Samuel Anyaso-Samuel

CONTACT INFORMATION Biostatistics Branch Division of Cancer Epidemiology & Genetics

National Cancer Institute 9609 Medical Center Drive Rockville, MD 20850 Cell: (208) 216-9207 Office: (240) 276-5168

Email: samuel.anyaso-samuel@nih.gov Website: samuelanyaso.github.io

RESEARCH INTERESTS

Microbiome, Statistical genetics, Biological networks, Cancer Epidemiology, Cluster-correlated data analysis, Informative Cluster Size, Univariate and Multivariate time-to-event data.

EDUCATION & TRAINING

Postdoctoral Research Fellow

September 2023 - Present

Biostatistics Branch,

Division of Cancer Epidemiology & Genetics, National Cancer Institute, Rockville, MD, USA

University of Florida, Gainesville, FL, USA

Doctor of Philosophy (Ph.D.), Biostatistics

August 2023

- Thesis Topic: Advances in cluster-correlated data analysis when cluster size is informative
- Advisor: Somnath Datta, Ph.D.

Boise State University, Boise, ID, USA

Master of Science (M.S.), Mathematics

May 2019

- Thesis Topic: Dynamic Sampling Versions of Popular SPC charts for Big Data Analysis
- Advisor: Partha Mukherjee, Ph.D.
- Area of Study: Statistics

Federal University of Technology, Owerri, Nigeria

Bachelor of Technology (BTech.), Statistics

December 2014

- Thesis Topic: Some contributions to the interpretation of Fuzzy Regression Intervals
- Advisor: Benson Onoghojobi, Ph.D.

Professional Experience

Biostatistician

August 2020 - August 2023

Brain Rehabilitation Research Center,

NF/SG VHS Malcom Randall VA Medical Center, Gainesville, FL, USA

Graduate Research and Teaching Assistant

August 2019 - August 2023

Department of Biostatistics,

University of Florida, Gainesville, FL, USA

Graduate Teaching Assistant

August 2017 - May 2019

Department of Mathematics,

Boise State University, Boise, ID, USA

HONORS AND AWARDS

- Selected for Multi Omics NETwork Analysis Workship (MONET) University of Colorado Anschutz Medical Campus (June 2025)
- Blackwell-Tapia Conference Travel Award (Nov. 2024)
- Southern Regional Council on Statistics Summer Research Conference Travel Award (May 2024).
- 2023 Lifetime Data Science Conference Student Poster Award (June 2023).
- 2023 Symposium on Data Science & Statistics Student & Early Career Travel Award (May 2023).

- 2023 Fostering Diversity in Biostatistics (ENAR) Student Scholarship (March 2023).
- DEI Poster Award (Data Science); 2022 College of Medicine Research Day (April 2022).
- UF Department of Biostatistics PhD Travel Award (March 2022).
- 2020 Intelligent Systems for Molecular Biology Fellowship Award (July 2020).
- 2019 JSM Diversity Workshop and Mentoring Program Student Scholarship (July 2019).
- 36th ASA Quality & Productivity Research Conference Student Scholarship (June 2019).
- 2019 ASA/IMS Spring Research Conference Student Scholarship (May 2019).
- Best Poster; Workshop on Emerging Data Science Methods for Complex Biomedical and Cyber Data, Department of Population Health Sciences, Medical College of Georgia (March 2019).
- Best Poster from College of Engineering; Boise State University service-learning student exhibition, *Boise State University* (Dec. 2018).
- ACM Richard Tapia Celebration of Diversity in Computing Conference Scholarship (Sept. 2018).
- Student Representative, American Statistical Association, Boise State University (June 2018).
- Graduate Summer Fellowship, Department of Mathematics, Boise State University (May 2018).
- Alfred M. Dufty Jr. Award, Boise State University (May 2018).
- Computing Research Association (CRA) Sponsorship for CRA URMD workshop (March 2018).
- Graduate Residential Scholars Program, Boise State University (Aug. 2017).

PUBLICATIONS

- 9. **Anyaso-Samuel S.**, and Datta S. Nonparametric estimation of a future entry time distribution given the knowledge of a past state occupation in a progressive multistate model with current status data. doi.org/10.48550/arXiv.2405.05781 (2025+).
- 8. **Anyaso-Samuel S.**, Datta S., Roos E., and Nevalainen J. Can the unit size predict outcomes? Testing for informativeness in three-level designs. *Statistics in Medicine*, (In Print) (2025).
- 7. **Anyaso-Samuel S.**, and Datta S. Testing for marginal covariate effect when the subgroup size induced by the covariate is informative. *Statistical Methods in Medical Research*, 33(7):1264-1277, (2024).
- 6. Sarkar S., **Anyaso-Samuel S.**, Qiu P., and Datta S. Multiblock Partial Least Squares and Rank Aggregation: Applications to Detection of Bacteriophages Associated with Antimicrobial Resistance in the Presence of Potential Confounding Factors. *Statistics in Medicine*, 43(13):2527-2546 (2024).
- 5. Leinen M., Grandy E., Gebel L., Santana T., Rodriguez A., Singh S., Fernandez M., Dalugdug J., Garcia-Colon E., Lybeshari K., Alexander D., Maura M., Gonzalez M., Almeida C., Anyaso-Samuel S., Datta S., and Schiefer M. Bilateral Subdiaphragmatic Vagal Nerve Stimulation Using a Novel Waveform Decreases Body Weight, Food Consumption, Adiposity, and Activity in Obesity-Prone Rats. *Obesity Surgery*, 34(1):1-14, (2024).
- 4. **Anyaso-Samuel S.**, Bandyopadhyay D, and Datta S. Pseudo-value regression of clustered current status data with informative cluster or subcluster sizes in a multistate model. *Statistical Methods in Medical Research*, 32(8):1494-1510, (2023).
- 3. **Anyaso-Samuel S.**, and Datta S. Adjusting for informative cluster size in pseudo-value based regression approaches with clustered time-to-event data. *Statistics in Medicine*, 42(13): 2162-2178 (2023).
- 2. **Anyaso-Samuel S.**, Sachdeva A., Guha S., and Datta S. Bioinformatics preprocessing of microbiome data with an application to metagenomic forensics. In *Statistical Analysis of Microbiome Data*, (pp. 45-78), Eds: S. Datta and S. Guha, Springer (2021).
- 1. **Anyaso-Samuel S.**, Sachdeva A., Guha, S., and Datta S. Metagenomic geolocation prediction using an adaptive ensemble classifier. *Frontiers in Genetics*, *12*, p.642282 (2021).

OTHER ABSTRACTS

- 3. Ashby F., **Anyaso-Samuel S.**, Gamlin P., Kabbej N., Andraka N., Mandel R., Riva A., Datta S., Heldermon C. AAV-barcoding for High-throughput Screening of Vector Transduction Efficiency in the CNS of Cynomolgus Macaques Compared to C57BL/6 Mice. *Florida Genetics Symposium*, Gainesville, FL, (Nov. 2022).
- 2. Kabbej N., Ashby F.J., Riva A., **Anyaso-Samuel S.**, Datta S., Heldermon C.D. Transcriptomic Disparities Between Male and Female Non-Human Primates Related to AAV Transduction Efficiency. *American Society of Gene & Cell Therapy (ASGCT) 25th Annual Meeting*, Washington DC, (May 2022).
- 1. Ashby F., Kabbej N., Riva A., Rouse C.J., Hawkins K., Andraka N., **Anyaso-Samuel S.**, Gamlin P., Mandel R., Kondratov O., Zolotukhin S., Datta S., Heldermon C. Genetic Barcoding Identifies Similar Transduction Efficiency Rankings within Disease Models of Sanfilippo Syndrome Type-B and Controls. *19th Annual WORLDSymposium*, Orlando, FL, (Feb. 2022).

PRESENTATIONS

Invited, Refereed & Seminar Talks

- Identifying High-Dimensional Genomic Factor Associations in Biological Networks, *Department of Quantitative Health Sciences at the University of Hawai'i* (April 2025).
- Estimating marginal association in clustered data with informative subgroups induced by a given covariate, *The 7th International Conference on Econometrics and Statistics*, BEIJING, CHINA (July 2024) [virtual presentation].
- Regression analysis for clustered multistate current status data using the pseudo-value approach, 2023 Symposium on Data Science & Statistics, St. Louis, MO (May 2023).
- Regression analysis of clustered time-to-event data when the cluster size is informative: a pseudo-value approach, *Department of Epidemiology and Biostatistics, West Virginia University* (March 2023).
- Pseudo-value-based regression analysis of clustered multistate time-to-event data when the cluster size is informative, *Biostatistics Branch*, *National Cancer Institute* (March 2023).
- Regression analysis of clustered time-to-event data when the cluster size is informative, *Division of Computing, Analytics, and Mathematics, University of Missouri*, KANSAS CITY (February 2023).
- Regression analysis of clustered time-to-event data when the cluster size is informative, *UFSTAT Student Seminar Series*, GAINESVILLE, FL (February 2023).
- Bioinformatics Pre-processing of Microbiome Data with an Application to Metagenomics Forensics, 2021 *Joint Statistical Meetings*, VIRTUAL CONFERENCE (August 2021).
- Metagenomic Geolocation Prediction Using an Adaptive Ensemble Classifier, 28th Conference on Intelligent Systems for Molecular Biology, VIRTUAL CONFERENCE (July 2020).
- Fuzzy Regression Intervals, Graduate Student Seminar, Department of Mathematics, Boise State University, BOISE, ID (December 2017).

Contributed Talks

- Can the unit size predict outcomes? Testing for informativeness in three-level designs, 2025 ENAR Spring meeting, NEW ORLEANS, LA (March 2025).
- Nonparametric estimation of a future state occupation given the knowledge of a past state occupation in a multistate model with current status data, 2024 Joint Statistical Meetings, PORTLAND, OR (August 2024).
- Testing for Marginal Covariate Effect when the Subgroup Size Induced by the Covariate is Informative, 2024 ENAR Spring meeting, BALTIMORE, MD (March 2024).

- Regression analysis of multistate current status data with informative cluster sizes: a pseudo-value approach, *UF PHHP Research Day* 2023, GAINESVILLE, FL (February 2023).
- Adjusting for Informative Cluster Size in Pseudo-Value-Based Regression Approaches with Clustered Time-to-Event Data, 2022 *Joint Statistical Meetings*, WASHINGTON DC (August 2022).
- Pseudo-value based regression for clustered time-to-event data when cluster size is informative, *UF PHHP Research Day* 2022, VIRTUAL CONFERENCE (February 2022).
- Bioinformatics Pre-processing of Microbiome Data with an Application to Metagenomics Forensics, *UF PHHP Research Day* 2021, VIRTUAL CONFERENCE (February 2021)

Posters

- Testing for Marginal Covariate Effect When the Subgroup Size Induced by the Covariate is Informative, *Blackwell-Tapia Conference*, PROVIDENCE, RI (Nov. 2024).
- Testing for Marginal Covariate Effect When the Subgroup Size Induced by the Covariate is Informative, *The Southern Regional Council on Statistics 59th Summer Research Conference*, CLEMSON, SC (June 2024).
- Pseudo-Value Regression of Clustered Current Status Data with Informative Cluster or Subcluster Sizes in a Multistate Model, 2023 Lifetime Data Science Conference, RALEIGH, NC (May 2023).
- Pseudo-Value Regression of Clustered Current Status Data with Informative Cluster or Subcluster Sizes in a Multistate Model, 2023 Annual ASA Florida Chapter Meeting, GAINESVILLE, FL (March 2023).
- Pseudo-Value Regression of Clustered Current Status Data with Informative Cluster or Subcluster Sizes in a Multistate Model, 2023 ENAR Spring meeting, NASHVILLE, TN (March 2023).
- Pseudo-value based regression for clustered time-to-event data when cluster size is informative, 2022 International Chinese Statistical Association (ICSA) Applied Statistics Symposium, GAINESVILLE, FL (June 2022).
- Pseudo-value based regression for clustered time-to-event data when cluster size is informative, *UF College of Medicine Research Day* 2022, GAINESVILLE, FL (April 2022).
- EWMA Control Chart with a Dynamic Sampling Scheme, 2019 Quality and Productivity Research Conference, WASHINGTION D.C. (June 2019).
- Dynamic Sampling Versions of Popular SPC charts for Big Data Analysis, 2019 IMS/ASA Spring Research Conference, BLACKSBURG, VA. (May 2019).
- Statistical Process Control Charts for Monitoring Big Data Streams, Workshop on Emerging Data Science Methods for Complex Biomedical and Cyber Data, AUGUSTA, GA. (March 2019).
- Using Data Science to help Idaho cities make hiring decisions, *Boise State University Service-learning student exhibition*, BOISE, ID. (December 2018).
- Some contributions to the interpretation of Fuzzy Regression Intervals, *Computing Research Association URMD Workshop*, SAN DIEGO, CA. (March 2018).
- 3. **U.S. Department of Veterans Affairs IPA**, "Brain Rehabilitation Research Projects". September 2022 August 2023. Role: Principal Investigator (Mentor: Somnath Datta; VA PIs: D. Clark, R. M. Bauer).
- 2. **U.S. Department of Veterans Affairs IPA**, "Brain Rehabilitation Research Projects". September 2021 August 2022. Role: Principal Investigator (Mentor: Somnath Datta; VA PIs: D.

GRANTS

Clark, R. M. Bauer).

1. **U.S. Department of Veterans Affairs IPA**, "Brain Rehabilitation Research Projects". September 2020 - August 2021. Role: Principal Investigator (Mentor: Somnath Datta; VA PIs: D. Clark, R. M. Bauer).

SOFTWARE

- 5. **Anyaso-Samuel S.** and Datta S. **crspack** R package to conduct inference based on rank-sum statistics for cluster-correlated data with informativeness of the total cluster size, informativeness of a binary covariate distribution or informativeness of a subject-level covariate distribution.
- 4. **Anyaso-Samuel S.**, Bandyopadhyay D., and Datta S. **mspack2**. R package for estimating several temporal functions (e.g. state occupation probabilities) for current-status data from of a general multistate model. The code estimates the SOP for the setting where the current-status data is either uncorrelated or cluster-correlated.
- Anyaso-Samuel S. and Datta S. pseudoReg-ICS. R program for estimating the state occupation probability for cluster-correlated data from a multistate model. The program allows for adjusting for informative cluster size.
- 2. **Anyaso-Samuel S.**, Sachdeva A., Guha S., and Datta S. **metagenomic_data_analysis** Suite of programs for the bioinformatics pre-processing and downstream analysis of raw sequence metagenomics data.
- 1. **Anyaso-Samuel S.**, and Mukherjee P. **DyAEWMA** R package for estimating the average time to signal (ATS) of an adaptive EWMA chart with a dynamic sampling scheme or the average run length (ARL) of the adaptive EWMA chart.

TEACHING EXPERIENCE

Short courses

- Division of Cancer Epidemiology & Genetics, National Cancer Institute Feb. 2025 Co-instructor for the course "Multistate models for studying the natural history of cancer" delivered to staff and fellows within the division as part of a lecture series on statistical modeling in cancer epidemiology.
- Fostering Diversity in Biostatistics, *ENAR*Co-instructor for the course "Introduction to Statistical Analysis using R" delivered to high school and undergraduate students.

Department of Biostatistics, University of Florida

Instructor Fall 2022

• STA 6177 - Applied Survival Analysis.

Guest lecturer

Spring 2021, Spring 2022

- PHC 7066: Large Sample Theory.
- Gave lectures on *Modes of convergence* and *Asymptotic normality of the MLE* to PhD students.

Teaching Assistant

Fall 2019 - Summer 2023

- PHC 6937: Bayesian Biostatistical Methods
- PHC 6089: Pubic health computing
- PHC 6937: Fontiers in Biostatistics

- PHC 6937: Introduction to Applied Biostatistical Computing Using SAS
- PHC 6052: Introduction to Biostatistical Methods
- PHC 6937: Data Visualization in the Health Sciences
- PHC 6092: Introduction to Biostatistical Theory

Department of Mathematics, Boise State University

Instructor

Fall 2017, Fall 2018, Spring 2019

- MATH 149: Pre-Calculus.
- MATH 108: Intermediate Algebra.

Tutor Spring 2018

• Tutored students enrolled in *Intermediate Algebra*, *College Algebra* and *Pre-calculus* classes.

CONSULTING EXPERIENCE

Division of Hematology and Oncology, University of Florida

Aug. 2021 - Aug. 2023

Provided statistical support for a study funded by the NIH grant (1R01NS102624-01) titled "Optimizing AAV Vectors for Central Nervous System Transduction" (PI: Coy Heldermon).

- Designed a suitable pipeline for the bioinformatics pre-processing of raw sequenced reads.
- Performed statistical analyses for analyzing mRNA and DNA profiles from different animal models. Primary statistical analyses include nonparametric tests, rank aggregation, and correlation analyses.
- Utilized modern machine learning techniques from data visualization software to provide statistical graphics.

Departments of **Psychiatry** and **Clinical Psychology**, University of Florida **Jan. 2021 - Aug. 2023** Provided statistical support for various research projects by Dr. John Williamson, Dr. David Clark, and their trainees.

- Developed statistical models to analyze longitudinal and high-dimensional cross-sectional data sets. Primary statistical analyses involved mixed effects modeling and penalized regression modeling.
- Developed computer programs for data visualization.
- Conducted power & sample size analyses and wrote the statistical plan for two grant proposals funded by the NIH.

U.S. Department of Veterans Affairs, Gainesville, FL USA Aug. 2020 - Aug. 2023 Provided analytical support for the Brain Rehabilitation Research Center (BRRC) housed at the VA.

- Performed statistical analyses for more than 10 different projects and grant proposals. Primary statistical analyses involved power and sample size calculations, regression analysis, analysis of variance, multivariate analysis of variance, and correlation analysis.
- Wrote the statistical plan section for four grants submitted by BRRC members to the Department of Veterans Affairs.
- Constant communication with clinicians and physicians on statistical design and methodologies for ongoing projects.

PROFESSIONAL DEVELOPMENT

Preparing Future Faculty

Center for Teaching Excellence, University of Florida

Aug. 2022 - Dec. 2022

- Competitive and selective semester-long workshop focused on preparing participants for future careers in various academic settings.
- Devoted emphasis on evidence-based teaching, learning practices, expanding mentoring team, and strategies for being a successful faculty member.

SERVICE

Scholarly Journal Refereeing

Referee for Biometrics (3), Statistical Methods in Medical Research (3), Statistics in Medicine (3), Journal of Applied Statistics (2), Biometrical Journal (1), Lifetime Data Analysis (1).

Conference Session Organizer/Chair

- 2025 Joint Statistical Meetings Innovations in Biological Network Modeling: Unraveling Omics Data Analysis
- 2025 Lifetime Data Science Conference Beyond Right-censoring: Unveiling New Insights with Interval-censored Data Models
- 2025 Eastern North American Region, Spring meeting Innovative Strategies for Integrating Epidemiology and Biostatistics in Cervical Cancer Screening
- 2024 Joint Statistical Meetings Recent advances in the analysis of multistate survival models
- 2024 Eastern North American Region, Spring meeting Advances in Epidemiologic Methods
- 2023 Symposium on Data Science & Statistics Methods in Health & Medical Research
- 2023 Eastern North American Region, Spring meeting Clustered data methods

Conference & Seminar Service

- Poster Judge for 2025 NIH Graduate Student Research Symposium
- Planning committee member for 2025 ENAR Fostering Diversity in Biostatistics Workshop
- Poster Judge for 2025 ENAR Spring Meeting's Student Poster Competition

University Service (University of Florida)

- Member, Student recruitment committee; Department of Biostatistics (Nov. 2021 April 2023).
- Vice President, Biostatistics Students' Organization (Sept. 2021 Sept. 2022).
- President, Biostatistics Students' Organization (Sept. 2022 April 2023).

PROFESSIONAL MEMBERSHIPS

American Statistical Association (ASA)

2018 - Present

- Lifetime Data Analysis Section
- Biometrics Section
- Section on Statistics in Epidemiology

Royal Statistical Society (RSS)	2014 - 2023
Institute of Mathematical Statistics (IMS)	2014 - 2023
International Biometric Society, Eastern North American Region (ENAR)	2022 - 2023
International Chinese Statistical Association (ICSA)	2022 - 2023
International Society for Computational Biology (ISCB)	2020 - 2021
Mathematical Association of America (MAA)	2017 - 2019

COMPUTER SKILLS General Software

- *Operating systems.* WINDOWS, LINUX and MACOS.
- Productivity applications. Advanced skills in WORD, EXCEL, and POWERPOINT.

Computing & Programming

- Parallel Computing in selected scripting languages.
- Extensive experience with **R**/RStudio, **C++**, MATLAB, PYTHON/JUPYTER, UNIX, LATEX.
- Intermediate experience with SAS, STATA, SPSS.
- Version control: GitHub user @samuelanyaso

Bioinformatics

- Extensive experience in building pipelines for pre-processing and analysis of large-scale sequencing data.
- Downstream analysis of -omics data.

REFEREES Available upon request.