RATUL CHOWDHURY, Ph. D

Assistant Professor

Department of Chemical and Biological Engineering Iowa State University. 5009 ATRB, 5672 Bissel Drive Ames, IA. 50011 Email: ratul@iastate.edu Tel: (814) 777 5355 ORCID 0000-0003-4522-6911 Google Scholar Profile

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♦ EDUCATION

The Pennsylvania State University (Costas Maranas Lab)

Ph.D. in Chemical Engineering + Minor in Computational Sciences

Thesis title: Computational redesign of channel proteins, enzymes, and antibodies

Jadavpur University, Kolkata, India

Bachelor of Science in Chemical Engineering

May/13

♦ RESEARCH INTERESTS

- Deep Learning on protein sequence, structure, and functional landscapes
- Activity and specificity engineering of proteins
- Protein interaction with proteins and non-proteins
- Protein-based biomaterials
- Drug-discovery in context of human proteome
- Structure-guided metabolic modeling, pathway design, and strain engineering

◆ PROFESSIONAL EXPERIENCE

Assistant Professor, Department of Chemical and Biological Engineering Jul/22-present Iowa State University, Ames, IA. Postdoctoral Fellow, Laboratory of Systems Pharmacology Jan/20-Jun/22 Harvard Medical School, Boston, MA. Advisors: Prof. Peter Sorger + Prof. Mohammed AlQuraishi Graduate Research Assistant, Department of Chemical Engineering Jul/13-Dec/19 The Pennsylvania State University, University Park, PA. Advisor: Prof. Costas D. Maranas Graduate Teaching Assistant, Department of Chemical Engineering Jan/17-May/17 The Pennsylvania State University, University Park, PA. Research Intern, Mechanobiology Institute Apr/12-Jul/12 National University of Singapore, 5A Engineering Drive 1, Singapore. REU Mentor: Prof. P. Kanchanawong Research Intern, Molecular Reproduction, Development and Genetics Apr/11-Jul/11 Indian Institute of Science, Biological Science Bldg., Karnataka 560012, India

♦ HONORS AND AWARDS

REU Mentor: Prof. R. Dighe

•	Protein structure prediction work featured in <i>Science</i> on Derek Lowe's blog (science.org)	/2021
•	Protein structure prediction work featured in Forbes alongside AlphaFold2 (forbes.ai)	/2021
•	Selected Speaker at SynBYSS Synthetic Biology Young Speaker Series 2022 (EBRC - https://ebrc.org/)	/2021
•	Invited Guest Editor for Special Issue in Frontiers in Biotech. & Bioengineering (Issue)	/2021
•	Alumni Spotlight Penn State University 2020 / Distinguished Alumni Series (Article link)	/2020
•	Cover article for Nature Materials, Volume 19, Issue 3	/2020
•	AIChE Separations Divisions, Dibakar Bhattacharyya Outstanding Graduate Researcher Award	/2019
•	Enzyme Engineering Conference XXV Conference Travel Award	/2019

•	Biochemical & Molecular Engineering XXI Conference Graduate Award	/2019
•	NAMS Outstanding Young Membrane Researcher Award (North American Membrane Society)	/2019
•	Graduate Research Travel Award for NAMS 2019 conference – Pittsburgh	/2019
•	Brown Graduate Award (Best Graduate Symposium Poster Award – Penn State)	/2018
•	Ward Graduate Award (Best Graduate Student Paper of the Year Award for PoreDesigner -Nature Comm)	/2018
•	Best Poster Award (Gordon Research Conference, Membranes, New Hampshire)	/2018
•	Best Oral Presentation Award (Gordon Research Seminar, Membranes, New Hampshire)	/2018
•	MDPI Feature Paper – OptMAVEn-2.0 published in MDPI journal Antibodies, Vol. 7	/2018
•	Best Oral Presentation in Rational Protein Engineering (AIChE Annual Meeting, Minnesota)	/2017
•	Summer Research Fellowship, Mechanobiology Lab, National University of Singapore	/2012
•	Distinguished Undergraduate Attendee at Thirteenth Marcel Grossmann Meeting, Stockholm	/2012
•	Summer Research Fellowship, Department of Science and Technology , Govt. of India	/2011
•	Best Presenter Award and Summer Cosmology Workshop Invitation (IUCAA, Pune, India)	/2010
•	Kishore Vaigyanik Protsahan Yojana (KVPY) – highest undergraduate research award from Govt. of India	/2010
◆ PUBLIC	ATIONS	
, 102210	Published/ Accepted/ Under review (* Equal contribution)	
•	Single-sequence protein structure prediction using language models from deep learning.	
	Biorxiv. (Under Review: Nature Biotechnology)	
	R Chowdhury*,, G Church, PK Sorger, M AlQuraishi. [Protein Structure Deep Learning]	/2022
•	Antiviral efficacy and mechanisms of plant-derived polyphenols for inactivation of Tulane virus,	
	a surrogate for human norovirus. (Accepted: ASM)	
	C Oh*, <u>R Chowdhury</u> *,, M Kumar. [Green chemistry Water Purification]	/2022
	Computational prediction of the effect of amino acid changes on the binding affinity	
_	between SARS-CoV-2 spike and human ACE2 receptor. PNAS	
	C Chen, VS Boorla, R Chowdhury,, CD Maranas. [Protein Structure SARS CoV-2]	/2021
•	Foldamer-based ultrapermeable and highly selective artificial aquaporins that exclude protons.	
	Nature Nanotechnology A Roy, J Shen, H Joshi, R Chowdhury,, M Kumar. [Nature-inspired Biophysics]	/2021
	A Roy, J Shen, 11 Joshi, K Chowdhury,, M Kumar. [Nature-inspired Biophysics]	/2021
•	De novo design of high-affinity antibody variable regions (scFv) against the SARS-CoV-2	
	spike protein. Biorxiv. (Under Review: Journal of Chemical Information and Modeling)	
	VS Boorla, <u>R Chowdhury</u> *,, JJ Gray, CD Maranas. [Protein Structure SARS CoV-2]	/2021
•	Recombination and lineage-specific mutations led to the emergence of SARS-CoV-2.	
	Genome Medicine JAP Galindo, I Filip, R Chowdhury,, R Rabadan. [Protein Structure SARS CoV-2]	/2021
	JAI Gainido, I l'inp, <u>R' Chowdhury</u> ,, R' Rabadan. [Flotelli Structure SARS Cov-2]	/2021
•	Computational biophysical characterization of the SARS-CoV-2 spike protein binding with the ACE2	
	receptor and implications for infectivity. Computational Structural Biotechnology Journal	
	R Chowdhury*, VS Boorla, and CD Maranas. [Protein Structure SARS CoV-2]	/2020
	Rapid fabrication of precise high-throughput filters from membrane protein nanosheets.	
-	Nature Materials * featured article	
	YM Tu, W Song,, R Chowdhury,, M Kumar. [Nature-inspired Biophysics]	/2020
•	IPRO+/- Computational protein design tool allowing for insertions and deletions.	
	Structure – Cell Press R Chowdhury,, CD Maranas. [Enzyme Engineering Protein Function]	/2020
	it chowenery,, ob interested, [Enzyme Engineering From Function]	, 2020

•	acid lactone and orsellinic acid. Protein Engineering Design and Selection	
	Z Wang, A Doshi, <u>R Chowdhury</u> ,, PC Cirino. [Biosensor Protein Function]	/2020
	2	, = 0 = 0
•	From directed evolution to computational enzyme engineering—A review.	
	AICHE Journal	
	R Chowdhury and CD Maranas. [Enzyme Engineering]	/2019
	Artificial water channels enable fast and selective water permeation through	
	water-wire networks. Nature Nanotechnology	
	W Song, H Joshi, R Chowdhury,, M Kumar. [Nature-inspired Biophysics]	/2019
	Directed evolution reveals the functional sequence space of an adenylation domain	
	specificity code. ACS Chemical Biology	
	K Throckmorton, V Vinnik, R Chowdhury,, BF Pfleger. [Enzyme Engineering Protein Function]	/2019
	7-log virus removal in a simple functionalized sandfilter.	
_	Environmental Science & Technology	
	L Samineni, B Xiong, R Chowdhury,, M Kumar. [Green Chemistry Water Purification]	/2019
	PoreDesigner for tuning solute selectivity in a robust and highly permeable outer membrane pore.	
	Nature Communications	
	R Chowdhury, T Ren,, CD Maranas. [Channel Protein Separations]	/2018
_	O-MAYE 20 D I in family with I in a single state of the	
•	OptMAVEn-2.0: De novo design of variable antibody regions against targeted antigen epitopes. Antibodies – MDPI * featured article	
	R Chowdhury, MF Allan, and CD Maranas. [Immunology Protein Structure]	/2018
	K Chowanary, Wir Anan, and CD Waranas. [minimiology Frotein Structure]	/2010
•	Highly active C8-acyl-ACP thioesterase variant isolated by a synthetic selection strategy.	
	ACS Synthetic Biology	
	N Hernandez, R Lai, <u>R Chowdhury</u> ,, BF Pfleger. [Enzyme Engineering Protein Function]	/2018
•	Facile affinity maturation of antibody variable domains using natural diversity	
	mutagenesis. Frontiers in Immunology	
	KE Tiller, R Chowdhury,, P Tessier. [Immunology Protein Structure]	/2017
	Using gene essentiality and synthetic lethality information to correct yeast and CHO cell	
	genome-scale models. Metabolites – MDPI	
	R Chowdhury, A Chowdhury, CD Maranas. [Metabolic Modeling Bio-networks]	/2015
_		
•	Conversion of slaughterhouse and poultry farm animal fats and wastes to biodiesel: Parametric sensitivity and fuel quality assessment. Renewable and Sustainable Energy Reviews	
	R Chakraborty, AK Gupta, R Chowdhury. [Biofuel Energy]	/2013
	R Charlacotty, 71R Gapat, R Chowanary. [Biotaer Energy]	/2013
•	Application of ANFIS model to optimize the photocatalytic degradation of chlorhexidine digluconate.	
	Royal Society of Chemistry – Advances	
	S Sarkar, <u>R Chowdhury</u> ,, C Bhattacherjee. [Process Optimization Neural Network]	/2013
	Interacting generalized cosmic Chaplygin gas in loop quantum cosmology:	
	Asingularity free universe. International Journal of Theoretical Physics	
	R Chowdhury, and P Rudra. [Cosmology Thermodynamics]	/2012
•	Influence of gauss-bonnet coupling parameter on the thermodynamic properties of Einstein-Gauss-Bonnet	
	and Einstein-Yang-Mills-Gauss-Bonnet black holes. International Journal of Theoretical Physics	/0015
	R Chowdhury, R Biswas,, S Chakraborty. [Cosmology Thermodynamics]	/2012

◆ PATENT

• US20190329189A1. Title: ANGSTROM-SCALE "SEPARATIONS BY DESIGN" USING PRECISION BIOMIMETIC MEMBRANE. (Penn State Invention Disclosure No. 2018-4764). [patents.google.com/patent/US20190329189A1/en]

◆ SELECTED MEDIA COVERAGE

◆ SELE	CTED MEDIA COVERAGE	
◆ SELF	 Science journal Blog by Derek Lowe Another Way to Do Protein Structure Prediction (Article link) ScienceMag Solving a Natural Riddle of Water Filtration (Article link) UT Austin Research Headlines Solving a Natural Riddle of Water Filtration (Article link) Alumni Spotlight Penn State University 2020 / Distinguished Alumni Series (Article link) Nature Nanotechnology News and Views A new type of artificial water channels. Nature Nanotechnology ScienceDaily Protein pores packed in polymers make super-efficient filtration membranes (Article link) Chemical & Engineering News (C&En) Filter from tropical tree seeds purges viruses from water (Article AIChE Separations Division Outstanding Researchers of 2019 (Article link) North American Membrane Society Student Fellows 2019 (Article link) UT Austin Research Headlines Bio-inspired new water filtration method (Article link) Penn State Chemical Engineering News Building better membranes (Article link) Penn State Research Headlines PoreDesigner process improves water treatment, bio-separations (Article Phys. Org PoreDesigner improves protein channel design for water treatment (Article link) Penn State Chemical Engineering News Research symposium – graduate student achievements (Article link) 	/2020 link) /2019 /2019 /2019 /2019 /2019 link) /2018 /2018
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	Single-sequence protein structure prediction and enzyme engineering Manifold Bio. Waltham, MA. Consequences of single-sequence protein structure prediction.	/2022
	Statistical Thermodynamics & Molecular Simulations Seminar Series. Yale University. CT.	/2022
	Single-sequence protein structure prediction and antimicrobial peptide design Dept. of Environmental Engineering. University of Illinois-Urbana Champaign. IL.	/2022
	Single-sequence protein structure prediction using deep learning Dept. of Environmental Engineering. University of Texas at Austin. TX.)21
	Protein design in the world of deep learning. Nested Therapeutics. Boston, MA.	/2021
	The future of protein engineering in the presence of AlphaFold2 and RGN2. Zymergen . Emeryville, CA.	/2021
	Novel protein engineering tool for predicting insertions and deletions along with substitutions. Enzyme Engineering Conference XXV . Whistler, British Columbia, Canada.	/2019
	Protein engineering for Separations. Popular lecture series Research Unplugged . Schlow Library. Penn State.	/2018
	New paradigms in protein design and precise aqueous separation devices. Bioinformatics and Genomics Annual Retreat. Huck Institute of Life Sciences. Penn State.	/2018
	CCTED ORAL AND POSTER PRESENTATIONS (speaker/ presenter) Structure-guided phylogeny of human kinome. CCSP-NIH. Virtual. USA.	/2022
	Single Sequence Protein Structure Prediction. CASP14 Conference. Virtual. USA.	/2020
	Single Sequence Protein Structure Prediction. NeurIPS Conference. Virtual. USA.	/2020
	PoreDesigner for precise tuning of OmpF for high solute selectivity STARTech Conference (NASA), TX. USA.	/2018
	PoreDesigner for engineering protein channels for precise angstrom scale bio-separations Gordon Research Conference, Membranes, NH. USA. *Best poster awardee	/2018
	Biomimetic pore design for precise angstrom scale bio-separations Gordon Research Conference, Membranes, NH. USA. *Best talk awardee	/2018

PoreDesigner for engineering protein channels for precise angstrom scale bio-separations Graduate Symposium Penn State University. PA. USA. *Best poster awardee	/2018
Computational substrate specificity switching of sorghum phenylalanine-ammonia lyase Center for Bio-Innovation, DOE Annual Meeting, TN. USA.	/2018
Redesign of E. coli water channel protein OmpF using Iterative Protein Redesign and Optimization Suite AIChE Annual Meeting , MN. USA. *Best talk awardee	/2017
Interacting Generalized Cosmic Chaplygin Gas in Loop Quantum Cosmology Marcel Grossmann Meeting, Stockholm. Sweden. *UG best poster awardee	/2012

◆ TEACHING AND MENTORING

During Ph.D.

• Several REU and graduate students mentored:

(a) describing project goals, (b) assigning deliverables, (c) setting up shared GitHub repositories for version control, (d) analyzing data, and (e) drafting manuscripts.

/2016-19

♦ SERVICE

Reviewer Activity

Nature Communications, Nature Scientific Data, Nature Scientific Reports, PNAS, Science, Cell Systems, Cell – Structure, PloS Computational Biology, ACS | Langmuir, Frontiers in Immunology, Metabolic Engineering, and Frontiers in Bioengineering and Biotechnology (Guest Editor: Special issue 2022 – Advances in Protein Structure, Function, and Design).