

(604) 710-7454  
Vancouver, BC  
kevinyang10@gmail.com

# Kevin Yang

## Data Scientist / Engineer

keviny2.github.io/  
github.com/keviny2  
linkedin.com/in/keyang2

### SKILLS

<b>Tools and Languages</b>	Python, R, PostgreSQL, Bash, AWS, snakemake, GATK, HPC (SLURM, LSF), Snowflake, Jenkins, MLflow, Airflow, Docker, tox, Git, Jupyter Notebook, Visual Studio, RStudio, pgAdmin, Vim, JIRA, CI/CD
<b>Packages</b>	PyTorch, NumPy, pandas, polars, huggingface, pysam, scikit-learn, statsmodels, SciPy, duckdb, psycopg, matplotlib, seaborn, altair, NumPyro, pytest, ggplot, dplyr, samtools
<b>Machine Learning</b>	Neural Networks/Deep Learning, Natural Language Processing (NLP), Transformers, Bayesian Networks, Computer Vision, XGBoost, Regression, Clustering and Classification, Decision Trees, Time Series, SVM

### TECHNICAL EXPERIENCE

#### Consultant @ Amaris Consulting

##### Senior Data Engineer

Genentech - A Member of the Roche Group

05/2023 — Present

San Francisco, CA (Remote)

- Generated a training set of over 35 million natural language sentences from single-cell metadata with Large Language Models (LLMs) for a multi-modal neural network model blending scRNA-seq data and structured metadata to characterize cells.
- Enhanced the performance of Large Language Models (LLMs) inference by over 1000 fold through strategic optimization leveraging NVIDIA's GPU Cloud (NGC), model quantization techniques, and efficient batch inference processes.
- Spearheaded the development of a software tool for accessing, querying, and analyzing a corpus of single-cell datasets, spanning 250 million cells with individual datasets exceeding 300GB, accelerating research by reducing metadata harmonization needs.
- Implemented an ETL workflow to migrate a 1TB single-cell metadata corpus from the cloud into an analytical database, resulting in a resource cost reduction of 70% and improved query speeds of 300%.
- Applied bioinformatics-specific Large Language Models to standardize over 18 million annotation terms, using biological ontologies as references, which encompassed over 20% of the entire single-cell database.

##### Data Scientist Intern

Intact Financial Corporation

01/2023 — 04/2023

Vancouver, BC

- Leveraged machine learning algorithms to optimize premium pricing, increasing profits by over 50%.
- Reduced feature engineering time by 70% using cluster computing and parallelization tools.
- Cut manual labor by 90% by developing automated pipelines for data ingestion, analysis, and model training.

##### Machine Learning Research Assistant

BC Cancer Research Centre

05/2020-12/2020

Vancouver, BC

- Implemented a clustering algorithm combining DNA-methylation and RNA-expression on pan-cancer datasets over 250GB in size.
- Refactored a pipeline to analyze high-dimensional DNA-methylation data, resulting in a >75% reduction in memory requirements through efficient data analysis and wrangling.

### PROJECTS (SEE GITHUB)

#### SummarizedExperiment

- Contributed methods to combine advanced data structures optimized for processing datasets exceeding 500GB.
- Developed flexible subsetting methods for data analysis, increasing developer capacity by 20%.
- Authored comprehensive unit tests, achieving a code coverage of >90%.

#### Bayesian Probabilistic Graphical Model for Early Cancer Detection

- Integrated genomic variant data from both liquid and solid tissue biopsies into a machine learning model, successfully reducing error by 75% compared to state-of-the-art methods for inferring cancer population and subpopulation structure.
- Developed data simulation pipelines, which downsample Whole Genome Sequencing (WGS) datasets, to generate realistic high-dimensional biological data from liquid and tissue biopsies for benchmarking of machine learning model performance.

#### 3D Image Reconstruction of Cancer Tumours

- Decreased validation error by 60% for spatial tissue analysis by implementing two cutting-edge deep learning models.
- Mitigated overfitting through weight decay and optimized loss functions, resulting in a 10% reduction in validation error.
- Accelerated neural network training over 100x using GPUs with CUDA in PyTorch.

### EDUCATION

**MSc. Bioinformatics - GPA: 4.0, The University of British Columbia**

2021-2022

- Canada Graduate Scholarships - Master's (CIHR) (\$17 500 over 1 year)

**BSc. Computer Science & Statistics - GPA: 3.9, The University of British Columbia**

2015-2020

- Dean's Honour List