

Nur M Shahir, PhD

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EDUCATION

University of North Carolina at Chapel Hill, Chapel Hill, NC, USA

- Ph.D. in Bioinformatics and Computational Biology Aug 2014 – May 2020
 - Dissertation: Inflammatory Bowel Disease Differentially Affects Region Specific Composition and Aerotolerance Profiles of Mucosally-Adherent Bacteria
 - Adviser: Dr. Terrence S. Furey, Dr. Shezad Z. Sheikh
 - Committee: Dr. Michael I. Love, Dr. Ian Carroll, Dr. Yufeng Liu
 - Focus: Inflammatory Bowel Disease, gut microbiota, bioinformatics, 16S amplicon sequencing.

University of Maryland, Baltimore County, Baltimore, Maryland, USA

- M.S. in Statistics Aug 2011 – Dec 2013
 - Track: Biostatistics
 - Thesis: Longitudinal Analysis of Urea Cycle Disorder Patients
 - Adviser: Dr. DoHwan Park

Massachusetts Institute of Technology, Springfield, Pennsylvania, USA

- B.S. in Mathematics 2006 – 2010

RELEVANT INDUSTRY EXPERIENCE

Booz Allen Hamilton Remote

- Lead Scientist Aug 2022 – Oct 2024
 - Served as a federal contractor bioinformatician, contributing to public health genomic surveillance, pathogen genomics, and large-scale human genomics projects across CDC and NIH.
 - Redesigned and optimized public health genomics workflows to align with Nextflow nf-core standards, implementing rigorous pipeline validation and automated unit testing. Streamlined collaboration through Git, ensured reproducibility via Docker containerization, and enhanced project delivery efficiency by integrating Agile tracking in JIRA.
 - Engineered reproducible bioinformatics pipelines with Snakemake on high-performance computing (HPC) clusters, enabling robust rare variant detection and uncovering genetic associations with ulcer development in sickle cell disease.
 - Designed and implemented bioinformatics pipelines in R and Python on Google Cloud Platform (GCP) via the NIH All of Us Researcher Workbench, spanning genomic data ingestion, QC, and advanced downstream analysis. Delivered cloud-native, production-ready workflows that accelerated large-scale genomic insights for precision medicine applications.

RESEARCH EXPERIENCE

Davenport Lab, Pennsylvania State University

- Postdoctoral Fellow Jun 2020 – Jul 2022
 - Piloted a benchmarking study on computational approaches to identify viral transcripts from bulk and single-cell RNA sequencing data, assessing the accuracy and precision of transcript identification.
 - Designed bioinformatics workflows in snakemake for efficient data processing on HPC computing environments, employing tools including samtools, bwa, bowtie2, Kraken2, and STAR, and utilizing R for downstream analysis and visualization.
 - Mentored undergraduate students, enhancing their research skills and academic performance.

Furey Lab, University of North Carolina at Chapel Hill

- Graduate Research Assistant Nov 2014 – May 2020
 - Identified key microbial associations using R for data analysis and visualization of 16S rRNA amplicon data from IBD patients and controls.
 - Developed a consensus analysis method in R with DESeq2 and Lefse, significantly enhancing the understanding of microbial dysbiosis in IBD.
 - Presented research findings at various conferences, including the American Society of Human Genetics, to engage the scientific community.
 - Authored a peer-reviewed journal article, contributing to the advancement of knowledge in the field of microbial dysbiosis.

PUBLICATIONS

Shahir, NM, et.al, “Crohn’s Disease Differentially Affects Region-Specific Composition and Aerotolerance Profiles of Mucosally Adherent Bacteria.” *Inflammatory bowel diseases*, vol. 26, no. 12, pp. 1843–1855, 2020.

PRESENTATIONS EXTERNAL TALKS

- *Crohn's Disease Differentially Affects Intestinal Region Composition and Aerotolerance Profiles of Mucosally-Adherent Bacteria*, Remote May 2020
Virtual Microbiome Summit
- *IBD differentially affects region specific composition and aerotolerance profiles of mucosal-adherent bacteria*, Remote May 2020
MIT and UNC Joint Virtual Microbiome Seminar Series

UNIVERSITY OF NORTH CAROLINA AT CHAPEL HILL

- *Crohn's Disease and the Intestinal Microbiota*, Chapel Hill, NC Dec 2016
Genetics Research Colloquium
- *Alterations in the Mucosal-Adherent Enteric Microbiota Between CD and nonIBD*, Chapel Hill, NC Oct 2016
Translations Medicine Closed Door Talks
- *A distinct microbiota signature characterizes patients with penetrating Crohn's disease*, Chapel Hill, NC Oct 2015
Center for Gastrointestinal Biology and Disease
- *Analysis of the Composition and Diversity of the Colonic Mucosa Microbiota in Crohn's Disease*, Chapel Hill, NC May 2015
Bioinformatics and Computational Biology Curriculum New Student Talks
- *Identification of SERPINA1 Splice Variants from Next-Gen Sequencing Data*, Chapel Hill, NC Oct 2014
Bioinformatics and Computational Biology Research in Progress Talks

HONORS & AWARDS

- NIH T32 Training Grant 2015
Bioinformatics and Computational Biology Predoctoral Training Grant

PROFESSIONAL AFFILIATIONS & ACTIVITIES

Joint Society of Earth Scientists and Global Think Tank on Climate Resiliency,
North Attleborough, Massachusetts, USA

- Member 2009 – Present

**CAMPUS
ACTIVITIES**

First Volunteers Club, First American University

Aug 2006 – Aug 2007

- President
 - Lorem ipsum dolor sit amet, consectetur adipiscing elit.
 - Curabitur vitae laoreet velit, vel ultricies est. Nam nec elit ac ante facilisis ultrices.
 - Wel turpis efficitur, non lacinia orci maximus.
 - Proin rhoncus, felis vel hendrerit lacinia.
 - Integer sit amet turpis dolor. Lorem ipsum dolor sit amet, consectetur adipiscing elit. Nunc at orci eu leo vulputate finibus sed et sem.
 - Suspendisse volutpat sapien et mi cursus, gravida ornare mauris sollicitudin.

**OTHER WORK
EXPERIENCE**

Alpha Engineering Firm, Oakland, Ohio, USA

Oct 2007 – Jan 2008

- Project Officer, Department of Meteorological Sciences, Research & Development Division
 - Nullam venenatis egestas nisl eget elementum.
 - Nulla finibus justo vel turpis efficitur, non lacinia orci maximus. Proin rhoncus, felis vel hendrerit lacinia, enim ipsum ultricies massa, sit amet interdum nisi massa sit amet justo.
 - Etiam vitae eros mollis, consectetur quam quis, molestie massa.

LANGUAGES

- English: Native language.
- Spanish: Fluent (speaking, reading, writing).
- Latin: Intermediate (reading); basic (speaking, writing).

SKILLS

T_EX, L^AT_EX, X_YL^AT_EX, MATLAB, Mathematica, Maple, R, Tableau, Adobe Photoshop, Adobe Illustrator, Microsoft Word, Microsoft Excel, Microsoft PowerPoint.

INTERESTS

Digital photography, typography, swimming.

REFERENCES

- **Professor Jonathan Public**
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- **Dr Alice Bob Carol**
Director, Research & Development
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