# Yasin Tepeli

→ +31 6 8248 3646 yitepeli@gmail.com yitepeli im yitepeli Ø yitepeli.github.io

My research interests lie in developing and studying AI methods to model complex biological systems, with a focus on oncology. My current work centers on therapeutic-target discovery and patient stratification for targeted cancer therapy, alongside diagnosing and mitigating bias in machine-learning models. Methodologically, I employ graph-based modeling and multi-omics integration across oncology applications, and address core ML challenges, selection bias and distribution shift, to deliver robust, translational results.

#### -EXPERIENCES -

#### **DELFT UNIVERSITY OF TECHNOLOGY**

Delft, NL

#### PhD / Postdoc Researcher, Delft Bioinformatics Lab

Sep 2020-present

- Developed computational tools for targeted cancer therapy: novel synthetic-lethal targets and patient stratification.
- Characterized selection bias in datasets/ML and proposed methods to mitigate it using semi-supervised learning.
- Supervised BSc, MSc, and PhD projects on DNA repair deficiency, survival modelling, bias mitigation, and mutational signatures.

#### **INSTITUTE OF HUMAN TECHNOPOLE**

Milan, IT

#### Visiting PhD Researcher, Computational Biology Research Center

Sep 2023-Jan 2024

- Built a framework to stratify oncogene-addicted cohorts by drug response.
- Prioritized synthetic-lethal gene pairs and co-designed validation experiments.

SABANCI UNIVERSITY Istanbul, TR

#### Research/Teaching Assistant, Computer Science & Engineering

Sep 2018–Sep 2020

- Identified cancer subgroups via functional graph kernels on multi-omics.
- · Predicted gene loss-of-function and essentiality using graph embeddings.
- TA for Artificial Intelligence, Algorithms, Data Science, and Machine Learning.

EPITA Paris, FR

**Erasmus Exchange, ME in Computer Engineering** 

Sep 2016-Feb 2017

## HAVELSAN Summer Intern, Big Data Department

Ankara, TR Jun 2017–Aug 2017

- Built an ELK-based log-analysis dashboard (Elasticsearch, Logstash, Kibana).
- Generated word-embedding vectors from English Wikipedia (web crawling, GloVe).

#### ERS YAZ. INT. TUR. SAN. TIC. A.S.

Antalya, TR

#### **Summer Intern, Web Development**

Jun 2016-Aug 2016

Built a web-based hotel management system (full stack: HTML, JavaScript, C#).

#### — EDUCATION —

#### PHD, DELFT BIOINFORMATICS LAB, DELFT UNIVERSITY OF TECHNOLOGY

Delft, NL

Computational Tools for Optimizing Targeted Cancer Treatments and Addressing Bias with J.P. de Gonçalves.

Sep 2020-Jun 2025

#### MSc (High Honors), Computer Science & Engineering, Sabanci University

Istanbul, TR

Discovering Cancer Patient Subgroups with Functional Graph Kernels with Ö. Taştan Okan.

Sep 2018-Sep 2020

#### BSc (High Honors), Computer Science, Bilkent University

Ankara, TR

AeroCast - Forecasting Flight Prices for Fixed Travel Days with Ç. Gündüz Demir.

Sep 2013-Sep 2018

#### -SKILLS —

Machine Learning scikit-learn, Keras, TensorFlow

Programming Python, R, MATLAB, C++, HTML, JavaScript, Bash, LaTeX

Software Linux, Git, Adobe Illustrator, Microsoft Office, Gurobi, MOSEK

Languages Turkish (native), English (fluent), Dutch (B1)

PUBLICATIONS & PREPRINTS —	
P8 <b>ELISL: early–late integrated synthetic lethality prediction in cancer Y. I. Tepeli</b> , C. Seale, J. P. Gonçalves. <i>Bioinformatics</i> , 40(1)	2024
P7 SNMF: Integrated Learning of Mutational Signatures and Prediction of DNA Repair Deficiencies S. Goossens, C. Seale, J. P. Gonçalves, Y. I. Tepeli. bioRxiv	2024
P6 Metric-DST: Mitigating Selection Bias Through Diversity-Guided Semi-Supervised Metric Learning Y. I. Tepeli, M. J. de Wolf, J. P. Gonçalves. <i>arXiv</i>	2024
P5 <b>DCAST: Diverse Class-Aware Self-Training Mitigates Selection Bias for Fairer Learning Y. I. Tepeli</b> , J. P. Gonçalves. <i>arXiv</i>	2024
P4 A Metagenomic Study of Antibiotic Resistance Across Diverse Soil Types and Geographical Locations S. Pillay, Y. I. Tepeli, P. van Lent, T. Abeel. <i>bioRxiv</i>	2024
P3 Overcoming selection bias in synthetic lethality prediction C. Seale, Y. I. Tepeli, J. P. Gonçalves. <i>Bioinformatics</i> , 38(18):4360–4367	2022
P2 <b>GEGE: Predicting Gene Essentiality with Graph Embeddings</b> H. I. Kuru, <b>Y. I. Tepeli</b> , Ö. Taştan. <i>Düzce University Journal of Science and Technology</i> , 10(1):88–97	2022
P1 <b>PAMOGK:</b> a pathway graph kernel-based multiomics approach for patient clustering <b>Y. I. Tepeli</b> , A. B. Ünal, F. M. Akdemir, Ö. Taştan. <i>Bioinformatics</i> , 36(21):5237–5246	2020
TALKS & CONFERENCES	
Leiden Computational Biology Center, Leiden, Netherlands. Invited talk (P5, P6).	2025
BYTE, Utrecht, Netherlands. Invited Talk.	2024
Human Technopole PhD Symposium, Milan, Italy. Talk (P8).	2024
BioSB, Egmond aan Zee, Netherlands. <i>Talk (P5)</i> .	2024
BioSB, Egmond aan Zee, Netherlands. <i>Poster (P6)</i> .	2023
ECCB, Barcelona, Spain. Poster (P8).	2022
BioSB, Lunteren, Netherlands. Talk (P8).	2022
BioDay, Delft, Netherlands. Poster (P8).	2022
OxfordML Summer School — Health Track, Oxford, UK. Attended.	2022
HIBIT, Istanbul, Turkey. Poster & pitch (P1).	2020
MLFPM Summer School, Basel, Switzerland. Attended.	2019
FENS Research Days, Sabanci University, Turkey. Poster (P2).	2019
AWARDS, HONORS, AND SCHOLARSHIPS	
Delft Bioinformatics Lab — Social Cohesion Award.	2024
TU Delft BioDay — Best Poster Award.	2022
HIBIT — 3rd Best Poster & Pitch.	2020
TUBITAK BİDEB — MSc 2210 National Scholarship.	2018
Bilkent University — BSc Comprehensive 100% Education Scholarship.	2013
OTHER UNPUBLISHED PROJECTS	
UNCOVERING AND EXPERIMENTALLY VALIDATING POTENTIAL GENE INTERACTIONS FOR CANCER • Prioritizing computationally derived potential synthetic lethal gene pairs and validating them in drug scree	2024 ens.
EARLY DETECTION OF SELF-HARM ON SOCIAL MEDIA <ul><li>Identified at-risk users from Reddit posts using classical ML and deep learning.</li></ul>	2019
SENTIMENT ANALYSIS OF TURKISH TWEETS  • Classified sentiment (positive/neutral/negative) using preprocessing word vectors. RNNs. and LSTMs.	2019

### DETECTING BRCA1 LOSS OF FUNCTION VIA GENE ALTERATIONS

2019

• Predicted BRCA1 loss of function using ML, Node2Vec, protein structures, and networks.

#### **ANIMAL VOICE RECOGNITION**

2017

• Extracted features with warbleR/seewave and trained SVM, Random Forest, and neural networks.

---INTERESTS ---

**Community engagement** Coordinated student-led retreats and volunteered initiatives (2023–present) **Hobbies** Cycling, running, Padel; photography