Yusuph Mavura, PhD, MS.

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Email: Yusuph.Mavura[at]ucsf.edu or yrmavura[at]gmail.com

Website: ymavura3.github.io

Areas of expertise: Causal inference using observational data, general epidemiology and study design, grant writing, technical writing, communication, exome/genome sequence data analysis, clinical/mendelian genetics, genetic epidemiology and complex disease genetics, population genetics, statistical genetics, RNASeq data analysis

Programming languages: Python, Bash Scripting, R, Stata, Perl.

Languages: English and Kiswahili

EDUCATION:

Ph.D., Epidemiology and Translational Science
 University of California, San Francisco
 Advisor: Neil Risch, PhD

• M.S., Bioinformatics

Georgia Institute of Technology, Atlanta

Dec 2018

B.S., Biochemistry and Molecular Biology
 Jomo Kenyatta University of Agriculture and Technology, Nairobi, Kenya

RESEARCH FUNDING:

 <u>Principal Investigator</u>: National Human Genome research institute (NHGRI) Predoctoral to Postdoctoral Transition Award for a Diverse Genomics Workforce (F99/K00)
 National Institutes of Health (NIH) Grant Title: "Assessing Clinical Utility of Polygenic Risk Scores in Ancestrally Diverse Real-World Cohorts"

Grant Number: F99HG013437

F99 Funding period: 01/01/2024 – 03/31/2025

K00 Funding period: Up to 3 years after starting postdoctoral research.

RESEARCH EXPERIENCE:

Graduate Research Assistant

Aug 2023 – Mar 2025

Department of Epidemiology and Biostatistics, University of California, San Francisco, CA *Advisor:* Dr. Akinyemi Oni-Orisan, PharmD, PhD

• Project lead in investigating the role of environmental, genetic factors, and dose-response in the effectiveness of antihypertensive drug therapy in individuals from ancestrally diverse cohort using AllofUS and Genetic Epidemiology research on Aging (GERA) data

Graduate Research Assistant/ Rotation Student

Aug 2020 – Dec 2020

Department of Epidemiology and Biostatistics,

University of California, San Francisco, CA

Advisor: Dr. John Witte, PhD, MS

• Investigated pleiotropy in cancer in a pan-cancer, cross-tissue transcriptome-wide association study (TWAS) using UKBiobank and Genetic Epidemiology research on Aging (GERA) cohort

Graduate Research Assistant

Mar 2020 - Dec 2024

Department of Epidemiology and Biostatistics, University of California, San Francisco, CA *Advisor:* Dr. Franklin Huang, MD, PhD

- Project lead in a study of the genomic rearrangement landscape of localized prostate cancer samples of Nigerian men of African descent
- Collaborated with researchers from Ahmadu Bello University, Zaria Nigeria

Graduate Research Assistant

Sep 2019 – Mar 2025

Department of Epidemiology and Biostatistics, and Institute of Human Genetics, University of California, San Francisco, CA *Advisor:* Dr. Neil Risch, PhD

- Project lead and Junior Principal investigator assessing Diagnostic Yield from Exome Sequencing
 in ancestrally diverse patients with suspected mendelian disorders in the P³EGS program, UCSF
- Junior Principal investigator and project lead in assessing Diagnostic Yield from Genome and Exome Sequencing in ancestrally diverse prenatal, neonatal intensive care unit (NICU), pediatric patients suspected mendelian disorders in various healthcare settings in the NHGRI Clinical Sequencing Evidence Generating research (CSER) Phase II program, involving centers and collaborators across the US
- Assessing association between genetic ancestry Diagnostic Yield from Exome Sequencing in ancestrally diverse newborns with suspected metabolic disorders in the Newborn Sequencing (NBSeqNext) program, UCSF
- Principal investigator, assessing Clinical Utility of Polygenic Risk Scores in Ancestrally Diverse Real-World Cohorts.

Intern Jun 2018 – Aug 2018

National Institutes of Health (NIH), Bethesda, MD

Graduate Summer Opportunity to Advance Research (G-SOAR)

Department of Cancer Epidemiology and Genetics (DCEG), National Cancer Institute (NCI)

Advisor: Dr. Laufey Amundadottir, PhD

• Conducted a Genome-Wide Association study (GWAS) which identified multiple new risk loci for pancreatic cancer in East Asian populations

Graduate Research Assistant

Aug 2017 – Dec 2018

School of Biological Sciences, Georgia Institute of technology, Atlanta, GA *Advisor:* Dr. John McDonald, PhD

• Analysis of mutations in cancer driver genes in matched primary, metastatic and recurrent ovarian cancer whole-exome sequencing samples (WES)

OTHER RESEARCH EXPERIENCE:

Mar 2016 - Jul 2017 Research Assistant

International Center of Insect Physiology and Ecology, Nairobi, Kenya Advisor: Dr. Daniel Masiga, PhD

- Primary data collection:
 - o Set traps and collected sandflies in Gilgil and Chemolingot, Kenya.
 - o Collected whole blood samples from cattle in Kwale, Kenya
- Molecular identification, genotyping, and characterization of Leishmania parasites and their sandfly vectors
- Screened for trypanosome parasite in whole blood samples from cattle
- Investigated the genetic diversity of trypanosome parasites, specifically *T. vivax* in cattle whole blood samples from Kwale, Kenya

Undergraduate Research Assistant

May 2015 – Dec 2015

Jomo Kenyatta University of Agriculture and Technology, Nairobi, Kenya Advisor: Dr. Daniel Kiboi, PhD

Investigated structural consequences of mutations in APIAP2 gene in Piperaquine and Lumefantrine resistant Plasmodium berghei Anka using 3D homology modeling

Research Intern Mar 2015 – May 2015

Department of Biology, Cell Biology, Konstanz, Germany

- Advisor: Dr. Christof Hauck, PhD
- Undertook intensive practical training course involving ligation independent cloning (LIC), bacterial transformation and antimicrobial selection, plasmid preparation, culture of human cell lines, sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), western blotting, light microscopy
- Investigated the effect of a SNP in the promoter of Protein Phosphatase (PPM1f) Gene on the binding of transcription factor Nkx2.2 in human cell lines

Research Intern Mar 2014 - May 2014

Government Chemists, Nairobi, Kenya

Screened for Narcotics and Aflatoxins in consumables using enzyme-linked immunosorbent assay (ELISA) and thin layer chromatography (TLC)

Research Intern Jan 2014 - Feb 2014

Africa Technical Research Center, Arusha, Tanzania Advisor: Dr. Johnson Ouma, PhD

Analyzed insecticide contents in treated mosquito nets for quality control using gas chromatography and UV spectrophotometry at the Analytical Chemistry lab

PUBLICATIONS: Refereed Journal Articles

Yusuph Mavura, Nuriye Sahin-Hodoglugil, Ugur Hodoglugil, Mark Kvale, Pierre-Marie Martin, Jessica Van Ziffle, W. Patrick Devine, Sara L. Ackerman, Barbara A Koenig, Pui-Yan Kwok, Mary E. Norton, Anne Slavotinek, Neil Risch. Genetic ancestry and diagnostic yield of exome sequencing in a diverse population. npj Genom. Med. 9, 1 (2024). https://doi.org/10.1038/s41525-023-00385-6

- Slavotinek A, Rego S, Sahin-Hodoglugil N, Kvale M, Lianoglou B, Yip T, Hoban H, Outram S, Anguiano B, Chen F, Michelson J, Cilio RM, Curry C, Gallagher RC, Gardner M, Kuperman R, Mendelsohn B, Sherr E, Shieh J, Strober J, Tam A, Tenney J, Weiss W, Whittle A, Chin G, Faubel A, Prasad H, Mavura Y, Van Ziffle J, Devine WP, Hodoglugil U, Martin PM, Sparks TN, Koenig B, Ackerman S, Risch N, Kwok PY, Norton ME. Diagnostic yield of pediatric and prenatal exome sequencing in a diverse population. NPJ Genom Med. 2023 May 26;8(1):10. doi: 10.1038/s41525-023-00353-0. PubMed PMID: 37236975; PubMed Central PMCID: PMC10220040.
- **Mavura Y**, Song H, Xie J, Tamayo P, Mohammed A, Lawal AT, Bello A, Ibrahim S, Faruk M, Huang FW. Transcriptomic profiling and genomic rearrangement landscape of Nigerian prostate cancer. Prostate. 2023 Jan 4;. doi: 10.1002/pros.24471. [Epub ahead of print] PubMed PMID: 36598071.
- Kamran SC, Xie J, Cheung ATM, **Mavura MY**, Song H, Palapattu EL, Madej J, Gusev A, Van Allen EM, Huang FW. Tumor Mutations Across Racial Groups in a Real-World Data Registry. JCO Precis Oncol. 2021 Nov;5:1654-1658. doi: 10.1200/PO.21.00340. PMID: 34994651.
- Oni-Orisan A, **Mavura Y**, Banda Y, Thornton TA, Sebro R. Embracing Genetic Diversity to Improve Black Health. N Engl J Med. 2021 Mar 25;384(12):1163-1167. doi: 10.1056/NEJMms2031080. Epub 2021 Feb 10. PMID: 33567186.

OTHER PUBLICATIONS: Non-Refereed Journal Articles and Book Chapters

Mavura MY, Huang FW. How Cancer Risk SNPs May Contribute to Prostate Cancer Disparities. Cancer Res. 2021 Jul 15;81(14):3764-3765. doi: 10.1158/0008-5472.CAN-21-1146. PMID: 34266915.

OTHER PUBLICATIONS: Manuscripts In Press, Under Review, or in progress

Yusuph Mavura, David Crosslin, Kathleen DM Ferar, John Greally, Lucia Hindorff, Gail P Jarvik, Sara Kalla, Barbara A Koenig, Mark Kvale, Pui-Yan Kwok, Mary Norton, Sharon E. Plon, Bradford C. Powell, Anne Slavotinek, Michelle L Thompson, Alice B Popejoy, Eimear E. Kenny, Neil Risch. Diagnostic Yield Results from the CSER 2 consortium: no evidence for differences by genetic ancestry. Under Review.

Beale, H. C., Tse, V., Lee, J. Y., Akutagawa, J., **Mavura, Y**., Saint-John, B., Cheney, A., Mulligan, D. R., Chacaltana, G., Gutierrez, M., Tenney, J., Shieh, J. T., Martin, P.-M., Yip, T., Hodoglugil, U., Fay, A. J., Brooks, A. N., Ziffle, J. V., Stone, M. D., ... Slavotinek, A. (2025). *A novel splice site variant in DEGS1 leads to aberrant splicing and loss of DEGS1 enzyme activity, a VUS resolved* (p. 2025.04.04.25325118). medRxiv. https://doi.org/10.1101/2025.04.04.25325118. Under Review.

ABSTRACTS:

Poster Presentations:

Yusuph Mavura, David Crosslin, Kathleen DM Ferar, John Greally, Lucia Hindorff, Gail P Jarvik, Sara Kalla, Barbara A Koenig, Mark Kvale, Pui-Yan Kwok, Mary Norton, Sharon E. Plon, Bradford C. Powell, Anne Slavotinek, Michelle L Thompson, Alice B Popejoy, Eimear E. Kenny, Neil Risch. Diagnostic yield of genome and exome sequencing in the ancestrally diverse CSER Phase II consortium is not associated with genetic ancestry in a variety of clinical settings. American Society of Human Genetics, Los Angeles, CA. October 2023.

Yusuph Mavura, Nuriye Sahin-Hodoglugil, Mark Kvale, Jessica Van Ziffle, W. Patrick Devine, Ugur Hodoglugil, Pierre-Marie Martin, Barbara A Koenig, Sara Ackerman, Anne Slavotinek, Pui-Yan Kwok, Mary E. Norton, Neil Risch. *No reduction in diagnostic yield of exome sequencing in prenatal and pediatric patients with non-European ancestries*. American Society of Human Genetics, Los Angeles, CA. October 2022.

PROFESSIONAL MEMBERSHIP:

American Society of Human Genetics.

2020- Present

PROFESSIONAL SERVICE:

American Society of Human Genetics career development committee.

Incoming Jan 2025

Manuscript Reviewer

American Journal of Epidemiology BMC Musculoskeletal Disorders

TEACHING EXPERIENCE:

Teaching Assistant Spring 2022

Course: Advanced Approaches to the Analysis of Observational Data (Biostats 215) Department of Epidemiology and Biostatistics, University of California San Francisco Course director: Tom Newman, MD, MPH

- Graded weekly assignments and tests for 14 MS and PhD students.
- Graded final exams for 14 MS and PhD students
- Led the review of previous week's assignments at the beginning of lectures
- Involved in formulating and editing the weekly and final exams questions
- Answered any questions or concerns from students

Teaching Assistant Winter 2021

Course: Molecular and Genetic Epidemiology (Epi 217)

Department of Epidemiology and Biostatistics, University of California San Francisco

Course director: John Witte, PhD, MS

- Graded weekly assignments and tests for 15 MS and PhD students
- Led the review of previous week's assignments at the beginning of lectures
- Guided students in their final project assignments
- Answered any questions or concerns from students

LEADERSHIP ROLES:

International Association for the Exchange of Students for Technical Experience (IAESTE) committee member.

Treasurer 2014-2015

Jomo Kenyatta University Students Governing Council Biochemistry Department Representative

2013-2014

REFERENCES:

Neil Risch, PhD

Lamond Family Foundation Distinguished Professorship in Human Genetics Professor, Department of Epidemiology and Biostatistics University of California San Francisco Email: Neil.Risch[at]ucsf.edu

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Maria Glymour, ScD Professor and Chair, Department of Epidemiology Boston University

Former Director of Epidemiology and Translational Science PhD Program at UCSF

Email: mglymour[at]bu.edu

Akinyemi Oni-Orisan, PharmD, PhD Associate Professor, Department of Clinical Pharmacy University of California San Francisco Email: Akinyemi.oni-orisan[at]ucsf.edu

Franklin Huang, MD, PhD
Associate Professor, Department of Medicine
University of California San Francisco
Email: Franklin.Huang[at]ucsf.edu