# Melih Yilmaz

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# Research Interests

- Machine Learning: seq2seq learning, self-supervised learning, representation learning
- Computational Biology: proteomics, mass spectrometry, drug discovery, biomedical data science

## **EDUCATION**

#### University of Washington

Seattle, WA

Ph.D. in Computer Science

Sep 2020-Current

Co-advisors: William Noble, Sewoong Oh

#### **Koc University**

Istanbul, Turkey

B.S. in Electrical and Electronics Engineering GPA: 4.00/4.00, Ranked 1st in the class

#### Osaka University

Osaka, Japan

2015-2020

Center for Japanese Language and Culture, MEXT Scholar

Apr 2016–Jan 2017

### RESEARCH EXPERIENCE

# University of Washington

Seattle, WA

Ph.D. Student, (Supervisors: William Noble, Sewoong Oh)

Sep 2020-Current

- Focusing on proteomics, my current research builds deep learning methods to analyze mass spectrometry data.
- Formulated *de novo* peptide sequencing as a supervised seq2seq learning task and trained a transformer model advancing the state of the art [1]. Exploring self-supervised learning tasks to train foundation models for mass spectra on repository-scale data sets

Novo Nordisk Seattle, WA

Machine Learning Research Intern, (Supervisors: Per Greisen, Kristine Deibler)

Summer 2022

 Developed a novel approach combining protein language models and molecular graph neural networks to learn representations for modified peptide drug candidates with the Computational Drug Discovery group

#### Stanford University

Stanford, CA

Research Intern, (Supervisor: Tina Hernandez-Boussard)

Summer 2019

- Modeled post-chemotherapy patient reported outcomes and electronic health records (EHRs) for cancer patients.
- Performed trajectory clustering and risk group classification to identify vulnerable patient populations [2].

#### Koc University

Istanbul, Turkey

Undergraduate Research Assistant, (Supervisor: Murat Tekalp)

Fall 2018, Fall 2019

- Developed deep learning models for image and video compression. Leveraged next video frame prediction as a self-supervised learning task to train an end-to-end compression model outperforming conventional codecs [3].

## **Publications**

[1] M. Yilmaz, W. Fondrie, W. Bittremieux, S. Oh, and W. Noble, "De Novo Mass Spectrometry Peptide Sequencing with a Transformer Model", International Conference on Machine Learning, 2022, [code] [talk].

- A. Azad, M. Yilmaz, S. Bozkurt, J. Brooks, D. Blayney, and T. Hernandez-Boussard, "Diverse Patient Trajectories during Cytotoxic Chemotherapy: Capturing Longitudinal Patient Reported Outcomes", Cancer Medicine, 2021.
- G. Ozsoy \*, M. Yilmaz \*, O. Kirmemis, and M. Tekalp, "New results in end-to-end image and video compression by deep learning", in IEEE Signal Processing and Communications Applications Conference (SIU), 2020.

# SCHOLARSHIPS AND AWARDS

• Paul G. Allen School First-Year Ph.D. Fellowship	2020
<ul> <li>Monbukagakusho (MEXT) Scholarship in Science</li> <li>Awarded by Japanese Ministry of Education, Culture, Sports, Science and Technology</li> </ul>	2016
<ul> <li>Turkish Government High Honour Scholarship         <ul> <li>Awarded based on ranking (14th out of 1.8 million students) in National University Entrance Exam</li> </ul> </li> </ul>	2015
<ul> <li>Koc University Suna Kirac Scholarship</li> <li>Full tuition waiver and stipend during the B.Sc. based on National University Entrance Exam rank</li> </ul>	2015

# LEADERSHIP EXPERIENCE

Nucleate Seattle, WA Apr 2022 - Current

Managing Director, Seattle

- Founded Pacific Northwest chapter of Nucleate, a global trainee-led nonprofit supporting emerging entrepreneurs in biotech and helping them spin out of academic labs. (as featured in Geek Wire)
- Headed a team of 10+ PhD/MBA/MD students and postdocs running the Activator incubator program to identify and support venture teams with early-stage technologies from research institutions across PNW.

#### Professional Services

- Conference and Invited Talks
- Reviewer: Genome Research, RECOMB 2022
- UW CSE: PhD Admission Committee Member (2021)
- De Novo Peptide Sequencing Transformer (slides)
  - ISMB CompMS 2022
  - Novo Nordisk Research Center Seattle 2022

#### SKILLS

- Programming Languages and Tools:
  - Python, R, SQL, Julia, MATLAB, Java, C, C++
- Libraries:
  - PyTorch, Keras, NumPy, Pandas, Scikit-Learn
- LANGUAGES
- Turkish: Native
- Japanese: Advanced (JLPT N1)
- French: Intermediate
- Spanish: Elementary