Samuel Sledzieski

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RESEARCH	**Flatiron Institute**	**New York, NY**
	Flatiron Research Fellow, Center for Computational Biology	Sep 2024 – Present
	Princeton University	**Princeton, NJ**
	Visiting Research Collaborator, Lewis-Sigler Institute for Integrative Genomics	Sep 2024 – Present
	Massachusetts Institute of Technology	**Cambridge, MA**
	Research Assistant, Computation and Biology Group	Feb 2020 – Aug 2024
	Centre Scientifique de Monaco	**Monaco**
	Visiting Researcher	Jan 2024 – Feb 2024
	Serinus Biosciences	**Cambridge, MA**
	Consultant	Feb 2023 – Dec 2023
	Microsoft Research	**Redmond, WA**
	Research Intern, AI For Good Lab	May 2023 – Oct 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

EDUCATION	**Massachusetts Institute of Technology**	**Cambridge, MA**
	PhD, Computer Science	2019 – 2024

- Minor in Biological Engineering
- Concentration: Protein language models, protein and drug interactions, protein structure
- Advisor: Dr. Bonnie Berger

	SM, Electrical Engineering and Computer Science	2019 – 2021
	University of Connecticut	**Storrs, CT**
	BS, Computer Science	2015 – 2019

- Minor in Molecular and Cellular Biology
- Concentration: Bioinformatics, Data Science
- Advisor: Dr. Mukul Bansal
- Magna Cum Laude, Honors Scholar

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	[9]	Kshirsagar, Meller, Humphreys, **Sledzieski**, Xu, Dodhia, Horvitz, Berger, Bowman, Lavista Ferres, Baker, Baek, "Rapid and accurate prediction of protein homo-oligomer symmetry," *Nature Communications*, In Press (2025)
	[8]	Singh, Im, Qiu, Macnkess, Gupta, Sorenson, **Sledzieski**, Erlach, Wendt, Nanfack, Bryson, Berger, "Learning the Language of Antibody Hypervariability," *Proceedings of the National Academy of Sciences*, 122.1 (2025): e2418918121
	[7]	**Sledzieski**, Kshirsagar, Baek, Berger, Dodhia, Lavista Ferres, "Democratizing Protein Language Models with Parameter-Efficient Fine-Tuning," *Proceedings of the National Academy of Sciences*, 121.26 (2024): e2405840121
	[6]	**Sledzieski***, Devkota*, Singh, Cowen, Berger, "TT3D: Leveraging Pre-Computed Protein Sequence Models to Predict Protein-Protein Interactions," *Bioinformatics*, 2023: btad663
	[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
	[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 20. doi: 10.1089/cmb.2021.0507. Epub ahead of print. PMID: 36125448
	[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
	[1]	**Sledzieski***, Singh*, Cowen, Berger, "D-SCRIPT translates genome to phenome with sequence-based, structure-aware, genome-scale predictions of protein-protein interactions," *Cell Systems*, 12.10 (2021): 969-982

CONFERENCE AND WORKSHOPS	[6]	**Sledzieski**, Versavel, Singh, Ocitti, Devkota, Kumar, Shhpilker, Roger, Yang, Lewinski, Putnam, Berger, Klein-Seetharaman, Cowen, "Decoding the Functional Interactome of Non-Model Organisms with PHILHARMONIC," *Conference on Research in Computational Molecular Biology (RECOMB)*, 2025
	[5]	**Sledzieski**, Kshirsagar, Baek, Berger, Dodhia, Lavista Ferres, "Parameter-Efficient Fine-Tuning of Protein Language Models Improves Prediction of Protein-Protein Interactions," *Machine Learning for Structural Biology Workshop at NeurIPS*, 2023
	[4]	**Sledzieski***, Singh*, Cowen, Berger, "Contrasting drugs from decoys," *Machine Learning for Structural Biology Workshop at NeurIPS*, 2022
	[3]	**Sledzieski***, Singh*, Cowen, Berger, "Adapting Protein Language Models for Rapid Drug-Target Interaction Prediction," *Machine Learning for Structural Biology Workshop at NeurIPS*, 2021
	[2]	**Sledzieski***, Singh*, Cowen, Berger, "Sequence-based prediction of protein-protein interactions: a structure-aware interpretable deep learning model," *Conference on Research in Computational Molecular Biology (RECOMB)*, 2021
	[1]	**Sledzieski***, Zhang, Mandoiu, Bansal, "TreeFix-TP: Phylogenetic Error Correction for Accurate Reconstruction of Viral Transmission Networks," *Pacific Symposium on Biocomputing (PSB)*, 2021

PREPRINTS	[2]	Vizgaudis, Kumar, Olaosebikan, Roger, Brenner, **Sledzieski**, Yang, Lewinski, Singh, Daniels, Cowen, Klein-Seetharaman, "Insulin Signaling and Pharmacology in Corals," *Authorea Preprints*, posted 31 January 2024, 10.22541/au.170666200.07483513/v1
	[1]	Kousi, Boix, Park, Mathys, **Sledzieski**, Peng, Bennett, Tsai, Kellis, "Single-cell mosaicism analysis reveals cell-type-specific somatic mutational burden in Alzheimer's Dementia," *bioRxiv*, posted 22 April 2022, 10.1101/2022.04.21.489103

PRESENTATIONS	Intelligent Systems for Molecular Biology (ISMB)	Jul 2022, Jul 2023, Jul 2024

	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, Dec 2023
	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 & FELLOWSHIPS	Flatiron Postdoctoral Research Fellowship	2024 - 2026
	National Science Foundation (NSF) Graduate Research Fellowship	2021 - 2024
	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

SOFTWARE	**D-SCRIPT**	https://github.com/samsledje/D-SCRIPT
	36k+ PyPI downloads	
	ConPLex	https://github.com/samsledje/ConPLex
	8k+ PyPI downloads	
	PHILHARMONIC	https://github.com/samsledje/philharmonic
	virDTL	https://github.com/suz11001/virDTL
	TreeFix-TP	https://github.com/samsledje/TreeFix-TP

MENTORING	MIT Undergraduate Research Opportunities Program (UROP) Advisor	2021 – 2024
	MIT Research Summer Institute (RSI) Advisor	2022 – 2023
	HackMIT Mentor	2019 – 2022

PEER REVIEW	*Nature, Nature Methods, Nature Communications, Current Opinions in Structural Biology, Bioinformatics, Journal of Computational Biology, PLOS Computational Biology, IEEE Transactions on Artificial Intelligence, NeurIPS, Machine Learning in Structural Biology (MLSB), RECOMB, ISMB*

MEMBERSHIPS & ACTIVITIES	International Society for Computational Biology (ISCB)
	Institute of Electronics Engineers (IEEE)
	Association for Computing Machinery (ACM)
	Tau Beta Pi, Engineering Honor Society (TBΠ)
	Eta Kappa Nu (IEEE-HKN)
	Upsilon Pi Epsilon, Computer Science Honor Society (UPE)

[*CV compiled on 2025-02-27*]