



Aleksandar Anžel

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Born 06.08.1995.

 AAnzel

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 AleksandarAnzel

 <https://aanzel.github.io>

WORK EXPERIENCE

July 2022 – present

Technical Lead

eMedicals Healthtech GmbH, Frankfurt am Main

- Mentor and guide team members to ensure the best productivity and overall satisfaction. Help with hiring new team members. Assess current technologies and advise on new ones the team could use. Determine potential risks and propose ways to mitigate them. Assert data compliance and ensure its proper adoption. Use Scrum methodology to help our product owner, and act as a Scrum Master. Design and implement application-critical parts of the kidiDNA and nephroDS. Review code of other team members and promote collaboration and engagement. Provide Architectural and Design directions for the company's products. Adhere the development to the multiple Software-as-a-medical-product standards and conduct internal audits.

October 2021 – July 2022

Data Scientist

eMedicals Healthtech GmbH, Frankfurt am Main

- Use analytical, statistical, and programming skills to collect, analyze, and interpret large medical and biological data sets. Manage algorithm development, machine learning workflows, and statistical techniques to produce solutions to problems quickly. Closely work with university partners (i.e., Technische Universität Darmstadt) and research organisations (i.e., Fraunhofer-Gesellschaft zur Förderung der angewandten Forschung e.V.) to improve and further develop algorithms relevant to the company.

December 2020 – present

Research assistant

Heider Lab, Philipps-Universität Marburg, Marburg

- Develop Bioinformatics and ML pipelines for Omics problems. Adapt existing and create new visualization methods for biological and medical data sets. Organize workshops and mentor multiple undergraduate and graduate students.

MOSLA, Philipps-Universität Marburg, Marburg

- Design, implement and evaluate automatic workflows for information storage systems based on the molecular storage media. Adapt existing and create new visualization methods and UIs for novel data storage technologies.

EDUCATION

2020 – present

Ph.D. degree in Computer Science

Philipps-Universität Marburg

August 2021

OxML summer school participant

Machine Learning summer school, University of Oxford

2018 – 2020

Master's degree in Mathematics

Module: Computer Science and Informatics

Faculty of Mathematics, University of Belgrade

- Average grade: 10.00 (out of 10.00)
- Thesis: *Determining protein N-glycosylation with machine learning methods*

2014 – 2018

Bachelor's degree in Mathematics

Module: Computer Science and Informatics

Faculty of Mathematics, University of Belgrade

- Average grade: 8.66 (out of 10.00)

SKILLS

Languages

Serbian — Native proficiency
English — Full professional proficiency

- Cambridge English: First (FCE): upper intermediate (B2 in CEFR)

German — Elementary proficiency
French — Elementary proficiency

Computer Science

Software Development

- C, Python, C++, Java, MATLAB, Shell, Haskell, Assembly IA-64, Assembly ARM-32

Machine Learning

- Keras, Tensorflow, Scikit-learn

Data Management

- SQL

Additional skills

Scientific and Cluster Computing, High-Level Visualization, CI/CD, Containerization (Docker), Azure, Agile development, Scrum, Code verification and validation

Document manipulation

LaTeX, Libre Office Suite, Microsoft Office Suite

Soft skills

- Excellent organizational and communication skills
- Ability to work collaboratively with people at all professional levels
- Thoroughness, with rigorous attention to both detail and quality

PROJECTS

Bioinformatics

- *Determining protein N-glycosylation with machine learning methods*
- *Modification and analysis of UPGMA algorithm while using different metrics*
- *MOVIS: A Multi-Omics Software Solution for Multi-modal Time-Series Clustering, Embedding, and Visualizing Tasks*
- *CMANGOES — Carbon-based Multi-level AtomicNeighborhood EncodingS*

Computer Science

- *Finding Waldo using various Machine Learning methods*
- *Image modification and correction with Python*
- *Determining integer variable ranges using Abstract Interpretation in C++ (LLVM, Clang)*
- *AVL trees in C programming language*

SELECTED EVENTS

- 2022 • *The 1st International Conference on Data Storage in Molecular Media*. Virtual. (attendee, co-host)
- 2021 • *IEEE Visualization Conference (VIS)*. Virtual. (attendee)
 - *Bio+Med+Vis Spring School*. Virtual. (attendee)
 - *Symposium on Interdisciplinary Bioinformatics and Biomedical Data Science (IBBMDS)*. Marburg, Germany. (presenter)
- 2020 • *IEEE Visualization Conference (VIS)*. Salt Lake City, Utah, USA. (attendee)
 - *Eurographics & Eurovis (EGEV)*. Norrköpping, Sweden. (attendee)

TEACHING

- 2022
- Guest lecture, *Molecules as storage media for long-term data storage*. Faculty of Mathematics, University of Belgrade. Belgrade, Serbia. (presenter)
 - Group student project (Projektarbeit), *DNA Storage Encodings*. Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany. (co-organizer, tutor)
 - Seminar, *Information Theory Tools for Visual Computing*. Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany. (co-organizer, presenter)
- 2021
- Group student project (Projektarbeit), *DNA Storage Encodings*. Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany. (co-organizer, tutor)
 - Seminar, *Information Theory Tools for Visual Computing*. Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany. (co-organizer, presenter)
 - Lecture, *Data Visualization*. Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany. (collaborator)

ADVISING & SUPERVISION

- 2022 – present
- Jing Chen, B.Sc., Bioinformatics (BI). Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany.
- 2021 – present
- Fabio Rougier, M.Sc., Computer Science (CS). Department of Mathematics and Computer Science, University of Marburg. Marburg, Germany.

PUBLICATIONS

- 2022
- Anžel, A., Heider, D., & Hattab, G. (2022). *MOVIS: A multi-omics software solution for multi-modal time-series clustering, embedding, and visualizing tasks*. Computational and Structural Biotechnology Journal, 20, 1044–1055. 10.1016/j.csbj.2022.02.012
- 2021
- Anžel, A., Heider, D., & Hattab, G. (2021). *The Visual Story of Data Storage: From Storage Properties to User Interfaces*. Computational and Structural Biotechnology Journal, 1(1), 1. 10.1016/j.csbj.2021.08.031

ADDITIONAL INFORMATION

Driving licence Category B (cars)

Interests Technology, Research, Computer Science, Bioinformatics, Linux, FOSS, Science Fiction, Fantasy, The Matrix, Video games, Hiking