

# Ekaterina Deyneka

\*\*\*\_\*\*\*\_\*\*\*\* | 92617, Irvine, CA, USA | [edeyneka@uci.edu](mailto:edeyneka@uci.edu) | [linkedin.com/in/edeyneka](https://www.linkedin.com/in/edeyneka)

## EDUCATION

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- University of California, Irvine (UCI)** Sep. 2020 – Jul. 2025  
Graduate Program in Computer Science Irvine, CA  
Average grade: 4.0/4.0
- Delft University of Technology (TU Delft)** Sep. 2018 – Jul. 2020  
Masters's Program in Nanobiology, Joint Degree with Erasmus Medical Center Delft, Netherlands  
Average grade: 8.2/10.0
- Moscow Institute of Physics and Technology (MIPT), State University** Sep. 2014 – Jul. 2018  
Bachelor's Program in Applied Mathematics and Physics Moscow, Russia  
Average grade: 4.8/5.0; top 1% of the department

## WORKING EXPERIENCE

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- Software Engineering Intern at Snap Inc.** Jun. 2021 – Sept. 2021  
Machine Learning Team Santa Monica, CA
- Denoising diffusion models application to solve a number of computer vision tasks: inpainting, adding noise and texture to synthetic data, adding noise and glare to assets

## RESEARCH EXPERIENCE

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- Graduate Student Researcher** June 2020 – Present  
Pierre Baldi's Group, Donald Bren School of Information and Computer Sciences, UCI Irvine, CA
- Neural networks to recover colors of the images taken in the infrared spectrum
  - Development of bioinformatics pipelines and tools for CircadiOmics
- Master Student Researcher** Sep. 2018 – Jul. 2020  
Pattern Recognition and Bioinformatics Lab, TU Delft Delft, Netherlands
- *De novo* genome assembly using long and short reads
  - Phylogenetic and variability analysis of proteins
  - Close interaction and collaboration with microbiologists, discussion of biological insights and presenting the results
  - Organizing a research agenda and managing fellows

## QUALIFICATIONS & SKILLS

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**Languages:** Python (numpy, sklearn, scipy, pandas, pytorch, tensorflow, keras, biopython, matplotlib, pyplot), Bash  
**General:** High performance computing cluster, AWS, L<sup>A</sup>T<sub>E</sub>X

## PUBLICATIONS

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- 2021 "Deep Learning to Enable Color Vision in the Dark, first author  
In progress
- 2020 "Comprehensive characterization of PE/PPE protein encoding genes in *M. tuberculosis* lineages through high-quality genome assemblies", first author  
In progress
- 2020 "Complete genome sequence of *Ralstonia pickettii* type strain DSM 6297 (ATCC 27511, NCTC 11149, K-288, NBRC 102503) and its conjugative plasmid determined by third-generation sequencing.", first author  
In progress

## SCHOLARSHIPS & AWARDS

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- 2020 The Donald Bren School of Information and Computer Sciences Dean's Award (10,000\$)
- 2019 The Next Generation Women Leaders award towards research, McKinsey & Company (2,000€)
- 2018–2020 Orange Tulip Scholarship for Master Program in Nanobiology (51,790€)