# Oleksandr Khoroshevskyi

## Python Software Engineer

## Personal information

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github.com/khoroshevskyi

### Skills

- Python (FastAPI, SQLAlchemy,
- scikit-learn, Pytest, async and others),
- · R, Bash, Rust
- Database Management: PostgreSQL, MongoDB
- Git, Docker, AWS
- Linux and HPC proficiency
- Pipeline development

Languages

English – pre-advanced
German – intermediate
Polish – advanced
Dutch – beginner

Russian – advanced
Ukrainian– native

## Profile

Scientific Programmer with more than four years of hands-on experience in web development, pipeline development, data processing, and manipulation, primarily utilizing Python. I am eager to embrace new challenges.

## Work Experience

## **Scientific Programmer**

University of Virgnia, USA

Sheffield lab of computational biology. databio.org

Jul 2021 - now

- Software engineering (Tech stack includes: Python, Rust, Bash, Docker, PostgreSQL, AWS)
- · Conducting data engineering and database construction
- RESTful API Development and optimization
- Development, maintenance, and optimization of bioinformatic pipelines
- Public projects: https://pephub.databio.org, https://bedbase.org
- Experience in conducting job interviews
- Project mentoring

## **Test Automation Engineer**

Nokia, Wroclaw, Poland

Jun 2020 - Feb 2021

- Writing and maintenance of software and test scripts (Python, Bash, Robot).
- Software verification.
- Configuration of automated regression test cases.

### Software developer, Internship

Webmil, Ivano-Frankivsk, Ukraine

Jul - Aug 2018

• Summer Internship in programming.

## Education

# **Master of Science in Bioinformatics**

Wroclaw University of Environmental and Life Sciences, Poland 2019-2023

**Bachelor in Bioinformatics** 

Wroclaw University of Environmental and Life Sciences, Poland 2016-2019

Erasmus+ exchange program

University of Lisbon, Portugal 2021

Erasmus+ exchange program

KU Leuven, Belgium 2018-2019

## Selected Publications

- Khoroshevskyi, O., LeRoy, N., Reuter, V. P., and Sheffield, N. C. (2023). GEOfetch: A command-line tool for downloading data and standardized metadata from GEO and SRA. Bioinformatics 39, btad069. doi: 10.1093/bioinformatics/btad069
- Xue B, Khoroshevskyi O, Gomez RA and Sheffield NC (2023), Opportunities and challenges in sharing and reusing genomic interval data. Front. Genet. 14:1155809. doi: 10.3389/fgene.2023.1155809