Jakub Giezgała

Bioinformatics and Systems Biology Student at the University of Warsaw

Date of birth: 08.12.2004

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Biology and computer science - life and computation - are related. I am confident that great discoveries await those who seek them at their interface.



Languages

- Polish (Native speaker)
- English B2
- German A2

Hard Skills

- Programming languages: Python, C++, Java, R, MATLAB
- Web development: HTML, CSS, Django
- Machine Learning in Python: NumPy, Scikit-learn, SciPy
- Deep learning in Python: PyTorch, Torchvision, JAX, Pyro, Deeptime
- Data analysis and visualization:
 - Python: pandas, matplotlib, plotly, seaborn, Pillow, PyQt6, MEFISTO, Scanpy
 - R: ggplot2, tidyr, plotly, qqplotr, caret, shiny
- Proficient in Microsoft Office
- Driving license Category B

Soft Skills

- Willingness for self-development and perfection
- Analytical thinking
- Teamwork skills
- Good time management
- Honesty and reliability

Courses

- NVIDIA Fundamentals of Deep Learning
- SANTANDER Fundamentals cybersecurity with Poznan University of Technology

Education

University of Warsaw

Faculty of Mathematics, Informatics and Mechanics Degree Course: Bioinformatics and Systems Biology

Third-year student

XVIII High School in Warsaw named after Jan Zamoyski

Math–Physics–Informatics profile Graduated with honors in 2022

Experience

2021 - Present

Private tutoring in mathematics, physics and informatics (elementary and high school level, including final exam preparation)

2022 - Present

Active membership of the Bioinformatics Student Club (popularisation of science at research picnics: $DOKO\ 2023/2024/2025$)

Hackathons:

• QL Future Hackathon 2024 - Tech category

Scientific Conferences:

- Biomeeting, 24–26 May 2024, Poster Abstracts Using CNNs and ATAC-STARR-seq uncover predictive opportunities in regulatory genomics & From Cooperation to Competition: revealing stable correlations between features in the evolutionary game theory
- International Student Biomedical Conference (*Biofusion*), 28-30 March 2025 Bioinformatics workshop on sc-RNA data analysis pipeline