KRISHNA CHOUDHARY

EDUCATION		
	University of California, Davis, USA	2014-19

Ph.D., Biomedical Engineering with Designated Emphasis in Biotechnology
Dissertation:
Statistical methods & software for comparative analysis of RNA structurome profiling data
Advisor: Sharon Aviran

Indian Institute of Technology, Delhi, India   2008-13
B. Tech & M. Tech (Dual Degree Program)
Biochemical Engineering & Biotechnology (Major)
Nano Science & Engineering (Minor)

RESEARCH
EXPERIENCE

Postdoctoral Scholar, University of California, San Francisco, USA   2021-
Advisor: Michael McManus
Developed an integrated computational and wet-lab platform for microscopy-based pooled CRISPR screens and applied it to quantify genetic interactions underlying DNA damage

Bioinformatician II, Gladstone Institute of Data Science & Biotechnology, USA   2020-21
Supervisor: Alexander R. Pico, Director, Bioinformatics Core
Delivered computational solutions for analysis of a large variety of deep-sequencing data sets in projects related to cardiovascular and neurological diseases

Biostatistician I, Gladstone Institute of Data Science & Biotechnology, USA   2019
Supervisor: Eva Wang, Director, Bioinformatics Core
Delivered computational solutions for analysis of a large variety of deep-sequencing data sets in projects related to cardiovascular and infectious diseases

PUBLICATIONS/
PREPRINTS

Post-Ph.D.   2019-

21. **Choudhary, K.**[#], McManus, M.T.[#], Scalable imaging-based profiling of CRISPR perturbations with protein barcodes. *bioRxiv*, 2025. ([#]co-corresponding author)

20. Krup, A. L., ... five authors ..., **Choudhary, K.**, ... four authors ..., Bruneau. B., A Mesp1-dependent developmental breakpoint in transcriptional and epigenomic specification of early cardiac precursors. *Development*, dev.201229, 2023.

19. Zhu, L., **Choudhary, K.**, ... sixteen authors ..., Bruneau, B. G., Steinmetz, L., Krogan, N. J., Pollard, K. S., Srivastava, D., Transcription factor GATA4 regulates cell type-specific splicing through direct interaction with RNA in human induced pluripotent stem cell-derived cardiac progenitors. *Circulation*, 146, 770-787, 2022.

18. Gonzalez-Teran, B., ... five authors ..., **Choudhary, K.**, ... fifteen authors ..., Conklin, B. R., Black, B. L., Bruneau, B. G., Krogan, N. J., Pollard, K. S., Srivastava, D., Transcription factor protein interactomes reveal genetic determinants in heart disease. *Cell*, 185, 1-21, 2022.

17. Abouleisa, R. R. E., ... ten authors ..., **Choudhary, K.**, ... eleven authors ..., Srivastava, D., Bolli, R., Mohamed, T. M. A., Transient cell cycle induction in cardiomyocytes to treat ischemic heart failure. *Circulation*, 145(17), 1339-1355, 2022.

16. **Choudhary, K.**[#], Pico, A. R.[#], Introducing R as a smart version of calculators enables beginners to explore it on their own. *F1000Research*, 10(859), 2021. ([#]co-corresponding author)

15. **Choudhary, K.**, ... two authors ..., Bader, G. D., Pico, A. R., Morris, J. H., scNetViz: from single cells to networks using Cytoscape. *F1000Research*, 10(448), 2021.

14. Taubes, A., ... eleven authors ..., **Choudhary, K.**, ... eleven authors ..., Sirota, M., Huang, Y., Experimental and real-world evidence supporting the computational repurposing of bumetanide for APOE4-related Alzheimer's disease. *Nature Aging*, 1, 932–947, 2021.

13. Gulbranson, D., ... seven authors ..., **Choudhary, K.**, Thomas, R., Mucke, L., Phenotypic differences between the Alzheimer's disease-related hAPP-J20 model and heterozygous Zbtb20 knockout mice. *eNeuro*, 8(3), 2021.

12. **Choudhary, K.**#, Narang, A.#, Urn models for stochastic gene expression yield intuitive insights into the probability distributions of single-cell mRNA and protein counts. *Physical Biology*, 17(6), 066001, 2020. (#co-corresponding author)

11. Garcia, P. D., ... two authors ..., **Choudhary, K.**, ... three authors ..., Zakian, V. A., Stability and nuclear localization of yeast telomerase depend on protein components of RNase P/MRP. *Nature Communications*, 11(1), 1-19, 2020. [recommended by *Faculty Opinions*]

10. **Choudhary, K.**#, Addition formulas for the $_pF_p$ and $_{p+1}F_p$ generalized hypergeometric functions with arbitrary parameters and their Kummer- and Euler-type transformations. 2020. (#corresponding author) [*arXiv*]

9. **Choudhary, K.**#, Narang, A.#, Analytical expressions and physics for single-cell mRNA distributions of the *lac* operon of *E. coli*. *Biophysical Journal*, 117(3), 572-586, 2019. (#co-corresponding author)

Ph.D. 2014-19

8. **Choudhary, K.**, Lai, Y. H., Tran, E., Aviran, S., dStruct: identifying differentially reactive regions from RNA structurome profiling data. *Genome Biology*, 20(1), 40, 2019. [open-source Bioconductor package]

7. Lai, Y. H., **Choudhary, K.**, Cloutier, S. C., Xing, Z., Aviran, S., Tran, E., Genome-wide discovery of DEAD-Box RNA helicase targets reveals RNA structural remodeling in transcription termination. *Genetics*, 212(1), 153-174, 2019.

6. Watters, K. E., **Choudhary, K.**, Aviran, S., Lucks, J. B., Perry, K. L., Thompson, J. R., Probing of RNA structures in a positive sense RNA virus reveals selection pressures for structural elements. *Nucleic Acids Research*, 46(5), 2573-2584, 2018.

5. **Choudhary, K.***, Deng, F.*, Aviran, S., Comparative and integrative analysis of RNA structural profiling data: current practices and emerging questions. *Quantitative Biology*, 5(1), 3-24, 2017. (*co-first author)

4. **Choudhary, K.**, Ruan, L., Deng, F., Shih, N., Aviran, S., SEQualyzer: interactive tool for quality control and exploratory analysis of high-throughput RNA structural profiling data. *Bioinformatics*, 33(3), 441-443, 2016.

3. **Choudhary, K.**, ... four authors ..., Aviran, S., Metrics for rapid quality control in RNA structure probing experiments. *Bioinformatics*, 32(23), 3575-3583, 2016.

Undergraduate/Masters -2014

2. **Choudhary, K.**, Oehler, S., Narang, A., Protein distribution from a stochastic model of *lac* operon with DNA looping: analytical expressions and comparison with experiments. *PLOS ONE*, 9(7), e102580, 2014.

1. Grover, A.*, Pande, A.*, **Choudhary, K.***, Gupta, K.*, Sundar, D., Re-programming DNA-binding specificity in zinc finger proteins for targeting unique address in a genome. *Systems and Synthetic Biology*, 4(4), 323-329, 2010. (*co-first author)

TEACHING EXPERIENCE [GRADUATE]	**Guest Instructor**, UC Davis Topic: Single-cell RNA-seq Data Analysis Course: Statistical Genomics (BIM254) IOR: Sharon Aviran ~10 students in class; positive feedback from IOR; not rated by students	2022
	Content Developer & Lead Instructor, Gladstone Data Science Training Program Topics: Single-Cell RNA-seq Analysis; Bulk RNA-seq Analysis; Data Analysis & Visualization Workshops attended by Gladstone/UCSF graduate students, postdoctoral scholars, and faculty ~100 hours of instruction, cumulative attendance ~1000, mean student rating 4.54/5	2019-21
	Guest Presenter & Discussion Lead, UC San Francisco Topic: Dimensionality Reduction Course: Statistical Methods in Bioinformatics (BMI206) IOR: Katherine Pollard ~20 students in class; positive feedback from IOR; not rated by students	2020
	Guest Instructor & Teaching Assistant, UC Davis Course: Genomic Big Data Analysis (BIM289C) IOR: Sharon Aviran ~20 students in class; positive feedback from IOR; not rated by students	2015-16
	Teaching Assistant, IIT Delhi Courses: Advanced Biochemical Engineering (BEL850); Microbial Engineering (BEL713) IOR: Atul Narang ~20-30 students in class; positive feedback from IOR; not rated by students	2012-13
TEACHING EXPERIENCE [UNDERGRAD]	**Teaching Assistant**, UC Davis Course: Probability & Statistics (BIM105) IOR: David Rocke ~70 students in class; mean student rating: 4.0/5	2018
	President & Tutor, *Students Tutoring Students* club, UC Davis Tutored ~10 students in lower-division chemistry and mathematics courses for free Recruited multiple volunteer tutors and connected them with students Faculty advisor: Andreas Toupadakis	2015-17
TEACHING EXPERIENCE [BROADER COMMUNITY]	**Instructor/Instruction team member**, UCSF AI4All Program Lectured on supervised learning & led group discussion with diverse 9th-12th graders Program Director: Marina Sirota	2021-22
	GED Tutor, Sacramento Public Library	2016
OTHER WORK EXPERIENCE	**Bioinformatics Intern**, Roche Molecular Systems, USA Developed a machine learning classifier to call somatic variants identified in liquid biopsy The method was integrated in Roche's pipeline for data analysis	2017
	Biopharmaceutical Production Scale-Up Intern, Biocon Limited, India Interfaced between Biocon's Mammalian Cell Culture Group and their Pilot Plant Modeled and explained anomaly in gas transfer that was hindering scale-up of CHO cell cultures	2011

MENTORING EXPERIENCE	**Graduate students**	
	Prathamesh Chati, Rotation student, Biomedical Informatics, McManus lab, UCSF	2023
	Yuhao Wang, Ph.D. candidate, Biomedical Sciences, McManus lab, UCSF	2021-2023
	Yongin Choi, Rotation student, Biomedical Engineering, Aviran lab, UC Davis	2018
	Undergraduate students	
	Richard Phouasalith, Researcher, Biomedical Engineering, Aviran lab, UC Davis	2017
	Kyle Van Housen, Researcher, Biomedical Engineering, Aviran lab, UC Davis	2017
	Cassidy Dzoan, Researcher, Biomedical Engineering, Aviran lab, UC Davis	2017
	Huan Chen, International summer intern, GREAT program, Aviran lab, UC Davis	2016
	Qianyu Gao, International summer intern, GREAT program, Aviran lab, UC Davis	2015
	High-school students	
	Nishant Pitta, Software Developer intern, McManus lab, UCSF	2023-2024
	Chubi Yambao, E-Mentor program, Sheldon High School Biotechnology Academy, CA	2017
	Jem Doan, E-Mentor program, Sheldon High School Biotechnology Academy, CA	2016
	Deirdre Willgohs, Intern, Aviran lab, UC Davis	2016

PEER REVIEWER	BMC Bioinformatics, Nature Communications, The Journal of Chemical Physics, Proceedings of the Royal Society A, Biophysical Journal, Physical Biology, PLOS ONE, Scientific Reports, Entropy, Cancers, BMC Systems Biology, Nucleic Acids Research [28 articles reviewed]	2019-

COMMUNITY OUTREACH	Application reviewer, UCSF's AI4All program targeting diverse high schoolers	2021
	Invited speaker, seniors at Avenidas Village, Palo Alto, CA via *Skype a Scientist*	2021
	Career counselor, Douglass Middle School, Woodland, CA	2016
	Music teacher, Davis Mosaics initiative of the Davis Community Church, Davis, CA	2016

INVITED TALKS	Department of Bioengineering, Indian Institute of Science, Bangalore, India	2025
	National Centre for Biological Sciences, Bangalore, India	2025
	Department of Biology, Indian Institute of Science Education & Research, Pune, India	2025
	Joint Seminars in Molecular Biology, University of California, Davis, USA	2023

CONFERENCE PRESENTATIONS	**Choudhary, K.**, & McManus, M., Array of minipools: cost-effective scale up of microscopy-based CRISPR screens. **Frontiers in Comparative Systems Virology**; San Francisco, CA, USA.	2025
	Choudhary, K., & McManus, M., Array of minipools: cost-effective scale up of microscopy-based CRISPR screens. **Veridical Data Science for Biology**; Berkeley, CA, USA.	2025
	Choudhary, K., & McManus, M., Array of minipools: cost-effective scale up of microscopy-based CRISPR screens. **Separating the Signal from the Noise: AI in Biology**; San Francisco, CA, USA.	2025
	Choudhary, K., & McManus, M., Array of minipools: cost-effective scale up of microscopy-based CRISPR screens. **Chan Zuckerberg Biohub Confab**; San Francisco, CA, USA.	2025
	Choudhary, K., … three authors …, & McManus, M., epitoPeR enables scale up of high-content CRISPR screens. **Systems Approaches to Cancer Biology conference**; Woods Hole, MA, USA. [*selected for lightning talk*]	2022
	Choudhary, K., & Aviran, S., dStruct: a Bioconductor package for differential analysis of RNA structurome profiling data. **Annual Meeting of the RNA Society**; Online.	2021
	Choudhary, K., & Narang, A., Urn models for stochastic gene expression. **Biophysical Society Annual Meeting**; Online.	2021

Choudhary, K., & Narang, A., Probability distributions of single-cell mRNA and protein counts derived by solving urn models for stochastic gene expression. **Biological Data Science conference of Cold Spring Harbor Laboratory**; Online.   2020

Choudhary, K., Shih, N., & Aviran, S., Noise in RNA structural profiling data and its impact on structure prediction. **Annual Meeting of the RNA Society**; Prague, Czech Republic. 2017

Choudhary, K., ... four authors ..., & Aviran, S., Methods for rapid and scalable quality assessment of RNA structure probing data. **Biological Data Science conference of Cold Spring Harbor Laboratory**; New York, USA.   2016

Choudhary, K., ... four authors ..., & Aviran, S. Methods for rapid quality assessment of RNA structure probing data. **Computational RNA Biology conference of Wellcome Genome Campus**; Cambridge, UK.   2016

AWARDS	Postdoc Travel Grant, Helen Diller Family Comprehensive Cancer Center, UCSF	2022
&	Conference Award (sponsored by NSF) to attend the RNA Society annual meeting	2021
FELLOWSHIPS	Graduate Student Travel Award, Graduate Studies, UC Davis	2017
	Travel fellowship to attend RNA Society annual meeting, RNA Society	2017
	Biomedical Engineering Graduate Group Travel Award, UC Davis	2016
	Travel bursary, Wellcome Genome Campus, Cambridge, UK	2016
	Scholarship for all-India rank 26 in GATE test, Government of India	2012-13
	Summer Undergraduate Research Award, Industrial R&D Unit, IIT Delhi	2010