

# Nil Sahin

---

Email: [nil\\_sahin@yahoo.com](mailto:nil_sahin@yahoo.com)

Website: <https://nsahin.github.io>

## Work Experience

- May 2023 – present      **Data Scientist at Recursion Pharmaceuticals**  
Implementing data solutions to industrialize drug discovery
- July 2021 – May 2023    **Computational Scientist at Cyclica, Inc.**  
Developed scientific software to streamline drug discovery

## Education

- 2015 – 2021              **University of Toronto**, Toronto, Ontario, Canada  
Ph.D., Department of Molecular Genetics, Faculty of Medicine  
Co-supervisors: Dr. Brenda J. Andrews, Dr. Quaid Morris
- 2011 – 2015              **Sabanci University**, Istanbul, Turkey  
BSc., Molecular Biology, Genetics and Bioengineering  
Faculty of Engineering and Natural Sciences
- 2006 – 2011              **Robert College**, Istanbul, Turkey

## Publications

- Yeung, C.H.L., **Sahin, N.**, and Andrews, B.J. (2022). Phenomics approaches to understand genetic networks and gene function in yeast. *Biochemical Society Transactions*. Accepted (Submission: BST-2021-0285CR1)
- Mattiazzi Ušaj, M.\*, **Sahin, N.\***, Friesen, H., Pons, C., Ušaj, M., Masinas, M., Shkurin, A., Morris, Q., Boone, C., and Andrews, B.J. (2020). Systematic genetics and single cell image analysis reveals widespread pleiotropy and cell-to-cell variability. *Molecular Systems Biology*, **16**(2), e9243.  
<https://doi.org/10.15252/msb.20199243> \*equal contribution
- Rubanova, Y., Rujan, S., Harrigan, C.F., Li, R., Wintersinger, J., **Sahin, N.**, Deshwar, A., Morris, Q. (2020). Reconstructing evolutionary trajectories of mutation signature activities in cancer using TrackSig. *Nature Communications*, **11**(1):731. doi: 10.1038/s41467-020-14352-7
- Sokolov, A., Ashenden, S., **Sahin, N.**, Lewis, R., Erdem, N., Ozaltan, E., Bender, A., Roth, F.P. and Cokol, M. (2019) Characterizing ABC-Transporter Substrate-Likeness Using a Clean-Slate Genetic Background. *Frontiers in Pharmacology*, **10**:448, doi: 10.3389/fphar.2019.00448
- Grys, B.T., Lo, D.S., **Sahin, N.**, Kraus, O.Z., Morris, Q., Boone, C., and Andrews, B.J. (2017). Machine learning and computer vision approaches for phenotypic profiling. *Journal of Cell Biology* **216**(1), 65–71. <https://doi.org/10.1083/jcb.201610026>
- Chandrasekaran, S., Cokol-Cakmak, M., **Sahin, N.**, Yilancioglu, K., Kazan, H. Collins, J.J. and Cokol, M. (2016). Chemogenomics and orthology-based design of antibiotic combination therapies. *Molecular Systems Biology* **12**(5):872, doi: 10.15252/msb.20156777

## **Research Experience**

- 2016 – 2021      **Andrews Laboratory, University of Toronto**  
Supervisor: Dr. Brenda J. Andrews  
**An Atlas of Morphological Phenotypes Associated with Gene Perturbations**  
Developed image analysis pipelines to identify genetic regulators of cellular morphology in the context of genome-wide single gene perturbations in yeast by applying machine learning and computer vision strategies
- 2015, Oct. – Dec.      **Morris Laboratory, University of Toronto**  
Supervisor: Dr. Quaid Morris  
**Mutational Signature Changes during Tumour Evolution**  
Implemented bioinformatics algorithms to quantify mutational signatures of 600 tumours from whole genome sequencing; identified significant signature differences between tumour types; created an opportunity for a new PhD thesis for a new student after this 5-week rotation project
- 2013 – 2015      **Cokol Laboratory, Sabanci University**  
Supervisor: Dr. Murat Cokol  
**Large-scale Experimental *E. coli* Drug Interactions Screen**  
Conducted drug interaction experiments among 25 antibiotics; analyzed results to find significantly synergistic drug pairs and relate them to mechanism of action of each antibiotic
- 2014, June – Aug.      **Roth Laboratory, University of Toronto**  
Supervisor: Dr. Frederick P. Roth  
**Drug Sensitivity Profiles of ABC Transporter deletion strains in *S. cerevisiae***  
Conducted screen to identify relationship between ABC transporters and drug sensitivity by using 16 *S. cerevisiae* strains and measured sensitivity of strains to 28 anti-fungal compounds with various mechanisms of action.

## **Conference Presentations**

Machine Learning and Computer Vision Approaches for Phenotypic Profiling in Yeast  
*Nil Sahin, Mojca Mattiazzi-Usaj, Quaid Morris, Charles Boone, Brenda J. Andrews*

- Dec. 2019      **Machine Learning in Computational Biology 2019**, Vancouver, BC, Canada  
Oct. 2019      **CytoData Symposium and Hackathon 2019**, Heidelberg, Germany  
Oct. 2018      **International Symposium on Health Informatics and Bioinformatics 2018**, Antalya, Turkey  
Aug. 2018      **Yeast Genetics Conference 2018**, Stanford, CA, USA  
Dec. 2017      **Medicine by Design Symposium 2017**, Toronto, ON, Canada  
Nov. 2017      **Genome Informatics 2017**, Cold Spring Harbor Laboratory, NY, USA  
Sept. 2017      **Society of Biomolecular Imaging and Informatics 2017**, San Diego, CA, USA  
May 2016      **Great Lakes Bioinformatics and the Canadian Computational Biology Conference**, Toronto, ON, Canada

Drug sensitivity profiles of ABC transporter deletion strains in *S. cerevisiae*  
*Nil Sahin, Frederick P. Roth, Murat Cokol*

- Sept. 2014      **EMBL Conference: Frontiers in Fungal Systems Biology**, Heidelberg, Germany

## **Awards and Scholarships**

- Jan. 2020      **2019-2020 Jennifer Dorrington Graduate Research Award**
- Sept. 2017      **2017-2018 University of Toronto Open Fellowship**
- Sept. 2017      **Poster Award (2<sup>nd</sup>)**  
Society of Biomolecular Imaging and Informatics, High Content 2017 Conference
- Sept. 2016      **2016-2017 Cecil Yip Doctoral Research Award**
- 2011 – 2015      **Sabanci University Full Honor Scholarship**  
Awarded with full tuition fee and monthly stipend for 4 years for being in the top thousand of more than a million students in the nationwide university entrance exam in 2011

## **Teaching Experience**

- Winter 2019      **University of Toronto – Teaching Assistantship in Computational Biology and Bioinformatics Graduate Course**  
Helped students at weekly office hours with their assignments and grading
- Fall 2017      **University of Toronto – Teaching Assistantship in Computer Science Course CSC120**  
Helped students at weekly lab hours, holding and marking course exams
- Fall 2014      **Sabanci University – Teaching Assistantship in Cell Biology Course BIO332**  
Evaluated student responses to innovative research papers from literature
- 2012 – 2015      **Sabanci University – Academic Support Program Education Coordinator & Executive Board Member**  
Held weekly meetings with 11 moderators on their teaching abilities
- 2011 – 2012      **Sabanci University – Academic Support Program Teaching Assistant**  
Tutored peers with Mathematics and Natural Sciences courses (Physics, Chemistry, and Biology) in weekly held study sessions after lecture hours for ten hours a week

## **Extracurricular Activities**

- 2018 – 2019      **University of Toronto – [Coders](#) Group Treasurer**  
Conducted accounting for the club  
Gave tutorials on machine learning and image processing in Python
- 2017 – 2018      **The Donnelly Centre Graduate Student Association Presidency**  
Established two seminar series on new technologies and trainee projects  
Organized the annual scientific conference for more than 200 members of the Donnelly Centre Research Institute

## **Languages**

- Programming      Current: Python (Advanced), Bash (Basic); Past: R , C++, MATLAB, Perl  
Spoken      English (Proficient), Turkish (Native)

**References:** Provided upon request, please send an email to [nil\\_sahin@yahoo.com](mailto:nil_sahin@yahoo.com)