Ekaterina Antonenko

Education

since 2020 **École Polytechnique**, *PhD candidate*.

Data Science and Mining (DaSciM) team, Laboratoire d'informatique (LIX)

Scientific advisor: Jesse Read

2014–2016 British Higher School of Art & Design, 2D-animation.

2012–2014 Moscow Bioinformatics School.

2009–2014 Moscow State University, Diploma in Mathematics.

Employment

since 2020 Data scientist, Digitalent (Paris, France).

2017–2019 Math instructor, Russian School of Mathematics (New York, USA).

2013–2016 Analyst, Moscow Center for Continuous Mathematical Education (Moscow, Russia).

Preprints

- 1. E. Antonenko, A.Carreño, J. Read, *Missing Value Imputation with Multi-label Random Forests*
- 2. M. Konnova, E. Antonenko, J. Read, *Missing Value Imputation for Genomics Data using a Sequence Based Generative Adversarial Network*
- 3. E. Antonenko, M. Mechenich, R. Beigaite, I. Zliobaite, J. Read, *Probabilistic prediction of land cover at the global scale: modeling absence of human activity*

Publications

- 1. E. Antonenko, J. Read, *Chains of Autoreplicative Random Forests for missing value imputation in high-dimensional datasets*, [Best paper award], Multi-Label Learning workshop at the ECML conference, 2022, https://arxiv.org/abs/2301.00595/.
- 2. E. Antonenko, J. Read, *Multi-modal ensembles of regressor chains for multi-output prediction*, Advances in Intelligent Data Analysis XXI 21st International Symposium, IDA 2022, 2022.
- 3. V. Ivanenko, E. Antonenko, M. Gelfand, J. Yager, F. Ferrari, Changes in segmentation and setation along the anterior/posterior axis of the homonomous trunk limbs of a remipede (Crustacea, Arthropoda), PeerJ, 2016, https://peerj.com/articles/2305/.

Talks

- 1. Chains of Autoreplicative Random Forests for missing value imputation in high-dimensional datasets, Multi-Label Learning workshop: current trends and open challenges, ECML PKDD 2022, Grenoble, France, September 2022.
- 2. *Multi-modal ensembles of regressor chains for multi-output prediction*, Intelligent Data Analysis XXI 21st International Symposium, IDA 2022, Rennes, France, April 2022.

Teaching

• CSE204 Machine Learning (Bachelor Programme), Teaching assistant, École Polytechnique, Spring 2022, 2023.

Mentoring students

Margarita Konnova (École Polytechnique), Bachelor thesis "Missing Value Imputation for Genomics Data using a Sequence Based Generative Adversarial Network", coadvised with Jesse Read, 2022.

Additional training

- Data Science: Multiple Imputation in Practice, *Utrecht University, The Netherlands*, 2022.
- Introduction to quantitative genetics, MIPT, Russia, 2021.
- Summer School on Machine Learning in Bioinformatics, HSE University, Russia, 2020.

Completed online-courses

- Al for Medicine Specialization, *Deeplearning.ai*, 2021, *Certificate*.
- Deep Learning Specialization, Deeplearning.ai, 2020, Certificate.
- Machine Learning, Stanford University, 2019, Certificate.
- Molecular Biology Part 3: RNA Processing and Translation, Massachusetts Institute of Technology, 2019, Certificate.
- Molecular Biology Part 2: Transcription and Transposition, Massachusetts Institute of Technology, 2019, Certificate.
- Molecular Biology Part 1: DNA Replication and Repair, Massachusetts Institute of Technology, 2018, Certificate.
- Quantitative Biology Workshop, Massachusetts Institute of Technology, 2018, Certificate.
- Introduction to Biology The Secret of Life, Massachusetts Institute of Technology, 2018, Certificate.
- Bioinformatics Algorithms (Part 1), University of California, San Diego, 2014, Certificate.

Additional skills

Programming: Python, R, MatLab

• Languages: Russian (native), English (fluent), French (intermediate)