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VIVEK RAI

EDUCATION

2022 University of Michigan, Ann Arbor

> Ph.D. in Bioinformatics + M.A. in Statistics

2017 Indian Institute of Technology, Kharagpur

> Dual Degree (B.Tech & M.Tech), Biotechnology GPA: 8.62/10

RESEARCH EXPERIENCE

University of Michigan, Ph.D. Student 2018-now

Ann Arbor, US

Mentor: Stephen CJ Parker, Ph.D

- Integrative analysis of chromatin accessibility profiles across skeletal, pancreatic, and adipose tissues to assess the functional effects of non-coding mutations in type-II diabetes (T2D) disease susceptibility.

2016-2017 Indian Institute of Technology, Kharagpur, Senior Thesis Kharagpur, India

Mentor: Amit K. Das. Ph.D.

- Characterized the roles of structural and functional patterns in the origin of knotted proteins and how they may influence the knotting mechanism. (Talk)

2016 **INSTITUTE FOR SYSTEMS BIOLOGY, Summer Intern** Seattle, US

Mentor: Gustavo Glusman, Ph.D

 Analyzed systematic autosomal sex-specific biases in sequence coverage profiles in more than 2000 individual whole-genomes and discover technical mapping biases due to segmental duplication between autosomes and sex chromosomes.

Indian Institute of Science, B.E.S.T Fellow

Bangalore, India

Mentor: Sandhya S. Visweswariah, Ph.D

 Studied the assembly mechanism of Cytolysin-A (ClyA), a bacterial pore-forming toxin, and demonstrated a lipid phase and C-terminal domain dependent kinetic behavior using synthetic lipid vesicles and calcein leakage assays. (Report)

2015-2016 **WIKIMEDIA** — Individual Engagement Grant

Team Leader: Maximilien Klein

- Quantified gender bias in Wikipedia corpus as a team member of \$22,500 grant using Wikidata through statistical analysis, developed website, and assisted in research, and paper writing, editing and review.

2015

PROGRAMMING EXPERIENCE

| 2016 | GOOGLE SUMMER OF CODE, OpenSNP — Student Developer |
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| | Mentors: Bastian Greshake, Philip Bayer |
| | Project: Linking Phenotypes to Genotypes in OpenSNP (opensnp.org) |
| 2015-2016 | SEQUENCESERVER — Wurm Lab, Queen Mary University of London, UK |
| | Mentors: Anurag Priyam, Yannick Wurm, Ph.D. |
| | Project: Sequenceserver: a modern graphical user interface for custom BLAST databases |
| 2015-now | CONTRIBUTOR — Open Source Projects |
| | Afra, Bioconda, Scikit-bio, CobraPy, awesome-biology, samtools |

MANUSCRIPTS

| 2019 | Rai V [†] , Quang DX [†] , Erdos MR, Cusanovich DA, Daza RM, Narisu N, et al. Single cell ATAC-seq in human pancreatic islets and deep learning upscaling of rare cells reveals cell-specific type 2 diabetes regulatory signatures. bioRxiv. 2019;749283. |
|------|--|
| 2019 | Priyam A, Woodcroft BJ, Rai V , Moghul I, Munagala A, Ter F, et al. Sequenceserver: A Modern Graphical User Interface for Custom BLAST Databases. Mol Biol Evol. 10.1093/molbev/msz185/5549819. |
| 2018 | Grüning B, Dale R, Sjödin A, Chapman BA, Rowe J, Tomkins-Tinch CH, Köster J, & The Bioconda Team et al. Bioconda: sustainable and comprehensive software distribution for the life sciences. Nature Methods. 2018;15:475. |
| 2017 | O'Neill K, Rai V, Kilpatrick AM. The International Society for Computational Biology and WikiProject Computational Biology: celebrating 10 years of collaboration towards open access. Bioinformatics. 2017;33:2429–30. |
| 2016 | Klein M, Gupta H, Rai V, Konieczny P, Zhu H. Monitoring the Gender Gap with Wikidata Human Gender Indicators. Proceedings of the 12th International Symposium on Open Collaboration - OpenSym '16. Berlin, Germany: ACM Press; 2016 p. 1–9. |

Talks & Posters

| 2019 | (Poster, MIDAS Annual Symposium) — Single cell ATAC-seq in human pancreatic islets and deep learning upscaling of rare cells reveals cell-specific type 2 diabetes regulatory signatures. |
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| 2019 | (Poster, Department of Computational Medicine & Bioinformatics Retreat) — Single cell ATAC-seq in human pancreatic islets and deep learning upscaling of rare cells reveals cell-specific type 2 diabetes regulatory signatures. |
| 2019 | (Talk, Midwest Islet Club 2019) — Single-nuclei ATAC-seq in human pancreatic islets reveals cell-type specific type 2 diabetes regulatory signatures. |
| 2018 | (Poster , Department of Human Genetics Retreat) — Single-nucleus ATAC-seq reveals cell-specific diabetes regulatory signatures in human pancreatic islets. |
| 2018 | (Poster , Department of Computational Medicine & Bioinformatics Retreat) — Single-nucleus ATAC-seq reveals cell-specific diabetes regulatory signatures in human pancreatic islets. |
| 2015 | (Talk, Journal Club, IIT Kharagpur) — Protein Knots: A Tangled Story |

Skills

| interests | Computational Genetics & Genomics, Statistical Analysis, Data Science |
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| languages | Python, R, Shell Scripting (BASH), Ruby, SQL |
| web | HTML/CSS, ReactJS, Static-site Development |
| tools | Snakemake, Nextflow, Unix CLI, Conda, Slurm, Singularity Docker |
| cloud | Google Cloud Platform, Amazon Web Services (AWS) |
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Fellowships & Grants

| 2019 | Rackham Student Research Grant (\$1,500) — University of Michigan |
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| 2017 | Simons-NCBS Travel Fellowship (\$100) — NCBS, Bangalore |
| 2016 | Simons Foundation Travel Fellowship ($$600$) — ISMB 2016, Orlando, Florida |
| 2016 | GATE Scholarship (\$2,500) — University Grants Commission, MHRD, India |
| 2012-2016 | Merit-cum-means Tuition Scholarship [4x] (\$4,000) — IIT Kharagpur |
| 2012-2016 | Sanmarg Relief Scholarship [4x] (\$2,000) — Harsh and Payal Foundation |

Honors & Awards

| 2016 | First place (\$300), Wikidata Competition — International Society for Computational Biology |
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| 2016 | Third place (\$200), Wikipedia Competition — International Society for Computational Biology |
| 2012 | First place (\$200), Woodstock (Virtual Stock Trading Competition) — Kshitij, IIT Kharagpur |
| 2010 | Ram Avatar Pratibha Puraskar (excellence in Hindi language) — Sanmarg Pvt. Ltd. |

Conferences & Workshops

| 2017 | 5^{th} Simons-NCBS Monsoon School, Physics of Life 2017 — NCBS, Bangalore |
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| 2016 | Intelligent Systems for Molecular Biology (ISMB), Orlando, Florida |
| 2016 | Workshop on Mathematical and Computational Biology (WMCB 2016) — IISER, Kolkata |
| 2015 | Winter school on Quantitative Systems Biology (QSB) 2015 — ICTS, Bangalore |
| 2014 | Short Term Course on Computational Systems Biology 2014 — IIT Kharagpur |

VOLUNTEER ACTIVITIES

| 2018 | Peer Mentor — Bioinformatics, Univ. of Michigan |
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| 2019 | Peer Mentor — Program in Biomedical Sciences (PIBS), Univ. of Michigan |
| 2018-2019 | Student Host (Ph.D. interview weekend) — DCM&B, Univ. of Michigan |
| 2018-2019 | Student Representative, Bioinformatics Seminar Committee — DCM&B, Univ. of Michigan |
| 2018-2019 | Girls Who Code – DCM&B, Univ. of Michigan |
| 2018 | Juror (India), Wiki Science Competition 2018 (wikisciencecompetition.org) — Wikimedia |
| 2018 | Contributor, Women in Science 2018 Editathon — AWIS, Univ. of Michigan |
| 2012-2018 | Contributor (40+ articles and 1800+ edits) — English Wikipedia |
| 2016 | Co-organizer and Contributor, Wikipedia Editathon — ISMB 2016 |
| 2016 | Python and Git Assistant, Software Carpentry Workshop 2016 — University of Washington |
| 2012-2014 | National Service Scheme — IIT Kharagpur |

Professional Activities

| 2018 | Course (Tech) Reviewer, Python Tips, Tricks and Techniques — Packt Publishing |
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| 2015 | Problem Author, Ruby and Python Challenge Section — HackerRank |

Memberships

| 2019 | Fellow, AAAS/Science Program for Excellence in Science |
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| 2018-now | Board Member, Michigan Argentine Tango Club (MATC) — Univ. of Michigan |
| 2016 | Co-founder, Co-maintainer, MetaKGP (metakgp.github.io) — IIT Kharagpur |
| 2015 | Core Team Member, iGEM Sythetic Biology Team — iGEM IIT Kharagpur |
| 2015-2016 | Captain, Opensoft — LBS Hall, IIT Kharagpur |
| 2014 | Core Team Member, Google Students Club — IIT Kharagpur |
| 2013 | English Team Member, Technology Literary Society (TLS) — IIT Kharagpur |
| 2013 | Product Design Team Member, ProDex — IIT Kharagpur |
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