# Anya Korsakova

anya@calicolabs.com | codehelix.ai | github.com/anyakors

#### Skills

**Programming:** python, TensorFlow, pyTorch, bash, C++, openCV

Tools: slurm, Google Cloud Platform, Docker, LaTeX

**Machine Learning (ML):** transformers, variational autoencoders (VAEs), sparse autoencoders (SAEs), energy-based models, convolutional neural networks (CNNs), decision trees, input importance attribution

# **Experience**

Postdoctoral researcher, Calico Life Sciences LLC (Alphabet) – San Francisco, CA

May 2023 - Present

- ML architecture design and data preparation for biological ML models.
- Enabled genetic variant effect prediction for structural variants using align-and-stitch method at inference and created new benchmarks for models while improving AUROC performance on existing benchmarks by 9% [1].
- Built a framework for mechanistic interpretability and extracted sequence insights from DNA sequence-based ML models using SAEs.

## Postdoctoral Researcher, Cancer Science Institute – Singapore

Oct 2022 - May 2023

- Devised and implemented a probabilistic mutational signature assignment method ALPS gitlab.com/PittGenomics/alps.
- Collaborated on building ensemble approaches to mutational signature assignment [2].

#### Quantitative Finance Developer, Juniper Investment Pte Ltd – Singapore

Feb 2022 - Sept 2022

• Programmed high yield multi-timeframe, multi-instrument trading bots for foreign exchange markets in C++ (MQL5) using technical and fundamental analysis.

## PhD Scholar, Nanyang Technological University – Singapore

Aug 2017 - July 2022

Developed machine learning frameworks for structural and cellular biology.

- Built a G-quadruplex DNA structure prediction model using convolutional neural networks and auxiliary inputs that achieved state-of-the-art performance [3].
- Implemented data pipelines and devised the architecture of an RNA splicing prediction framework with an energy-based model augmented with RNA-binding protein levels [4].

#### Junior Researcher, Lebedev Physical Institute RAS – Moscow, Russia

Jan 2014 – Aug 2017

• Modeled diffusive-thermal instabilities in hydrogen-air flames via solving systems of partial differential equations in Mathcad and MATLAB; reported instability onset regimes in journal publications [5].

#### Junior C++ Developer, NRNU MEPhI - Moscow, Russia

Dec 2012 - Dec 2013

• Implemented an algorithm for eye iris recognition and tracking in live video stream with C++ and openCV.

#### **Projects**

### **Attractors in Neural Networks**

codehelix.ai/blog/attractors

• Visualizing and analyzing attractors in feedforward neural network circuits, featured in Towards Data Science.

#### Education

Nanyang Technological University, Singapore – PhD in Biophysics	July 2022
NRNU MEPHI, Moscow, Russia – MS in Applied Mathematics and Physics (First Class Honors)	June 2016
NRNU MEPhI, Moscow, Russia – BS in Applied Mathematics and Physics	June 2016

## **Awards**

#### **Peer Review**

Nature Machine Intelligence ISSN: 2522-5839, reviewer, 2024

# **Research Internships**

Wan Lab of RNA structuromics, Genome Institute of Singapore

Mar 2021 - June 2021

RNA G4 structure probing in long synthetic RNA with NAI-SHAPE and nanopore sequencing.

Lab of Biology and Applied Pharmacology, École Normale Supérieure Paris-Saclay

June 2018 – Aug 2018

Geometrical improvement of NMR RDC data usage for G4 structure resolution.

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**Institute of Technical Thermodynamics**, Karlsruhe Institute of Technology.

Oct 2015 - Nov 2015

# **Teaching and Supervision**

NTU School of Physical and Mathematical Sciences, Singapore

• Undergraduate student mentor

2018, 2020, 2021

Supervised students on an ML-genomic project and a nanopore sequencing project.

• From synthesis to quantification of DNA using UV absorption, circular dichroism and fluorescence spectroscopy (PH3399)

2019, 2020

• Silicon Charge Particle Detectors (PH3199)

2018, 2019, 2020

• Fabrication Laboratory (PH3199)

#### **Conference Presentations**

- Korsakova A et al., "Shift augmentation for improved indel scoring in DNA sequence-based ML models", ASHG 2024 Annual Meeting, Denver.
- Srivastava D, Yuan H, Korsakova A et al., "Borzoi-guided fine mapping improves variant and gene prioritization in GWAS", ASHG 2024 Annual Meeting, Denver.

#### **Selected Publications**

- [1] *Korsakova A et al.*, "Shift augmentation for improved indel scoring in DNA sequence-based ML models" to be submitted to Nat Methods.
- [2] Wu AJ, Perera A, Kularatnarajah L, Korsakova A, Pitt JJ, "Mutational signature assignment heterogeneity is widespread and can be addressed by ensemble approaches" in Briefings in Bioinformatics, 2023, DOI:10.1093/bib/bbad331.
- [3] *Korsakova A, Phan AT*, "Prediction of G4 formation in live cells with epigenetic data: a deep learning approach" in NAR Genomics and Bioinformatics, 2023, DOI:10.1093/nargab/lqad071.
- [4] Chan A, Korsakova A et al., "RNA alternative splicing prediction with discrete compositional energy network" at ACM CHIL '21, DOI:10.1145/3450439.3451857.
- [5] **Google Scholar:** scholar.google.com/citations?user=5A3VUGMAAAAJ.