

# Arnoneel Sinha, PhD

*Post-Doctoral Fellow, CBGM, Augusta University*

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Augusta, GA

## Professional Summary

Ph.D.-trained Biostatistician and Postdoctoral Fellow at Augusta University with expertise in bioinformatics, statistical methodology, and data science education. My research bridges methodological development and collaborative applications in multi-omics and translational health data. I have experience leading analytical workflows, mentoring graduate students, and co-authoring NIH and NSF grant proposals. I am passionate about building an independent research program that unites statistical innovation with biomedical discovery while contributing to teaching and mentorship in biostatistics and data science.

## Skills

### Technical Skills

**Statistical Analysis:** Hypothesis testing, regression analysis, sample size estimation, experimental design, ANOVA, GLM.

**Multi-omics Data Analysis:** Analysis of transcriptomics (bulk RNA-seq, scRNA-seq) and proteomics (mass spectrometry); experience with data integration and normalization across platforms.

**Single-Cell Analysis:** End-to-end scRNA-seq workflows including QC, normalization, clustering, differential expression, cell type annotation (automated and manual), cell-cell communication (CellChat), and metabolic flux estimation (scFEA).

**Bioinformatics Tools:** Proficient in Seurat, DESeq2, Cell Ranger, STAR, HTSeq, Bioconductor, Enrichr, GSEA.

**Machine Learning:** Experience with supervised/unsupervised algorithms, time-to-event modeling, changepoint detection, survival analysis.

**Data Management:** Handling large-scale biomedical datasets, data cleaning, preprocessing, transformation, integration.

**Data Visualization:** Building plots, dashboards, and interactive tools using ggplot2, Shiny, matplotlib, and custom scripts.

**Statistical Modeling:** Frequentist and Bayesian approaches, mixed effects models, time series analysis.

**Protocol and Analysis Plans:** Designing statistical protocols for studies and grants; experienced in writing statistical analysis plans for collaborative biomedical projects.

**High-Performance Computing:** Experience running bioinformatics workflows on HPC systems using SLURM, Linux, and Bash scripting.

### Softwares

R (Seurat, DESeq2, ggplot2, Shiny), Python (pandas, sklearn, matplotlib), SAS (macros), L<sup>A</sup>T<sub>E</sub>X, Bash

### Platforms & Tools

Git/GitHub, Microsoft Office, SLURM, HPC environments, RStudio, Jupyter, VSCode

### Languages

English (professional), Hindi, Bengali

## Professional Experience

### Post-Doctoral Fellow

*Dr. Richard McIndoe lab, Center for Biotechnology & Genomic Medicine, Augusta University*

**Aug 2023 - Present**

*Augusta, GA*

- Led analyses of more than 15 scRNA-seq and bulk RNA-seq datasets across multiple biomedical studies, including tumor and pancreatic tissues; developed end-to-end bioinformatics pipelines in R for differential expression, cell type annotation (manual and automated), and pathway enrichment using Seurat, DESeq2, and Loupe Browser
- Collaborated with interdisciplinary investigators to deliver advanced statistical and bioinformatics support using R and Python; co-authored multiple grant proposals (NIH, NSF) and research manuscripts involving scRNA-seq and multi-omics data analysis
- Mentored graduate students in computational genomics projects

**Graduate Research Assistant (Teaching Assistant)**  
*Augusta University*

**Aug 2018 - July 2023**  
*Augusta, GA*

- Provided statistical consulting on multidisciplinary projects, taught undergraduate math/statistics, and gained expertise in clinical trials with applications of machine learning and advanced modeling.

**Senior Subject Matter Expert**  
*Nerdy Turtlez*

**Dec 2017 - Mar 2018**  
*Kolkata, India*

- Led and executed data analytics and visualization projects, conducting comprehensive statistical analyses to extract insights.
- Developed customized reports and presentations, creating informative visualizations tailored to client-specific needs.

#### **Research Mentor, Undergraduate Training Program**

*Statistics Undergraduate Research Experience (SURE), Southern Regional Council on Statistics (SRCOS)*

- Mentored undergraduate STEM majors during the SURE Summer Research Conferences (SRCOS 2022 and 2023), providing guidance on graduate school and career opportunities in statistics.
- Collaborated with session leaders to facilitate discussions and address student questions during mentoring sessions.

## **Teaching Experience**

Experienced in teaching and mentoring undergraduate and graduate students in statistics, mathematics, and computational genomics. Instructional philosophy emphasizes reproducible research, applied data analysis, and interdisciplinary learning.

Taught undergraduate and graduate-level courses in statistics, mathematics, and computational genomics in both in-person and hybrid formats. Responsibilities included lecture and slide preparation, grading, office hours, one-on-one mentoring, and collaborative instruction with faculty. Experience spans traditional classroom teaching as well as project-based, hands-on computational training in genomics and bioinformatics.

#### **Graduate Courses:**

##### **Co-Instructor, Translational Genomics and Proteomics (GNMD 8051)**

Spring 2025

- Co-instructed a graduate-level course for 2nd year biomedical students on advanced genomic and proteomic technologies and their applications in translational medicine
- Delivered a lecture on genomic approaches to disease classification and prediction, integrating clinical relevance with computational methods
- Supported students in critical reading of omics literature, project-based assignments, and discussions on emerging genomic tools

##### **Co-Instructor, Computational Methods in Genomics and Genetics (GNMD 8050)**

Fall 2024, 2025

- Co-taught a 4-credit graduate-level course combining lectures and supervised labs, introducing students to computational and statistical techniques in genomic and genetic research
- Led multiple sessions covering R programming, data visualization, statistical methods, and machine learning techniques including regression, clustering, dimensionality reduction, and neural networks
- Collaborated with faculty to design and evaluate student projects and assessments, and supported hands-on learning through case-based data analysis

#### **Undergraduate Courses:**

##### **• Elementary Statistics (MATH 1401)**

Hypothesis testing, regression, ANOVA, CLT, etc.

Fall 2021

##### **• Pre-Calculus Mathematics (MATH 1113)**

Polynomial/exponential functions, trigonometry, unit circle

Fall 2020, Spring 2021

# Publications & Presentations

## Publications

- **Arnoneel Sinha**, Richard A. McIndoe. *cellivista: An Interactive Seurat-Based R Shiny App for Single-Cell RNA-seq Analysis.* (Research article, under preparation)
- Kimberly A. Rivera-Caraballo, Tae Jin Lee, **Arnoneel Sinha**, Marco Orecchioni, Rafal Pacholczyk, Karina Vázquez-Arreguín, Shilpa Sharma, Kimya Jones, Kailash Vemuri, Upasana Sahu, Sara A. Murphy, Bangxing Hong, Ravindra Kolhe, Ashok Sharma, Balveen Kaur; Oncolytic HSV-1-Mediated JAG1 Blockade Induces Glioma Senescence-Associated Secretory Phenotype to Increase Macrophage Activation and Cetuximab-Mediated Senolysis. *Cancer Res* 2025; <https://doi.org/10.1158/0008-5472.CAN-25-1402>
- **Sinha A**, Chen J. *On the Detection of Multiple Change Points in a Sequence of Bivariate Normal Random Vectors with Application to COVID-19 Data.* (Research article, under review)
- **Sinha A**, Chen J. *On the Detection of Multiple Change Points in a Sequence of Bivariate Negative Binomial Random Vectors with Application to COVID-19 Data.* (Research article, under preparation)
- James JN, Frazier KB, Looney SW, **Sinha A**, Faigen AB. *Does a Difference Exist in Comprehensive Basic Science Examination Scores of 4-Year versus 6-Year Oral-Maxillofacial Surgery Residents?* *J Oral Maxillofac Surg.* 2020 Sep;78(9):1459.e1-1459.e6. doi: 10.1016/j.joms.2020.04.007. Epub 2020 Apr 18. PMID: 32413336.

## Presentations

- **Arnoneel Sinha**, Richard A. McIndoe. *cellivista: An Interactive Seurat-Based R Shiny App for Single-Cell RNA-seq Analysis.* **Health Sciences Fair and Expo**, Augusta University, GA, 2025. (Poster presentation)
- **Arnoneel Sinha**, Jie Chen. *Detecting Multiple Change Points in Bivariate Negative Binomial Model with Application to COVID-19 Incidence and Death Counts.* **Summer Research Conference, Southern Regional Council on Statistics**, Waco, TX, 2023. (Poster presentation)
- **Arnoneel Sinha**, Jie Chen. *Detecting Multiple Change Points in Bivariate Negative Binomial Model with Application to COVID-19 Incidence and Death Counts.* **The 3rd Annual Workshop: Emerging Data Science Methods for Complex Biomedical and Cyber Data**, Augusta University, Augusta, GA 2023. (5 minute presentation)
- **Arnoneel Sinha**, Jie Chen. *Detection of Multiple Change Points in Bivariate Data: Application to COVID-19 Incidence and Death Rates.* **Summer Research Conference, Southern Regional Council on Statistics**, Jekyll Island, GA, 2022. (Poster presentation)

## Education

**Doctor of Philosophy (Biostatistics)**  
*Augusta University*

**Aug 2018 - July 2023**

**Dissertation Title:** On the Detection of Multiple Change Points in A Sequence of Bivariate Random Vectors with Application to COVID-19 Data

**Master of Science (Statistics)**  
*Visva Bharati University*

**2015 - 2017**

**Thesis Title:** An overview of Process Capability Analysis

**Bachelor of Science (Statistics)**  
*Visva Bharati University*

**2012 - 2015**

**Thesis Title:** Juvenile Apprehension in India (2001-2012): A Statistical Perspective

# Academic Projects & Theses

## Selected Academic Projects & Theses

- **A Review on Gene Set Analysis**

*'Omics Data Analysis' course final project*

- **A Comparison of Five Machine Learning Techniques Predicting Time to Failure in Laboratory Earthquakes**

*'Statistical and Machine Learning for Big Data' course final project*

- **An overview of Process Capability Analysis**

*Masters' Thesis*

- **Juvenile Apprehension in India (2001-2012): A Statistical Perspective**

*Bachelor's Thesis*

## Current Projects

- Downstream Single Cell RNA-Seq Data Analysis with RShiny: A User-Friendly Tool for Biologists
- Comparison of Existing and LLM-based Automated Cell Type Annotation Methods for Single-Cell RNA-Seq Data

# Awards & Honors

- Winner of '*People's Choice Award*' in the 3MT competition at Augusta University (03/29/2023)
- '*NSF Travel Award*' winner at the 3rd Annual Data Science Workshop, Augusta University (03/17/2023)
- '*Outstanding Student of the Year*' award at Biostatistics & Data Science Division, Augusta University (2021-2022)
- '*Boyd Harshburger Travel Award*' winner at the Summer Research Conference, SRCOS (10/04/2022)
- Member of the *Augusta University Post-Doctoral Association*, Augusta University (Fall 2024 - Present)
- Member of the *Student Advisory Council*, Enrollment & Student Affairs, Augusta University (Fall 2024 - Present)
- Member of the *Graduate Student Council* (GSC), Augusta University (Fall 2021 - Summer 2023)
- GSC representative on the *The Graduate School - Faculty Council*, Augusta University (Fall 2022 - Summer 2023)

# Training / Certificates

**One year Master of Business Administration (MBA)**  
*National Institute of Business Management (NIBM)*

**2017**

Specialization: Market Research

**Business Analytics training course**  
*CONCEPTlytics Institute & Technologies*

**2017**