Samuel Anyaso-Samuel, Ph.D.

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

Work Authorization: Eligible to work in the U.S. without employer sponsorship

RESEARCH INTERESTS

Statistical Genomics, Integrative Analysis of Sequencing and Multi-Omics Data, Graphical Models, Cancer Epidemiology, Cluster-correlated data analysis, Informative Cluster Size, Univariate and Multivariate time-to-event data.

EDUCATION

University of Florida, Gainesville, FL, USA
Doctor of Philosophy (Ph.D.), Biostatistics   **Aug. 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   **Dec. 2014**

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

PROFESSIONAL EXPERIENCE

Postdoctoral Research Fellow   **Sept. 2023 - Present**
Biostatistics Branch
Division of Cancer Epidemiology & Genetics
National Cancer Institute, Bethesda, MD, USA

Biostatistician   **Aug. 2020 - Aug. 2023**
Brain Rehabilitation Research Center
NF/SG VHS Malcom Randall VA Medical Center, Gainesville, FL, USA

Graduate Research and Teaching Assistant   **Aug. 2019 - Aug. 2023**
Department of Biostatistics
University of Florida, Gainesville, FL, USA

Graduate Teaching Assistant   **Aug. 2017 - May 2019**
Department of Mathematics
Boise State University, Boise, ID, USA

HONORS AND AWARDS

- Selected for Multi Omics NETwork Analysis (MONET) Workshop at 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).
- Poster Award (Data Science); 2022 College of Medicine Research Day (Apr. 2022).
- UF Department of Biostatistics PhD Travel Award (Mar. 2022).
- 2020 Intelligent Systems for Molecular Biology - Fellowship Award (July 2020).
- 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* (Mar. 2019).
- Best Poster from College of Engineering; Boise State University service-learning student exhibition, *Boise State University* (Dec. 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 (Mar. 2018).
- Graduate Residential Scholars Program, *Boise State University* (Aug. 2017).

PUBLICATIONS

10. McElderry, J., Zhang, T., Zhao, W., Hoang, P., **Anyaso-Samuel, S.**, ..., Landi, M. Comprehensive microbiome analysis of 940 lung cancers in never-smokers. *Accepted at Nature Communications*.

9. Wei, Z., Zhang, T., ..., **Anyaso-Samuel, S.**, ..., Shi, J., Landi, M. A prognostic signature for lung adenocarcinoma in people who have never smoked. *Accepted at Cancer Discovery*.

8. **Anyaso-Samuel S.**, Datta S., Roos E., Nevalainen J. Can the unit size predict outcomes? Testing for informativeness in three-level designs. *Statistics in Medicine*, 44(6), p.e70041, (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., ..., **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).

IN REVISION/REVIEW/PREPARATION:

12. Abubakar, M., Shahin, S., ..., **Anyaso-Samuel, S.**, ..., Shi, J., Yang, X. Multi-platform spatial profiling reveals intra-tumor heterogeneity in immune and tumor markers in breast tumors from Kenyan patients. *Submitted to Cancer Research*.

11. Qin, F., Hua, X., ..., **Anyaso-Samuel, S.**, ..., Shi, J., Yu, K. Identification of immune cell type-specific susceptibility genes in multiple cancers using transcriptome-wide association studies. *Submitted to JNCI*.

10. Thakur, R., Xu, M., Thornock, A., Yon, J., **Anyaso-Samuel, S.**, ..., Shi, J., Brown, K. Functional characterization of the 9q34.13 melanoma risk locus identifies *RAPGEF1* as a melanoma risk and nevus gene linked to RAS activation. *Submitted to Cancer Discovery*.

9. **Anyaso-Samuel S.**, Albert, P., Shi, J. **MoSCNet**: a computational tool to uncover Genetic Modulators of Gene Expression Networks in Single-Cell RNA-Seq studies.

8. Thong, L., **Anyaso-Samuel, S.**, ..., Choi, J. Single-cell eQTL dataset of lung tissues from Asian never-smokers highlight the roles of alveolar epithelial cells in lung cancer etiology.

7. Ke, C., **Anyaso-Samuel S.**, Bandyopadhyay, D. Kernel-based Sufficient Dimension Reduction For Single-index Survival Model.

6. **Anyaso-Samuel, S.** and Ramos, M. Self-Reported Statistical Literacy and Conditional Willingness to Use Statistics in Everyday Decision-Making: Evidence from a National Survey of U.S. Adults. *Submitted to PlosOne*.

5. **Anyaso-Samuel S.**, Li, S., ..., Albert, P., Shi, J. Identifying High-Dimensional Genomic Factors Modulating Biological Networks Across Multi-Omic Data. *Submitted to Cell Genomics*.

4. Li, F., **Anyaso-Samuel, S.**, ..., Vogtmann, E. Association between diet quality and the oral microbiome in three US cohort studies. *Submitted to The American Journal of Clinical Nutrition*.

3. Bather, J., **Anyaso-Samuel, S.**, Chen, Y., Elliott, L., Bennett, A., and Goodman, M. Cluster-weighted modified Poisson regression for estimating risk ratios in longitudinal data with informative cluster sizes. doi.org/10.1101/2025.05.23.25328253 *Submitted to BMC Medical Research Methodology* (2025+).

2. **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 *Under revision at Lifetime Data Analysis* (2025+).

1. **Anyaso-Samuel S.**, Bandyopadhyay, D., Datta, S. **mspack**: An R package for nonparametric estimation of temporal functions in a progressive multistate model with current status data.

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).

<div style="display:flex">
<div style="width:20%">PRESENTATIONS</div>
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Invited Talks & Mini-symposiums

- From Graduate School to the Next Chapter: Navigating the NIH Postdoctoral Experience, *Department of Biostatistics and Data Science, University of Texas Health Science Center at Houston* (Jan. 2026).

- Uncovering high-dimensional genomic signals modulating biological networks via Gaussian graphical models, *19th International Joint Conference on Computational and Financial Econometrics and Computational and Methodological Statistics,* LONDON, U.K (Dec. 2025).

- Identifying high-dimensional genomic factors associated with biological networks, *Department of Biostatistics at Virginia Commonwealth University* (Aug. 2025).

- Inference for current status observations in a multi-state setting, *Department of Statistical Sciences and Operations Research at Virginia Commonwealth University* (Aug. 2025).

- Early career panel: Transition from graduate school to the workplace, *2025 Diversity Mentoring Workshop at the Joint Statistical Meetings,* NASHVILLE, TN (Aug. 2025).

- Nonparametric estimation of a future state occupation given the knowledge of a past state occupation in a multistate model with current status data, *2025 Lifetime Data Science Conference,* BROOKLYN, NY (May 2025).

- Identifying High-Dimensional Genomic Factor Associations in Biological Networks, *Department of Quantitative Health Sciences at the University of Hawai'i* (Apr. 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* (Mar. 2023).

- Pseudo-value-based regression analysis of clustered multistate time-to-event data when the cluster size is informative, *Biostatistics Branch, National Cancer Institute* (Mar. 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 (Feb. 2023).

- Regression analysis of clustered time-to-event data when the cluster size is informative, *UFSTAT Student Seminar Series*, GAINESVILLE, FL (Feb. 2023).

- Bioinformatics Pre-processing of Microbiome Data with an Application to Metagenomics Forensics, *2021 Joint Statistical Meetings*, VIRTUAL CONFERENCE (Aug. 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 (Dec. 2017).

Contributed Talks

- Identifying high-dimensional genomic factors associated with biological networks, *2025 Joint Statistical Meetings,* NASHVILLE, TN (Aug. 2025).

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- Can the unit size predict outcomes? Testing for informativeness in three-level designs, *2025 ENAR Spring meeting,* NEW ORLEANS, LA (Mar. 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 (Aug. 2024).

- Testing for Marginal Covariate Effect when the Subgroup Size Induced by the Covariate is Informative, *2024 ENAR Spring meeting,* BALTIMORE, MD (Mar. 2024).

- Regression analysis of multistate current status data with informative cluster sizes: a pseudo-value approach, *UF PHHP Research Day 2023*, GAINESVILLE, FL (Feb. 2023).

- Adjusting for Informative Cluster Size in Pseudo-Value-Based Regression Approaches with Clustered Time-to-Event Data, *2022 Joint Statistical Meetings*, WASHINGTON DC (Aug. 2022).

- Pseudo-value based regression for clustered time-to-event data when cluster size is informative, *UF PHHP Research Day 2022*, VIRTUAL CONFERENCE (Feb. 2022).

- Bioinformatics Pre-processing of Microbiome Data with an Application to Metagenomics Forensics, *UF PHHP Research Day 2021*, VIRTUAL CONFERENCE (Feb. 2021)

Posters
- Identifying high-dimensional genomic factors associated with biological networks, *NIH Research Festival,* BETHESDA, MD (Sept. 2025).

- 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 (Mar. 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 (Mar. 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 (Apr. 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. (Mar. 2019).

- Using Data Science to help Idaho cities make hiring decisions, *Boise State University Service-*

learning student exhibition, BOISE, ID. (Dec. 2018).

- Some contributions to the interpretation of Fuzzy Regression Intervals, *Computing Research Association URMD Workshop*, SAN DIEGO, CA. (Mar. 2018).

GRANTS

Brain Rehabilitation Research Projects   **Aug. 2020 - Aug. 2023**
Studies Agency: U.S. Department of Veterans Affairs (PI: D. Clark, R. M. Bauer)
Grant Type: IPA
Grant Role: Statistician

WISE II - Obesity and Type-2 Diabetes:
Bariatric Surgery Effects on Brain Function   **Nov. 2022 - Aug. 2023**
Studies Agency: NIH/NIDDK (PI: E. Porges)
Grant Type: R01 DK099334-06A1
Grant Role: Graduate Research Assistant

SOFTWARE

6. **Anyaso-Samuel S.**, Albert, P.S., Shi, J. **GFBioNet** is an R package that learns a network for biological traits via Gaussian graphical modeling and maps genomic factors modulating specific edges using node-wise regression. It reports *trait–trait–factor* triplets with robust false-discovery control, even under strong predictor correlation.

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.

3. **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 Excellence in Biostatistics**, *ENAR*   **Mar. 2025**
  Co-instructor for the course "Introduction to Statistical Analysis using R" delivered to high school and undergraduate students.

Regular courses

- **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.

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, Statistical Methods in Medical Research, Statistics in Medicine, Journal of Applied Statistics, Biometrical Journal, Lifetime Data Analysis, Sankhya B, BMC Medical Research Methodology, Pharmaceutical Statistics.*

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*
- Judge for *2025 ASA Lifetime Data Science Section Student Paper Awards*

Department Service

- *Fellows' representative*, Biostatistics Branch, DCEG, NCI (Aug. 2025 - Present)
- *Member*, Student recruitment committee; Department of Biostatistics, UF (Nov. 2021 - Apr. 2023).
- *Vice President*, Biostatistics Students' Organization, UF (Sept. 2021 - Sept. 2022).
- *President*, Biostatistics Students' Organization, UF (Sept. 2022 - Apr. 2023).

PROFESSIONAL MEMBERSHIPS	American Statistical Association
	Eastern North American Region, International Biometric Society

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.**