

Nima Sarfaraz, Ph.D.

Molecular and Cell Biology and Genetics | Wet Lab and Bioinformatics | Project Leadership

Email & Tel. available upon request – www.nima.bio/contact

Molecular biologist combining wet lab experimental expertise with advanced computational analysis and enterprise-level project management from big data consulting. Specialized in cancer biology, virology, and multi-omics approaches with proven ability to translate complex research into actionable therapeutic targets. Track record of high-impact deliverables, cross-functional leadership, and mentoring in academic and commercial environments.

SKILLS

Wet lab:

Molecular Biology: Vector design and implementation (adenoviral and lentiviral packaging, plasmids, siRNAs, transposons), molecular cloning, RT-qPCR, PCR, blotting (Western, Northern, Southern), RACE, extractions (protein, DNA, RNA)

Cellular Techniques: BSL-2 mammalian cell culture (immortalized cell lines, primary hepatocytes), transfections, transductions, viral infections, stable cell line engineering and generation, banking, aseptic technique, bacterial transformations and culture

Analytical Assays: Gene perturbation experiments, metabolic flux analysis, plate-based assays (ELISA, OPP-incorporation protein synthesis, luciferase), colony formation, drug and compound treatments

Computational and Bioinformatics:

Programming: R, Python, UNIX/Linux, Bash scripting, SQL, NoSQL, high-performance computing environments, Slurm, Git

Analyses & pipelines: NGS / Next-generation sequencing (Illumina, Oxford nanopore), transcriptomics, proteomics, Nextflow, nf-core, quality control, alignment, quantification, batch effect integration, differential expression, pathway enrichment, hierarchical clustering, data visualization, biomarker and target identification

Databases: TCGA, GTEx, GEO, ICGC, BioMart, genome browsers (UCSC, NCBI, Ensembl)

Software: GraphPad Prism, ImageJ, IGV, ELNs, Adobe (Photoshop, Illustrator), MS Office Suite, local/commercial LLMs

Project Management: Agile and Scrum methodologies, sprint planning, stakeholder management, cross-functional team collaboration, requirements gathering, documentation

RESEARCH EXPERIENCE

DREXEL UNIVERSITY COLLEGE OF MEDICINE, BOUCHARD LAB

Philadelphia, PA

Doctoral Researcher

2020-2025

Project 1: Hepatic Tumor Suppressor Target Identification and Characterization

- **Identified and functionally validated a tissue-specific long non-coding RNA (lncRNA) as a therapeutic target and biomarker in liver cancer**, narrowing candidates through in vitro validation and database mining that correlated with significant clinical patient survival outcomes
- Elucidated novel mechanism of action through adenovirus design, transductions, RNA-seq experiments, TRENDS, pathway enrichment analyses, and protein synthesis assays in both cancer cell lines and primary human hepatocytes
- Engineered tetracycline-inducible stable cell lines using lentiviral and PiggyBac transposon systems, enabling temporal gene expression control for mechanistic studies and establishing renewable cell line resources for the lab

Project 2: Transcriptomic Benchmarking for Optimal Cancer Model Selection

- **Developed computational framework analyzing 541 bulk RNA-seq samples to systematically compare liver cancer cell lines against primary tumor subtypes**, enabling evidence-based experimental model selection and addressing a critical gap in cancer research reproducibility
- Pioneered analysis pipeline integrating GEO database mining, quality control, batch correction, differential expression analysis, and hierarchical clustering in high-performance computing environments to establish comprehensive molecular maps of heterogeneous liver cancer subtypes
- Identified critical mischaracterizations in widely-used cell lines, helping researchers improve reliability and avoid incorrect contexts and errors

Lab Development:

- Mentored 15 rotation and visiting students in laboratory techniques and experimental design, with 5 progressing to co-authorship on manuscripts
- Established 2 key cross-lab research collaborations, wrote joint R01 and R21 grants
- Initiated and developed a lab-wide digital centralized repository for common protocols and metadata, standardizing procedures for better reproducibility between colleagues and improved onboarding efficiency for new lab members

OTHER PROFESSIONAL EXPERIENCE

ACCENTURE

Washington, D.C.

Technology Consultant (Hybrid Scrum Master / Product Owner)

2017-2019

Led a 12-person Agile-based development team delivering custom big data software and dashboard integration projects, serving as a requested primary interface to stakeholders from a 'diamond client' (\$100+ million revenue per year) account during the software development lifecycle

- Leveraged system architecture expertise to develop user requirements, technical specifications, and business logic for new features while ensuring seamless integration with existing databases, APIs, and data flows
- Authored monitoring scripts that were adopted into the company's 24/7 production dashboard metrics, improving visibility into system performance and early issue detection
- Oversaw production releases and documentation, provided on-call emergency support

Technology Consulting Senior Analyst

2015-2017

- Mentored 4 junior analysts on backend testing methodologies; developed standardized workflow adopted as official protocol for developer-to-tester handoffs that reduced error rates leading up to production deployments
- Served as subject matter expert for critical system components, consulted by stakeholders and leadership; generated and maintained interface control documentation for core system components
- Performed root cause analysis using SQL/CQL and data analytics

Technology Consulting Analyst

2014-2015

- Conducted backend testing for feature implementations across database storage, JSON messaging, and system logic
- Obtained public trust clearance

EDUCATION

DREXEL UNIVERSITY

Philadelphia, PA

Ph.D., Molecular and Cell Biology and Genetics

- Awards: Drexel University Graduate College Outstanding Mentor Award; 1st Place: Best Trainee Poster Presentation (Sydney Kimmel Cancer Center Trainee Retreat)
- Elective coursework in Biochemical Data
- GPA: 4.00

M.S., Molecular and Cell Biology and Genetics

- Elective coursework in Cancer Biology, Drug Discovery and Development
- GPA: 4.00

UNIVERSITY OF MARYLAND COLLEGE PARK

College Park, MD

B.S., Bioengineering

- Dean's Scholarship Award

FIRST-AUTHOR PUBLICATIONS

- Sarfaraz, N., Kaur, R., Harper, S., Oni, L., Somarowthu, S., & Bouchard, M. J. (2025). **The Liver-Enriched Long Non-Coding RNA FAM99A Suppresses Tumorigenesis through Negative Regulation of Protein Synthesis.** *Under Review* *Journal of Molecular Biology*. <https://doi.org/10.1101/2025.04.29.651319>
- Sarfaraz, N., Winkler, T., Patel, R., Rosen, G., Somarowthu, S., & Bouchard, M. J. (2025). **Transcriptomic Benchmarking of Liver Cancer Cell Line Models Reveals Optimal Matches for Tumor Subtypes.** *Under Review* *Scientific Reports*.
- Sarfaraz, N., Somarowthu, S., & Bouchard, M. J. (2022). **The interplay of long noncoding RNAs and hepatitis B virus.** *Journal of Medical Virology*, 95(2), e28058. <https://doi.org/10.1002/jmv.28058>