# Melih Yilmaz

Email: melih@cs.washington.edu Website: melihyilmaz.github.io

## Research Interests

- Machine Learning: sequence learning, multimodal LLMs, self-supervised learning, representation learning
- Computational Biology: proteomics, protein design, regulatory genomics, biomedical data science

# **EDUCATION**

#### University of Washington

Seattle, WA

2015-2020

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

# Experience

#### University of Washington

Seattle, WA

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

Sep 2020-Current

- My current research builds deep learning methods to analyze mass spectrometry data in proteomics
- Formulated *de novo* peptide sequencing as a supervised seq2seq learning task and trained a transformer model advancing the state of the art [1-4]. Exploring self-supervised learning tasks to train foundation models for mass spectra on repository-scale data sets

Amazon Seattle, WA

Applied Scientist Intern, (Supervisors: Bella Dubrov, Tommi Jaakkola)

Summer 2024

 Developed novel multimodal protein language models for a confidential project within the Grand Challenge organization

#### Alphabet (Calico Life Sciences)

South San Francisco, CA

Machine Learning Research Intern, (Supervisor: David Kelley)

Summer 2023

 Explored model ensembling and knowledge distillation strategies to improve gene expression prediction with state-of-the-art DNA sequence models

Nucleate Seattle, WA

Managing Director, Seattle

Apr 2022 - Sep 2023

- 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 GeekWire)
- Recruited and led a team of 12 PhD/MBA/MD students and postdocs running the Activator incubator program to identify and support venture teams with early-stage technologies

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, CA Summer 2019

Research Intern, (Supervisor: Tina Hernandez-Boussard)

- 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 [5]

## **PUBLICATIONS**

- [1] M. Yilmaz, W. Fondrie, W. Bittremieux, C. Melendez, R. Nelson, V. Ananth, S. Oh, and W. Noble, "Sequence-to-sequence translation from mass spectra to peptides with a transformer model", *Nature Communications*, 2024, [code].
- [2] C. Melendez, J. Sanders, M. Yilmaz, W. Bittremieux, W. Fondrie, S. Oh, and W. Noble, "Accounting for digestion enzyme bias in Casanovo", Journal of Proteome Research, 2024.
- [3] V. Ananth, J. Sanders, M. Yilmaz, B. Wen, S. Oh, and W. Noble, "A learned score function improves the power of mass spectrometry database search", *Bioinformatics*, 2024.
- [4] 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 (ICML)*, 2022, [code][talk].
- [5] 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.

#### Conference and Invited Talks

- Translating from mass spectra to peptides with a transformer model (slides)
  - ASMS 2023 (Oral Presentation and Evening Workshop)
- De Novo Peptide Sequencing Transformer (slides)
  - ISMB CompMS 2022
  - Novo Nordisk Research Center Seattle 2022

#### SCHOLARSHIPS AND AWARDS

• ]	Paul G. Allen School First-Year Ph.D. Fellowship	2020
• ]	Monbukagakusho (MEXT) Scholarship in Science  – Awarded by Japanese Ministry of Education, Culture, Sports, Science and Technology	2016
• '	Turkish Government High Honour Scholarship  - Awarded based on ranking (14th out of 1.8 million students) in National University Entrance Exam	2015
• ]	Koc University Suna Kirac Scholarship	2015

# SERVICE

• Program Committee: CompMS (ISMB 2024) • Reviewer: Genome Research, PLOS Computational Biology, Journal of the American Society for Mass Spectrometry, ISMB (2024), RECOMB (2022, 2024) • UW CSE: PhD Admission Committee Member (2021)

- Full tuition waiver and stipend during the B.Sc. based on National University Entrance Exam rank

# SKILLS

- Programming Languages and Tools:
  - Python, R, SQL, Julia, MATLAB, Java, C, C++
- Libraries:
  - PyTorch, Hugging Face, Keras, NumPy, Pandas, Scikit-Learn

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## LANGUAGES

• Turkish: Native

• Japanese: Advanced (JLPT N1)

• French: Intermediate

• Spanish: Elementary