

Anna Toidze

MASTER'S STUDENT IN COMPUTATIONAL BIOLOGY AND BIOINFORMATICS

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Education

Msc in Computational Biology and Bioinformatics at ETH Zürich, D-BSSE

- **ESOP Scholarship:** top 3 % of all incoming MSc students at ETH Zürich

Zürich, Switzerland

Sept 2023 - now

BSc in Biochemistry at Ruhr University Bochum, Faculty of Chemistry and Biochemistry

- **Wilke Award:** ranked 1st in Bachelor's degree in Biochemistry, GPA 1.0/1.0

Bochum, Germany

Oct 2019 - Sept 2022

Bioinformatics for Biologists, Additional Education at Bioinformatics Institute

- **Award:** ranked 1st out of 53 of the year, GPA 5.0/5.0

Saint Petersburg, Russia

Sept 2021 - May 2022

High School at Authorized School No. 6

- **International Biology Olympiad:** Bronze medals in 2018 and 2019

Tbilisi, Georgia

Sept 2016 - May 2019

Research

Internship in pRED Ophthalmology at Roche at Roche, pRED

- scRNA analysis in a pre-clinical mouse model of subretinal fibrosis in Python and R

Basel, Switzerland

Apr 2025 - Now

Internship in Laboratory of Prof. Dr. Sebastian Bonhoeffer at ETH Zürich, D-USYS

- Dynamically modeling the spread of plasmids carrying antibiotic-resistance genes in R using ODEs

Zürich, Switzerland

Nov 2024 - Mar 2025

Internship in Laboratory of Prof. Dr. Andreas Wagner and Dr. Aleks Nivina at UZH, IEU

- Stochastically modelled the recombination of antibiotic-producing polyketide system in *Streptomyces* in Python

Zürich, Switzerland

Sept 2024 - Oct 2024

Laboratory Technician in Laboratory of Prof. Dr. Fyodor Kondrashov at OIST, Japan

- Created *E. coli* mutant libraries of GFPs, sorted them using FACS, improved prediction of function proteins through ML
- Devised and implemented a populations genetics model for epistasis in Rust
- Guided an intern in preparation of *E. coli* GFP library, sorting using FACS and successful preparation for sequencing

Onna, Japan

Oct 2022 - Aug 2023

Internship in Laboratory of Prof. Dr. Fyodor Kondrashov at ISTA, Austria

- Cloned and purified green fluorescent proteins with varying sequence divergence from the wild-type sequence
- Experimentally assessed the physical stability of the proteins and correlated it with the mutational robustness using Python ([GitHub](#))

Klosterneuburg, Austria

Apr 2022 - Sept 2022

Internship in Laboratory of Prof. Dr. Maria Logacheva at Bioinformatics institute

- Investigated the expression patterns of egg-cell specific genes in crops using Python, R, phylogenetics tools

Online

Feb 2022 - May 2022

Internship Laboratory of Prof. Dr. Ute Krämer at RUB, MGPP

- Implemented a pipeline to investigate the contribution of CNV to metal tolerance in *A. halleri* using bioinformatics tools, R

Bochum, Germany

Jan 2021 - Jan 2022

Student at School of Molecular and Theoretical Biology

- Investigated intracellular parasites *B. mallei* using Python, bioinformatics tools in Laboratory of Prof. Dr. Mikhail Gelfand in 2019

Barcelona, Spain - Online

Aug 2019

Teaching and Volunteering

Teaching Assistant in Computational Systems Biology at ETH Zürich

- Writing lecture notes for computational systems biology class under Prof. Dr. Jörg Stelling

Zürich, Switzerland

Oct 2024 - Now

Teaching Assistant in Maths and Supervisor in Laboratory practical at RUB

- Supervised tutorial classes, graded assignments and exams for the maths courses for first-year (bio-)chemistry students
- Monitored experiments and graded reports of 20 students the introductory chemistry practical

Bochum, Germany

Oct 2020 - Sept 2022

Staff member and Mentor at School of Molecular and Theoretical Biology

- Introduced machine learning, Python, Git to high-school students in Laboratory of Prof. Dr. Olga Kalinina in 2022
- Mentored 12 students throughout their stay at the online summer school and beyond, and organized school-wide social activities

Online

Aug 2020 - Aug 2022

Skills and Honors

Experimental Bioinformatics Conferences

Cloning; bacterial culture; electrophoresis; Western Blot; PCR; extractions; transformation; microscopy; FACS; Benchling
Python; R; version control; HPC computing; NGS analysis; phylogenetics/-dynamics; data analysis; ML; protein modeling
SMBE 2025, Young Investigator Award: *Hiking Peaks and Valleys of GFP Fitness Landscapes for Data-Driven Protein Design*

Summer Schools

Guarda Summer School in Evolutionary Biology (2024), School of Molecular and Theoretical Biology (2019-2022)

Languages

Georgian (native), English (fluent), German (fluent), Russian (fluent)