

PINAR DEMETCI

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EDUCATION

Ph.D. Computational Biology

Brown University 2018 - 2023 (Expected)
GPA: 3.90/4.00

M.Sc. Computer Science

Brown University 2018 - 2020 (Expected)
GPA: 4.00/4.00

B.Sc. Bioengineering

Olin College of Engineering 2013 - 2017
GPA: 3.67/4.00

AWARDS

2020	ICML WBC Best Poster Award Awarded for research project
2020	ICML WBC Fellowship Conference fellowship awarded to cover attendance fees
2016	MCM/ICM Meritorious Winner Interdisciplinary Contest in Mathematical Modeling (top 10%)
2015-2017	Olin Alumni Merit Scholarship Award covered miscellaneous living costs on campus
2013-2017	Sunlin Chou International Scholarship Award covered 50% of tuition
2013-2017	Olin Merit Scholarship Award covered 50% of tuition
2013	Honorable Mention First Step to Nobel Prize in Physics Research Project Contest
2013	First Place MEF Research Project Contest

TEACHING EXPERIENCE

Grad. Teaching Assistant	Spring 2019
CSCI2820 Advanced Algorithms in Comp. Bio. & Medical Bioinfo. at Brown University	
Underg. Teaching Assistant	Fall 2016
AHSE1515 Products & Markets at Olin College	
SCI1240 Designing Better Drugs at Olin College	

COMMUNITY SERVICE & MEMBERSHIPS

2020 -	Ph.D. Admissions Committee Computational Biology Program at Brown University
2020	Reviewer for MLCB Conference Machine Learning in Computational Biology Conference
2020 -	Peer Mentor for Incoming International Graduate Student at Brown University
2020 -	Society for Industrial and Applied Mathematics (SIAM) Student member
2018 -	International Society for Computational Biology (ISCB) Student Member
2018 -	Models, Inference & Algorithms (MIA) at Broad Institute of MIT and Harvard Member
2018 -	Graduate Women in Science and Engineering (GWISE) at Brown U. Student Member

SELECTED EXPERIENCE

Brown University

Graduate Research & Teaching Assistant

Microsoft Research

Research Intern (Genomics)

Massachusetts Institute of Technology

Research Associate

Olin College of Engineering

Undergraduate Research & Teaching Assistant

Sep 2018 - Present

Providence, RI

June 2020 - Sep 2020

Redmond, WA

May 2017 - Aug 2018

Cambridge, MA

Sep 2015 - May 2017

Needham, MA

PUBLICATIONS & PRE-PRINTS

* Denotes equal contribution

6. **P Demetci***, R Santorella*, B Sandstede, W Stafford Noble, R Singh. Gromov-Wasserstein optimal transport to align single-cell multi-omics data (2020). *bioRxiv*.
5. R Singh, **P Demetci**, G Bonora, V Ramani, C Lee, H Fang, Z Duan, X Deng, J Shendure, C Distche, W Stafford Noble. Unsupervised manifold alignment for single-cell multi-omics data (2020). *IEEE/ACM Transactions on Computational Biology and Bioinformatics*.(in press)
4. B Alpay*, **P Demetci***, S Istrail, D Aguiar. Combinatorial and statistical prediction of gene expression from haplotype sequence (2020). *Bioinformatics*.36:Supplement-1: i194-i202.
3. **P Demetci**, W Cheng, G Darnell, X Zhou, S Ramachandran, L Crawford. Multi-scale genomic inference using biologically annotated neural networks (2020). *bioRxiv*. (currently under review at *Nature Communications*).
2. D Parker*, **P Demetci***, G W Li. Rapid accumulation of motility-activating mutations in resting liquid culture of *Escherichia coli* (2019). *Journal of Bacteriology*. 201(19):e00259-19
1. **P Demetci**, C Nichols, Y V Zastavker, J D Stolk, A Dillon, M Gross. Externalization and internalization in the classroom: How do they emerge and why is it important? (2016). *IEEE Frontiers in Education Conference*

SELECTED CONFERENCES & INVITED TALKS

* Denotes equal contribution, Presenters underlined

2020	Machine Learning in Computational Biology: Oral Presentation (Acceptance rate: 15%) Gromov-Wasserstein optimal transport to align single-cell multi-omics data <u>P Demetci*</u> , R Santorella*, B Sandstede, W S Noble, R Singh
2020	Workshop on Optimal Control, Optimal Transport University of Minnesota - Institute for Mathematics and Its Applications Gromov-Wasserstein optimal transport to align single-cell multi-omics data <u>P Demetci*</u> , R Santorella*, B Sandstede, W S Noble, R Singh
2020	ICML Workshop on Computational Biology: Spotlight Talk & Poster (Acceptance rate: 21%) Gromov-Wasserstein optimal transport to align single-cell multi-omics data <u>P Demetci*</u> (spotlight), <u>R Santorella*</u> (poster), B Sandstede, W S Noble, R Singh
2020	ISMB (ML in CSB Track): Spotlight Talk & Poster (Acceptance rate: 25%) Gromov-Wasserstein optimal transport to align single-cell multi-omics data <u>P Demetci*</u> (poster), <u>R Santorella*</u> (spotlight), B Sandstede, W S Noble, R Singh
2020	ISMB Proceedings: Oral Presentation (Acceptance rate: 19%) Combinatorial and statistical prediction of gene expression from haplotypes <u>B Alpay*</u> , <u>P Demetci*</u> , S Istrail, D Aguiar
2020	ACM-BCB Proceedings: Oral Presentation (Acceptance rate: 27%) Unsupervised manifold alignment for single-cell multi-omics data <u>R Singh</u> , <u>P Demetci</u> , G Bonora, V Ramani, C Lee, H Fang, ..., W Stafford Noble.
2020	Computational Intelligence & Applications: Invited Talk (at Brown U.) Gromov-Wasserstein optimal transport to align single-cell multi-omics data <u>P Demetci*</u> , R Santorella*, B Sandstede, W S Noble, R Singh
2019	CCV-Con: Invited Talk (at Brown U.) Biologically Annotated Neural Networks for Multi-Scale Genomic Discovery <u>P Demetci</u> , W Cheng, S Ramachandran, L Crawford
2016	Frontiers in Education (FIE): Oral Presentation (Acceptance rate: 48%) Internationalization and Externalization in the Classroom: <u>P Demetci</u> , <u>C Nichols</u> , YV Zastavker, JD Stolk, A Dillon, M Gross