

Dr. Bora Uyar

Senior Bioinformatics Scientist
@ MDC-Berlin

Hannoversche Str. 28
Berlin 10115
Germany

✉ bora.uyar@mdc-berlin.de
🌐 <https://borauyar.github.io/>
👤 borauyar
👤 bora-uyar-11050425



Summary

I am a Senior Staff Scientist at the Bioinformatics & Omics Data Science platform of the Max-Delbrück-Center for Molecular Medicine. My current research is focused on the development of deep learning-based multi-modal data integration tools for precision oncology. My other roles in the platform include running research collaborations and supporting other researchers in the form of consultations, mentorships, workshops, and user-friendly software development for wet-lab researchers. My work in the last 15 years spanned various topics such as comparative genomics, protein sequence analysis in the context of molecular interactions and disease mechanisms, RNA bioinformatics, and omics data science with a focus on development of reproducible software for the bioinformatics community.

Experience

- 2015–Present **Bioinformatics Scientist**, *Max-Delbrück-Center for Molecular Medicine*, Berlin, Germany
- 2014–2015 **Postdoctoral Fellow**, *European Molecular Biology Laboratory*, Heidelberg, Germany

Education

- 2011–2014 **Ph.D. in Bioinformatics**, *European Molecular Biology Laboratory*, Heidelberg, Germany
- 2008–2010 **M.Sc. in Bioinformatics**, *Simon Fraser University, BC Cancer Agency*, Vancouver, Canada
- 2003–2008 **B.Sc. in Biological Sciences**, *Sabancı University*, Istanbul, Turkey

Profiles

- Google Scholar scholar.google.com/YEZr1LUAAAAJ
- GitHub github.com/borauyar
- ORCID orcid.org/0000-0002-3170-4890
- LinkedIn linkedin.com/in/bora-uyar-11050425

Selected Publications

Nature Communications, Flexynesis: A deep learning toolkit for bulk multi-omics data integration for precision oncology and beyond
2025

Genome Biology, Identifying tumor cells at the single-cell level using machine learning
2022

Cell Reports, Parallel genetics of regulatory sequences using scalable genome editing in vivo
2021

Aging Research Reviews, Single-cell analyses of aging, inflammation and senescence
2020

GigaScience, PiGx: reproducible genomics analysis pipelines with GNU Guix
2018

Cell, Mutations in Disordered Regions Can Cause Disease by Creating Dileucine Motifs
2018

Nucleic Acids Research, RCAS: an RNA centric annotation system for transcriptome-wide regions of interest
2017

My research in the news

- Using deep learning for precision cancer therapy (mdc-berlin.de)
- New textbook for computational genomics (mdc-berlin.de)
- Searching the sewers for viruses (bionity.com)
- AI identifies cancer cells (bionity.com)
- Parallel Genome Editing in Microscopic Worms Maps Regulatory Genomic Elements to Physiology (genengnews.com)
- Stray proteins cause genetic disorders (sciencedaily.com)
- A genetic chaperone for healthy aging? (bionity.com)