Vivek Ruhela

Postdoctoral Research Scientist, Columbia University

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Research Statement

Computational biologist investigating how genomic variation and artificial intelligence converge to decode the biology of complex diseases. I develop scalable and reproducible pipelines for genetics (Runs of Homozygosity (ROH), Genome-wide Association Study (GWAS), and expression quantitative trait loci (eQTL) finemapping) and genomic modeling (bio-inspired GCN frameworks, integrative transcriptomic–genetic analysis, and end-to-end bioinformatics workflows) to uncover disease mechanisms across diverse populations.

Education

2018 – 2024 **P.hD., Computational Biology**, IIIT-Delhi

Research Area - Cancer Genomics.

Thesis title: Design and development of AI-based computational tools for identifying predictive biomarkers and signaling pathways for blood cancer.

Advisors: Prof. Anubha Gupta (IIIT Delhi) and Prof. (Dr.) Ritu Gupta (AIIMS, Delhi)

2014 – 2016 M.Tech., Instrumentation & Control ZHCET, Aligarh Muslim University.

CPI – 9.0 (First Class Honours)

Thesis title: Detection of Brain tumor in Brain MRI (Magnetic Resonance Imaging).

Advisor: Prof. Yusuf Uzzaman Khan

2006 – 2010 **B.Tech. Electrical Engineering**, BSACET College, AKTU.

Marks - 68.78%

2003 – 2004 **12th Std., P.D.D.S.V.M, Vrindavan**.

Board – C.B.S.E.

Marks - 74.40%

2001 – 2002 **I** 10th Std., P.D.D.S.V.M, Vrindavan.

Board – C.B.S.E. Marks – 79.20%

Research Experience

May 2024 - *Till Now*

Postdoctoral Research Scientist

Department of Neurology, Columbia University Irving Medical Center (CUIMC), New York, USA.

Project: Genetic fine-mapping of Alzheimer's disease using multi-omics integration, Runs of Homozygosity (ROH), and eQTL analysis across diverse populations. *Key Contributions:* Developed scalable pipelines for ROH detection, ancestry-specific meta-analysis, and eQTL–GWAS colocalization across populations.

Research Experience (continued)

2018 - 2024

■ Doctoral Researcher (Ph.D. in Computational Biology)

Indraprastha Institute of Information Technology Delhi (IIIT-Delhi), India. *Thesis:* AI-driven biomarker discovery and genomic modeling in Multiple Myeloma.

Key Contributions: Developed miRPipe, miRSim, and Bio-DGI frameworks for small-RNA sequencing and AI-based multi-omics integration to identify novel biomarkers distinguishing MGUS and MM.

2015 - 2016

Graduate Researcher (M.Tech. Thesis)

Aligarh Muslim University (AMU), Aligarh, India.

Thesis: Detection of brain tumors in MRI using image processing and machine learning techniques.

Tools: MATLAB, SVM, and feature extraction methods for biomedical imaging.

Teaching

Jan 2017 – Dec 2017 Assistant Professor Electrical and Electronics Dept., PSIT.

Aug 2016 – Dec 2016 Assistant Professor. Electrical Department, Sanskriti University.

Oct 2012 – March 2013 Course Instructor (Ad-hoc) Electrical Department, ADKMP College.

Jan 2013 – March 2013 Lecturer(Ad-hoc) Electrical Department, BSA College.

Aug 2010 – Sept 2012 Lecturer. Electrical Department, Mangalayatan University.

Research Publications

Journal Articles

- 1 Cieza, B., Pandey, N., **Ruhela, Vivek**, Ali, S., & Tosto, G. (2025). **SAGA** (**S**implified **A**ssociation **G**enomewide **A**nalyses): A user-friendly pipeline to democratize genome-wide association studies. *bioRxiv*, 2025–08.
 6 doi:https://doi.org/10.1101/2025.08.25.672146
- Ruhela, Vivek, Gupta, R., Oberoi, R., & Gupta, A. (2025). A comprehensive targeted panel of 295 genes: Unveiling key disease initiating and transformative biomarkers in multiple myeloma. *Computers in Biology and Medicine*, 196, 110619. Odoi:https://doi.org/10.1016/j.compbiomed.2025.110619
- Farswan, A., Gupta, A., Jena, L., **Ruhela, Vivek**, Kaur, G., & Gupta, R. (2022). Characterizing the mutational landscape of mm and its precursor mgus. *American journal of cancer research*, 12(4), 1919. Retrieved from 6 https://pubmed.ncbi.nlm.nih.gov/35530275/
- **Ruhela, Vivek**, Gupta, A., Sriram, K., Ahuja, G., Kaur, G., & Gupta, R. (2022). A unified computational framework for a robust, reliable, and reproducible identification of novel mirnas from the rna sequencing data. *Frontiers in Bioinformatics*, *2*, 842051. 6 doi:10.3389/fbinf.2022.842051
- Vivek Ruhela, Gupta, R., Krishnamachari, S., Ahuja, G., & Gupta, A. (2021). miRSim: Seed-based Synthetic Small Non-coding RNA Sequence Simulator. Zenodo. https://doi.org/10.5281/zenodo.6546356. Odoi:https://doi.org/10.5281/zenodo.6546356
- Ruhela, Vivek, Farswan, A., Gupta, A., Sriram, K., Kaur, G., & Gupta, R. (2021). P-035: Ai-based models for the identification of critical genetic biomarkers to distinguish mm from mgus using the wes data. Clinical Lymphoma Myeloma and Leukemia, 21, S57. Odoi:10.1016/S2152-2650(21)02169-8

Kaur, G., **Ruhela, Vivek**, Rani, L., Gupta, A., Sriram, K., Gogia, A., ... Gupta, R. (2020). Rna-seq profiling of deregulated mirs in cll and their impact on clinical outcome. *Blood cancer journal*, 10(1), 1–9. Odoi:10.1038/s41408-019-0272-y

Conferences

- Pandey, N., **Ruhela, Vivek**, Cieza, B., Barral, S., Samper-Ternent, R., Montesinos, R., ... Tosto, G. (2025). Genome-wide meta-analysis of cognitive performances in hispanics/latinos identifies novel rare variant locus pcats. In *Alzheimer's & parkinson's diseases conference*. Retrieved from http://dx.doi.org/10.13140/RG.2.2.24659.90407
- Ruhela, Vivek, Cieza, B., Mayeux, R., Reyes-Dumeyer, D., Teich, A. F., & Tosto, G. (2025). Genetic colocalization of expression quantitative trait loci (eqtl) mapping and gwas in a multiethnic brain bank: An insight into ancestry-specific regulatory architecture in alzheimer's disease. In *Alzheimer's association international conference*. ALZ. Retrieved from https://alz.confex.com/alz/2025/meetingapp.cgi/Paper/107199
- Vivek Ruhela, Panday, N., Cieza, B., Barral, S., Samper-Ternent, R., Montesinos, R., ... Tosto, G. (2025). Fine mapping of regions of homozygosity and their role in alzheimer's disease: Insights from the peruvian and mexican populations. In *Alzheimer's & parkinson's diseases conference*. Retrieved from ₱https://doi.org/10.13140/RG.2.2.15642.15048
- Ruhela Vivek, Yang, Z., Jacobson, S. W., Jacobson, J. L., Meintjes, E. M., Tosto, G., & Carter, R. C. (2024). Gene by prenatal alcohol exposure interaction effects on growth and cognition in mother-child dyads in a south african birth cohort. In *American society of human genetics (ashg)*. Retrieved from http://dx.doi.org/10.13140/RG.2.2.17434.56006

Skills

Languages Strong reading, writing and speaking competencies for English.

Coding MATLAB, Python, R, Bash, Awk, Lager MATLAB, Python, R, Bash, Awk, Lager Matlabet Matla

Misc. Academic research, teaching, training

Miscellaneous Experience

Awards and Achievements

2010 Competition Prize for Outstanding Performance in MATLAB Code Challange, BSA Engg.

Certification

2017 Medical Image Analysis. Elite certification in NPTEL Online certification.

Enhancing Soft Skills and Personality. Elite certification in NPTEL Online certification.

References

Available on Request