# Metin Balaban

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2017 – present Ph.D. Bioinformatics and Systems Biology UC SAN DIEGO (UCSD), SAN DIEGO, CA, US

Advisor: Siavash Mirarab. Expected graduation date: Dec. 2021.

2015 – 2017 M.Sc. Computer Science EPFL, SWITZERLAND

GPA: 5.54/6.00. Advisor: Patrick Thiran

2010 – 2015 B.S Computer Engineering (Valedictorian) MIDDLE EAST TECHNICAL UNIVERSITY, ANKARA, TURKEY

GPA: 3.97/4.00

SCHOLARSHIPS AND AWARDS

2020 UCSD Center for Microbiome Innovation Grand Challenge Award (\$67K grant)

2015-2016 EPFL Graduate Research Scholarship

2010-2015 Turkey State Outstanding Academic Success Scholarship

EXPERIENCE

Dec 2016 – Jun 2017 Research Intern Swisscom Digital Lab Lausanne, VD, Switzerland

Thesis: Real-time Traffic Estimation on Highways Using Cellular Data

Sep 2015 – Dec 2016 Research Scholar Laboratory of Computational Biology and Bioinformatics. EPFL, SWITZERLAND

Mentor: Bernard Moret

ORAL PRESENTATIONS

2020 ISMB Phylogenetic double placement of mixed samples Montreal (Virtual), Canada

2019 RECOMB APPLES: Fast Distance-Based Phylogenetic Placement. DISTRICT OF COLUMBIA, US

Won NSF travel fellowship award.

2018 GLBIO TreeCluster: Clustering biological sequences using phylogenetic trees. Madison, WI, US

Won ISCB travel fellowship award.

### **PUBLICATIONS**

### CONFERENCE & JOURNAL PUBLICATIONS AND PREPRINTS

Metin Balaban, Daniel Roush, Qiyun Zhu, and Siavash Mirarab (2021). "APPLES-2: Faster and More Accurate Distance-based Phylogenetic Placement using Divide and Conquer". In: bioRxiv. DOI: 10.1101/2021.02.14.431150.

Yueyu Jiang, <u>Metin Balaban</u>, Qiyun Zhu, and Siavash Mirarab (2021). "DEPP: Deep Learning Enables Extending Species Trees using Single Genes". In: *bioRxiv*. DOI: 10.1101/2021.01.22.427808.

Shahab Sarmashghi, <u>Metin Balaban</u>, Eleonora Rachtman, Behrouz Touri, Siavash Mirarab, and Vineet Bafna (2021). "Estimating repeat spectra and genome length from low-coverage genome skims with RESPECT". In: *bioRxiv*. DOI: 10.1101/2021.01.28.428636.

Eleonora Rachtman, Metin Balaban, Vineet Bafna, and Siavash Mirarab (2020). "The impact of contaminants on the accuracy of genome skimming and the effectiveness of exclusion read filters". In: Molecular Ecology Resources 20.3, pp. 649–661. DOI: https://doi.org/10.1111/1755-0998.13135.

Metin Balaban Curriculum Vitæ

Metin Balaban, Niema Moshiri, Uyen Mai, Xingfan Jia, and Siavash Mirarab (Aug. 2019). "TreeCluster: Clustering biological sequences using phylogenetic trees". In: *PLOS ONE* 14.8, pp. 1–20. DOI: 10.1371/journal.pone.0221068.

Metin Balaban, Shahab Sarmashghi, and Siavash Mirarab (Sept. 2019). "APPLES: Scalable Distance-Based Phylogenetic Placement with or without Alignments". In: Systematic Biology 69.3, pp. 566–578. ISSN: 1063-5157. DOI: 10.1093/sysbio/syz063.

Daniel Doerr, <u>Metin Balaban</u>, Pedro Feijão, and Cedric Chauve (2017). "The gene family-free median of three". In: *Algorithms for Molecular Biology* 12.1, p. 14.

### **TEACHING**

2020-2021 2012-2013 **Teaching Assistant** Intro to Computer Engineering **Teaching Assistant** C Programming

UC San Diego

MIDDLE EAST TECHNICAL UNIVERSITY

# SERVICE TO THE SCIENTIFIC COMMUNITY

2020 - present

# Peer reviewer for life-sciences journals

Briefings in Bioinformatics