Samuel Sledzieski

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Cambridge, MA

Massachusetts Institute of Technology

EDUCATION

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	PhD, Electrical Engineering and Computer Science • In Progress	2019 – 2024	
	 Concentration: Protein language models, protein and drug interactions, protein structure Advisor: Dr. Bonnie Berger 	e	
	SM, Electrical Engineering and Computer Science	2019 – 2021	
	University of Connecticut	Storrs, CT	
	BS, Computer Science • Minor in Molecular and Cellular Biology • Concentration: Bioinformatics, Data Science • Advisor: Dr. Mukul Bansal • Magna Cum Laude, Honors Scholar	2015 – 2019	
RESEARCH	Massachusetts Institute of Technology	Cambridge, MA	
	Research Assistant, Computation and Biology Group	Feb 2020 – Present	
	Microsoft Research	Redmond, WA	
	Research Intern, AI For Good Lab	May 2023 – Aug 2023	
	Cellarity	Cambridge, MA	
	Machine Learning Intern, Perturbation Biology Group	May 2021 – Aug 2021	
	MIT Lincoln Laboratory	Lexington, MA	
	Summer Research Program, Advanced Lasercom Systems Group	May 2019 – Aug 2019	
	University of Connecticut	Storrs, CT	
	Undergraduate Research Assistant, Computational Biology Lab	Jan 2017 – May 2019	
	Software Developer, Jackson Laboratory for Genomic Medicine	Aug 2018 – May 2019	
	Undergraduate Research Assistant, Nelson Lab	Oct 2015 – Dec 2016	
TEACHING	Massachusetts Institute of Technology	Cambridge, MA	
	Teaching Assistant, Machine Learning in Genomics (6.878)	Fall 2021	
	Teaching Assistant, Intro to Deep Learning (6.S191)	Winter 2021, 2022, 2023	
	University of Connecticut	Storrs, CT	
	Teaching Assistant, Theory of Computation	Spring 2018	
JOURNAL PUBLICATIONS	[5] Singh*, Sledzieski *, Bryson, Cowen, Berger, "Contrastive learning in protein language space predicts interactions between drugs and protein targets", Proceedings of the National Academy of Sciences 120.24 (2023): e2220778120.		
	[4] Kumar, Brenner, Sledzieski , Olaosebikan, Lynn-Goin, Putnam, Yang, Lewinski, Singh, Daniels, Cowen, Klein-Seetharaman, "Transfer of knowledge from model organisms to evolutionarily distant non-model organisms: The coral Pocillopora damicornis membrane signaling receptome," Plos one 18.2 (2023). 10.1371/journal.pone.0270965		

20. doi: 10.1089/cmb.2021.0507. Epub ahead of print. PMID: 36125448.

[3] Zaman*, **Sledzieski***, Wu, Bansal, "virDTL: Viral recombination analysis through phylogenetic reconciliation and its application to sarbecoviruses and SARS-CoV-2," J Comput Biol. 2022 Sep

[2] Singh*, Devkota*, **Sledzieski**, Berger, Cowen, "Topsy-Turvy: integrating a global view into sequence-based PPI prediction," Bioinformatics, 38.Supplement 1 (July 2022): i264–i272.

	sequence-based, structure-aware, genome-scale predictions of pro <i>Cell Systems</i> 12.10 (2021): 969-982.		
CONFERENCE AND WORKSHOPS	Sledzieski* , Singh*, Cowen, Berger, "Contrasting drugs from decoys" NeurIPS Workshop on Machine Learning for Structural Biology (MLSB) 2022.		
	[3] Sledzieski* , Singh*, Cowen, Berger, "Adapting protein language models for rapid DTI prediction" NeurIPS Workshop on Machine Learning for Structural Biology (MLSB) 2021.		
	 [2] Sledzieski*, Singh*, Cowen, Berger, "Sequence-based prediction of p a structure-aware interpretable deep learning model," <i>Conference on F Molecular Biology</i> (RECOMB) 2021. [1] Sledzieski, Zhang, Mandoiu, Bansal, "TreeFix-TP: Phylogenetic Erro Reconstruction of Viral Transmission Networks," <i>Pacific Symposium on</i> 	Research in Computational or Correction for Accurate	
	Proceedings, pages 119-130.		
PREPRINTS	[1] Kousi, Boix, Park, Mathys, Sledzieski , Peng, Bennett, Tsai, Kellis, "Sing reveals cell-type-specific somatic mutational burden in Alzheimer's Den April 2022, 10.1101/2022.04.21.489103		
	Intelligent Systems for Molecular Biology (ISMB)	Jul 2022, Jul 2023	
	Cold Spring Harbor Laboratory Meeting on Network Biology	Mar 2021, Mar 2023	
	Machine Learning in Structural Biology (MLSB) Workshop at NeurIPS	Dec 2021, Dec 2022	
	Research on Computational Molecular Biology (RECOMB)	Apr 2019, May 2022	
	Pacific Symposium on Biocomputing (PSB)	Jan 2021	
	IEEE ICCABS Workshop on Computational Advances for Next Generation Sequencing Oct 2018		
	UConn Fall Frontiers in Undergraduate Research	Oct 2018	
	University of Connecticut Bioinformatics Seminar	Mar 2018, Oct 2018	
AWARDS &	National Science Foundation (NSF) Graduate Research Fellowship	2021 - 2024	
FELLOWSHIPS	First Place, MIT Intro to Deep Learning Final Project Competition	2020	
	New England Scholar, University of Connecticut	2017 – 2019	
	Dean's List, College of Liberal Arts and Sciences, School of Engineering	2015 – 2019	
	Academic Excellence Scholarship, University of Connecticut	2015 – 2019	
	National Merit Scholarship Finalist	2014	
MEMBERSHIPS & ACTIVITIES	International Society for Computational Biology (ISCB)		
& ACTIVITIES	Institute of Electronics Engineers (IEEE)		
& ACTIVITIES	Association for Computing Machinery (ACM)		
& ACTIVITIES			

[1] Sledzieski*, Singh*, Cowen, Berger, "D-SCRIPT translates genome to phenome with

[CV compiled on 2023-10-04]

Upsilon Pi Epsilon, Computer Science Honor Society (UPE)