

# Khachatur Dallakyan

Room 1507, Studentbacken 23 / 1507, Stockholm, Sweden

+46 734836391 | [khachatur.dallakyan@gmail.com](mailto:khachatur.dallakyan@gmail.com) | <https://khachdallak02.github.io/>

## EDUCATION

<b>Karolinska Institute, Stockholm University</b>	Sweden
Joint Master's Programme in Health Informatics	2024-2026 (expected)
Awards: The Swedish Institute Scholarship for Global Professionals	
<b>The Chinese University of Hong Kong</b>	Hong Kong
Bachelor of Sc. in Artificial Intelligence: Systems and Technologies (GPA: 3.29 / 4.00)	2020-2024
Awards: Second class upper division with Honours, Belt and Road Scholarship	
<b>International Baccalaureate Diploma Programme at Quantum College</b>	Armenia
Total points: 37/45	2018-2020
Subjects: Mathematics, Computer Science, ITGS, Armenian, Economics, English.	

## RESEARCH AND WORK EXPERIENCE

<b>Exploring the Enteric Nervous System (ENS) through Single-Cell Transcriptomics</b>	2025
Main objective of the research is to find regulatory mechanisms in embryonic ENS, by exploring spatio-temporal differences for ENS cells, namely progenitors, neuroblasts and SCPs between regions and timepoints.	
<ul style="list-style-type: none"><li>Finetuned various foundation models (scGPT, cell2sentence) with custom learning objectives to create region and timepoint classifiers.</li><li>Utilized the classifiers to simulate in-silico perturbation of various candidate genes and the attention architecture to study gene-to-gene relations.</li><li>Applied SCENIC to infer gene regulatory networks and identify transcription factor motifs, though database-derived predictions produced many false positives. As a next step, we plan to integrate scATAC-seq data from our lab to improve motif detection together with conventional cell fate prediction models like RegVelo to improve in-silico perturbation predictions.</li></ul>	
<b>Biopsy QR Code Management System</b>	2025
<ul style="list-style-type: none"><li>Developed biopsy management system for Freiburg's medical center.</li></ul>	
<b>Benchmarking Single Cell Foundation Models (FM) for Retrieval</b>	2024-2025
Benchmarking traditional methods and FMs across diverse biological tasks.	
<ul style="list-style-type: none"><li>Benchmarked retrieval accuracy, diversity, biomarker identification, and novel cell type detection across multiple platforms, species, and technologies.</li><li>Evaluated and compared traditional methods with foundation models (scGPT, Geneformer, scMulan, UCE, SCimilarity, scFoundation), designing benchmarks for complex scenarios with ambiguous or incomplete ground truth.</li></ul>	
<b>Prediction of Cellular Morphology with Image-to-image Translation Models</b>	2023-2024
<ul style="list-style-type: none"><li>To benchmark MorphDiff established baseline using the DMIT and DRIT++ models.</li><li>Adapted the model architectures to process 5 channel cellular morphology imaging data.</li></ul>	
<b>Internship at Centre for Perceptual and Interactive Intelligence (CPII)</b>	2023
AI research company targeting sectors of healthcare, urban services, and reindustrialization	
<ul style="list-style-type: none"><li>Developed a prototype for speech-based interaction with LLMs without relying on ASR or TTS.</li><li>Utilized speech representations from Hubert pretrained model to finetune LLAMA.</li></ul>	
<b>Internship at ONEFi (iPYGG)</b>	2022
Fintech company aiming to connect Asia-Pacific banking into a single application.	
<ul style="list-style-type: none"><li>Developed ML models for transaction classification and description summarization.</li></ul>	

## PREPRINTS AND JOURNAL PUBLICATIONS

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### Prediction of Cellular Morphology Change Under Perturbations with Transcriptome Guided Diffusion Model [\[paper\]](#)

2025

Xuesong Wang, Yimin Fan, Yucheng Guo, Chenghao Fu, Kinhei Lee, [Khachatur Dallakyan](#), Yaxuan Li, Qijin Yin, Yu Li, and Le Song

### SCMBench: Benchmarking Domain-specific and Foundation Models for Single-cell Multi-omics Data Integration [\[preprint\]](#)

2025

Yixuan Wang, Yimin Fan, Xuesong Wang, Tingyang Yu, Yongshuo Zong, Xinyuan Liu, Meitong Liu, Qing Li, Kin hei Lee, [Khachatur Dallakyan](#), Junjie Huang, Gengjie Jia, Jiao Yuan, Ting-Fung Chan, Xin Gao, Irwin King, Yu Li

## EXTRACURRICULAR ACTIVITIES

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### KI-Mayo 30th annual scientific meeting

2024

- Helped to organize main and breakout sessions for “AI and health” section.

### Videogame modding

2024

- Modified WW2 videogame resource trading algorithm for AI to prioritize historical allies and trade partners.

### Armenian Red Cross Society (ARCS)

2019-2020

- Volunteered as part of the “Social Support and Health Care” team.

### Global Innovation Forum

2019

- Volunteered for GIF during Aurora Humanitarian Initiative
- Helped to organize a talk about the scientific computing framework Torch and its use cases.

### Competitive programming

2017-2019

- Codeforces: <https://codeforces.com/profile/khachdallak>
- Developed and tested competitive programming problems for Armenian SPOJ: <https://am.spoj.com/problems/TIGRANMEC/>

## ACHIEVEMENTS

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### The Swedish Institute Scholarship for Global Professionals

2024

- Awarded for demonstrated academic excellence and strong leadership potential.

### CUHK Data Hack 2024

2024

- Champion team during the Data Solutions for Social Good Hackathon.
- Implemented random forest algorithm for early detection and prevention of drug abuse.

### “Belt and Road” scholarship

2020

- Scholarship for outstanding new students from the Belt and Road Region countries.

### “Best student” state prize

2018

- Republic of Armenia State Prize in Information Technologies.

### “Algorithmization Methods” Programming Contest

2017

- Second and third places (top 0.5%) in Autumn and Winter Cup contests respectively.