Last Updated: April 16, 2021

# Education

2017– Joint Computational Biology Ph.D., Carnegie Mellon University-University of Pitts-

Present burgh, Pittsburgh, PA.

GPA: 3.84/4.0

2012-2017 B.Sc., Materials Science and Engineering, Sabanci University, Istanbul, Turkey.

Minor: Physics GPA: 3.6/4.0

# Research Experience

2018– Carvunis Lab, University of Pittsburgh, Pittsburgh, PA, Graduate Research Assistant.

Present Advisor: Dr. Anne-Ruxandra Carvunis

Projects:

- Developed simulation methods to analyze large yeast genetic interaction dataset using graph theory.
- Developed multiple sequence alignment analysis and clustering pipeline in R and Python to identify evolutionary changes in open reading frames.
- Used membrane molecular dynamics and structural prediction tools to understand the evolution of transmembrane domains in *de novo* genes.
- 2016 Computational Molecular Biology Lab, Freie Universität Berlin, Berlin, Germany, Undergraduate Researcher.

Advisor: Dr. Christoph Wehmeyer

Developed algorithms for automated umbrella sampling simulations on Markov Chain Monte Carlo systems as part of summer internship project.

2013–2017  $\ \, \textbf{MIDSTLAB}, \, Sabanci \, \, University, \, Istanbul, \, \textbf{Turkey}, \, \textbf{Undergradute Research Assistant}. \,$ 

Advisor: Dr. Canan Atilgan

Projects:

- Applied statistical physics and molecular dynamics to understand self-assembly of lipidoids to develop robust drug delivery systems.
- Used molecular dynamics, free energy perturbations, and steered molecular dynamics to study DHFR mutations causing antibiotic resistance in E.Coli.

## Awards

### Travel grants

- 2021 University of Pittsburgh Biomedical Graduate Student Association (BGSA) Travel grant Fellowships
- 2012–2017 Sabanci University B.Sc. Merit Scholarship covering full tuition upon success in nation-wide university entrance exam

### Honors

2014-2017 Placed in Dean's High Honor list (7 semesters)

# Work Experience

2015 Tusas Engine Industries (TEI), Eskisehir, Turkey, Intern.

Developed Microsoft Excel macros to automate analysis of mechanical properties of alloys

# Technical Experience

Extremely Proficient With

languages Python, R

technologies BioConductor, Tidyverse, BioPython, Scikit-Learn, Igraph, Networkx, LATEX, Bash Scripting, Git, Vim, MS Office Products

Have Experience With

languages PHP, C++

technologies MySQL, RShiny, NAMD, Amber, PyMol, VMD, Adobe Illustrator

## **Publications**

### Journal Articles

- [1] Nikolaos Vakirlis, **Omer Acar**, Brian Hsu, Nelson Castilho Coelho, S. Branden Van Oss, Aaron Wacholder, Kate Medetgul-Ernar, Ray W. Bowman, Cameron P. Hines, John Iannotta, Saurin Bipin Parikh, Aoife McLysaght, Carlos J. Camacho, Allyson F. O'Donnell, Trey Ideker, and Anne-Ruxandra Carvunis. De novo emergence of adaptive membrane proteins from thymine-rich genomic sequences. *Nature Communications*, 11(1):781, Feb. 2020.
- [2] Haleh Abdizadeh, Yusuf Talha Tamer, Omer Acar, Erdal Toprak, Ali Rana Atilgan, and Canan Atilgan. Increased substrate affinity in the Escherichia coli L28R dihydrofolate reductase mutant causes trimethoprim resistance. *Physical Chemistry Chemical Physics*, 19(18):11416-11428, 2017.

## Talks

Network Biology (Virtual), Cold Spring Harbor Labs, Cold Spring Harbor, NY, Detecting effector and sensor genes in genetic networks.

#### Posters

- [1] Omer Acar, Aaron Wacholder, and Anne-Ruxandra Carvunis. Exploring the evolution of young gene function through genetic interaction networks. In *CSHL Network Biology Conference*, Cold Spring Harbor, NY, March 2019.
- [2] Omer Acar, Aaron Wacholder, and Anne-Ruxandra Carvunis. Exploring de novo gene emergence through genetic interaction networks. In *Evolution in Philadelphia Conference*, *EPiC*, Philedelphia, PA, September 2019.
- [3] Omer Acar, Aaron Wacholder, and Anne-Ruxandra Carvunis. Exploring de novo gene

emergence through genetic interaction networks. In CMU-University of Pittsburgh Computational Biology Program Retreat, Pittsburgh, PA, August 2019.