Implemented an ensemble of neural networks for stock market prediction.
- RSNA Breast Cancer Detection (Rank 60/1,687) [Link]: Fine-tuned pretrained computer vision models (EfficientNet) and inference using NVIDIA TensorRT for medical image analysis.

**Selected Publications** (A full list of publications is available on my <u>Personal Site</u>):

- Qi, Bill, and Yannis J. Trakadis. "Graph Representation Learning for the Prediction of Medication Usage in the UK Biobank Based on Pharmacogenetic Variants."
  Bioengineering 12.6 (2025): 595. [Link]
- Qi, Bill, and Yannis J Trakadis. "Advancing Clinical Psychiatry: Integration of Clinical and Omics Data Using Machine Learning." Biological psychiatry (2023). [Link]
- Qi, Bill, Janani Ramamurthy, Imane Bennani, and Yannis J. Trakadis. "Machine learning and bioinformatic analysis of brain and blood mRNA profiles in major depressive disorder: A case-control study." American Journal of Medical Genetics Part B: Neuropsychiatric Genetics (2021). [Link]