[Long CV]

# Mike Nsubuga

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Website: <a href="https://mikensubuga.github.io/">https://mikensubuga.github.io/</a>

## **EDUCATION**

2023 – 2027 University of Bristol, UK Ph.D.

Research: Using machine learning to trace gastrointestinal disease outbreaks and antimicrobial

resistance. Supported by MRC Ph.D. Studentship

Supervisors: Sion Bayliss, Lauren Cowley, Andrew Dowsey, and Kristen Reyher

2020 – 2022 Makerere University, Uganda M.Sc. in Bioinformatics (First Class Honors)

Dissertation Title: A machine learning approach to predict *E. coli* antibacterial resistance using whole-genome sequencing data in Uganda. Supported by **NIH-Fogarty EANBIT M.Sc. Fellowship** 

Supervisors: Daudi Jjingo and Gerald Mboowa

2016 – 2019 Makerere University, Uganda B.Sc. in Information Technology (First Class Honors)

#### RESEARCH EXPERIENCE

2023 – PRESENT **Jean Golding Institute of Data Science,** University of Bristol, UK

Ask-JGI Research Data Scientist

2023 – 2023 **Division of Human Genetics,** University of Cape Town, South Africa

Research Fellow, Sickle in Africa Consortium

2022 – 2023 Department of Entomology, Uganda Virus Research Institute, Uganda

Research Fellow, Open Data Science Platform - eLwazi

2020 – 2023 Infectious Diseases Institute, African Center of Excellence in

**Bioinformatics, Makerere University, Uganda** Research Fellow & Bioinformatics Trainee

#### **AWARDS, HONORS AND FELLOWSHIPS**

2024 – Travel Award: Awarded a travel grant by the Medical Research Council (MRC) to attend the 2024

Computational Genomics Summer Institute at University of California, Los Angeles

2023 – Ph.D. Studentship: Awarded a studentship from the Medical Research Council (MRC) to undertake

a 4-year Ph.D. at University of Bristol.

2022 – Travel Award: Awarded a travel scholarship by Open Science Grid funded by National Science

Foundation (NSF) to attend OSG school at the University of Wisconsin-Madison.

2021 – M.Sc. Fellowship: Awarded a Fellowship under EANBIT project from the National Institutes of Health

(NIH) Fogarty to undertake a 2-year Master's degree program in Bioinformatics at

Makerere.

2020 - Dissertation Award: Graduated top of undergraduate class, 2019

#### **PUBLICATIONS**

#### **Published**

2025

- Gregorova, M., Santopaolo, M., Garner, L. C., Hayati, R. F., Diamond, D., Ramamurthy, N., Tran, V. T., Nguyen, N. M., Heesom, K. J., Nguyen, V. L., Jones, E., **Nsubuga, M**., Luscombe, C., Vo, H. T. M., Ho, C. Q., Nguyen, C. T. X., Dong, T. T. H., Huynh, D. T. L., Cao, T. T., ... Rivino, L. (2025). Early NK-cell and T-cell dysfunction marks progression to severe dengue in patients with obesity and healthy weight. Nature Communications, 16(1), 5569. <a href="https://doi.org/10.1038/s41467-025-60941-9">https://doi.org/10.1038/s41467-025-60941-9</a>
- Please, H., **Nsubuga, M.,** Kintu, T. M., Bakesiga, A., Stewart, K., & Navarro, S. (2025). 462 Powering Precision Medical Education Through Al-Generated Country-Specific Global Surgery Podcasts: A Low-or-Middle-Income-Country Case Study. *British Journal of Surgery*, https://doi.org/10.1093/bjs/znaf128.052
- Kintu, T., **Nsubuga, M.,** Bakesiga, A., Please, H., Stewart, K., & Navarro, S. M. (2025). 369 Precision education and generative AI in surgery utilization study: A framework for global surgical education. *Journal of Clinical and Translational Science*, 9(s1), 114–114. https://doi.org/10.1017/cts.2024.994
- Bolton, W., Nathani, P., Please, H., Philomen, J., Kahar, N. A., Aruparayil, N., Bandyopadhyay, S., Boodhoo, V., Magoha, **M., Nsubuga**, M., Mishra, A., Jani, P., Burke, J., Mathew, R., & Culmer, P. (2025). Accelerating innovation by integrating artificial intelligence into a global surgery hackathon. *International Health Trends and Perspectives,* 5(1), Article 1. <a href="https://doi.org/10.32920/ihtp.v5i1.2332">https://doi.org/10.32920/ihtp.v5i1.2332</a>
- **Nsubuga, M.,** Kintu, T. M., Please, H., Stewart, K., & Navarro, S. M. (2025). Enhancing trauma triage in low-resource settings using machine learning: A performance comparison with the Kampala Trauma Score. *BMC Emergency Medicine*, 25(1), 14. https://doi.org/10.1186/s12873-025-01175-2
- Nabisubi, P., Kanyerezi, S., Kebirungi, G., Sserwadda, I., **Nsubuga, M.,** Kisitu, G., Nahirya, P. N., Mulindwa, B., Akabwai, G. P., Nantongo, S., Kekitiinwa, A., Kigozi, E., Luutu, N. M., Katabazi, F. A., Kalema, L., Katabalwa, A., Jjingo, D., & Mboowa, G. (2025). Beyond the fever: Shotgun metagenomic sequencing of stool unveils pathogenic players in HIV-infected children with non-malarial febrile illness. *BMC Infectious Diseases*, 25(1), 96. https://doi.org/10.1186/s12879-025-10517-1

2024

- **Nsubuga, M.,** Galiwango, R., Jjingo, D., & Mboowa, G. (2024). Generalizability of machine learning in predicting antimicrobial resistance in *E. coli*: A multi-country case study in Africa. *BMC Genomics*, *25*(1), 287. https://doi.org/10.1186/s12864-024-10214-4
- Babirye, S. R., **Nsubuga, M**., Mboowa, G., Batte, C., Galiwango, R., & Kateete, D. P. (2024). Machine learning-based prediction of antibiotic resistance in Mycobacterium tuberculosis clinical isolates from Uganda. BMC Infectious Diseases, 24(1), 1391. <a href="https://doi.org/10.1186/s12879-024-10282-7">https://doi.org/10.1186/s12879-024-10282-7</a>
- **Nsubuga, M**., Mutegeki, H., Jjingo, D. *et al.* The Ugandan sickle Pan-African research consortium registry: design, development, and lessons. *BMC Med Inform Decis Mak* **24**, 212 (2024). <a href="https://doi.org/10.1186/s12911-024-02618-9">https://doi.org/10.1186/s12911-024-02618-9</a>
- Gregorova, M., Santopaolo, M., Garner, L. C., Diamond, D., Ramamurthy, N., Vi, T. T., Nguyen, N. M., Vuong, N. L., Jones, E., **Nsubuga, M.,** Luscombe, C., My, H. V. T., Chanh, H. Q., Chau, N. T. X., Tam, D. T. H., Le, D. H. T., Tam, C. T., Klenerman, P., Yacoub, S., & Rivino, L. (2024). Early NK-cell and T-cell dysfunction marks progression to severe dengue in patients with obesity and healthy weight (p. 2024.09.06.611687). bioRxiv. https://doi.org/10.1101/2024.09.06.611687

- Mboowa, G., Kakooza, F., Egesa, M., Tukwasibwe, S., Kanyerezi, S., Sserwadda, I., Kidenya, B. R., Kabahita, J. M., Namaganda, M. M., **Nsubuga, M.,** Nabisubi, P., Ayitewala, A., Kebirungi, G., Nakafu, E., & Akwii, N. P. (2024). The rise of pathogen genomics in Africa (13:468). *F1000Research*. https://doi.org/10.12688/f1000research.147114.1
- Please, H., Narang, K., Bolton, W., **Nsubuga, M.,** Luweesi, H., Richards, N. B., Dalton, J., Tendo, C., Khan, M., Jjingo, D., Bhutta, M. F., Petrakaki, D., & Dhanda, J. (2024). Virtual reality technology for surgical learning: Qualitative outcomes of the first virtual reality training course for emergency and essential surgery delivered by a UK–Uganda partnership. *BMJ Open Quality*, *13*(1). <a href="https://doi.org/10.1136/bmjoq-2023-002477">https://doi.org/10.1136/bmjoq-2023-002477</a>

2022

Buyego, P., Katwesigye, E., Kebirungi, G., **Nsubuga, M.,** Nakyejwe, S., Cruz, P., McCarthy, M. C., Hurt, D., Kambugu, A., Arinaitwe, J. W., Ssekabira, U., & Jjingo, D. (2022). Feasibility of virtual reality based training for optimising COVID-19 case handling in Uganda. *BMC Medical Education*, 22(1), 274. <a href="https://doi.org/10.1186/s12909-022-03294-x">https://doi.org/10.1186/s12909-022-03294-x</a>

## **Accepted**

Lujumba et al., (2025) A practical guide to identifying associations between tandem repeats and complex human traits using consensus genotypes from multiple tools. *Nature Protocols* 

#### **Other Scholarly Works (Posters)**

- **Nsubuga M,** Yi Ling Tam, Malaka de Silva, James Hall, Lauren Cowley, Claire Jenkins, Kate Baker, Sion Bayliss. (2024). Mapping the distribution of AMR in *Shigella sonnei*. NIHR HPRU in Gastrointestinal Infections Conference, Birmingham, UK [Poster]. <a href="http://hprugi.nihr.ac.uk/media/f1uc4ayb/mike-nsubuga.pdf">http://hprugi.nihr.ac.uk/media/f1uc4ayb/mike-nsubuga.pdf</a>
- **Nsubuga, M**. (2023). A machine learning approach to predict E. coli antibacterial resistance using whole-genome sequencing data [Thesis, Makerere University]. <a href="http://makir.mak.ac.ug/handle/10570/13162">http://makir.mak.ac.ug/handle/10570/13162</a>
- **Nsubuga M,** Kristen Reyher, Andrew Dowsey, Lauren Cowley, Marie A. Chattaway, Sion Bayliss. (2025). Characterising regionally associated AMR lineages from a decade of genomic surveillance in *Salmonella* enterica serovar Enteritidis. I3S Congress(International Symposium on Salmonella and Salmonellosis), Saint-Malo, France [Poster].

#### **INVITED TALKS**

04/03/2022 -

<b>Upcoming Keynotes</b>	
17 - 21/04/2026 —	Is AI the key to acceleration of Antimicrobial Stewardship in LMICs? ESCMID Global,
	Germany
12/11/2025 -	What can we expect from AI to better survey and control AMR particularly in LMICs,
	AMR, a one health challenge course by Fondation Mérieux, France
Completed	
30/05/2025 –	Utilizing Machine Learning for tracing gastrointestinal outbreaks, UKHSA PhD Student
	Day, Harwell Science & Innovation Campus, UK
13/04/2024 -	Combatting AMR using AI in low resource settings, AMR, a one health challenge course
	by Fondation Mérieux, France
20/03/2024 -	Utilising Machine Learning for tracing gastrointestinal outbreaks and antimicrobial
	resistance, MRC GW4 DTP, UK
23/08/2023 -	Empowering low middle income countries against antimicrobial resistance with AI and
	Whole-Genome Sequencing, AMR Force, University of Bristol
23/07/2022 -	Ugandan AI COVID-19 chatbot for automated and personalized symptom assessment in
	Luganda & English. University of Wisconsin-Madison, Open Science Grid School, US

End to end AI and data systems for targeted surveillance and management of COVID-19

and future pandemics affecting Uganda (COAST), Uganda

08/04/2022 — Optimizing the SickleInAfrica Registry Data Collection Workflow Using Site-specific

Clinical Processes, Sickle in Africa consortium meeting, South Africa

26/04/2022 – Virtual Reality in Medicine and Surgery Conference (VRiMs), Uganda

#### **POSTER PRESENTATIONS**

2025 – Characterising regionally associated AMR lineages from a decade of genomic

surveillance in Salmonella enterica serovar Enteritidis Genome Science, Newcastle, UK

2025 – Characterising regionally associated AMR lineages from a decade of genomic

surveillance in Salmonella enterica serovar Enteritidis I3S Congress (International

Symposium on Salmonella and Salmonellosis), Saint-Malo, France

2024 – Mapping the distribution of AMR in Shigella sonnei, NIHR HPRU in Gastrointestinal

Infections Conference, Birmingham, UK

2024 – Mapping the distribution of AMR in Shigella sonnei, NIHR HPRU genomics and Enabling

Data Conference, London, UK

## **TEACHING EXPERIENCE**

2024 – School of Engineering Mathematics & Technology, University of Bristol

Teaching Assistant - EMATM0047, Data Science Project

Teaching Assistant - SEMT20003, Methods of Artificial Intelligence

Teaching Assistant - EMATM0048, Software Devt: Programming & Algorithms

2021 – 2022 H3ABioNet and Wellcome Connecting Science

Teaching Assistant – Next Generation Sequencing short course

2018 – 2019 **Department of Computer Science,** Makerere University

C & Java programming tutorial assistant

#### **WORK EXPERIENCE**

2020 – 2022 Supporting African Math Initiatives (SAMI), U.K

Software Developer (Volunteer to SAMI Math Charity)

2020 – 2020 Statistics for Sustainable Development (STATS4SD), Reading, UK

**Programming Intern** 

#### **CO-SUPERVISION**

Juliet Nabateesa, MSc in Bioinformatics, Makerere University, 2023 – 2026 Mansi Chandra, MSc in Bioinformatics, University of Bristol, 2024 – 2025

Florence Nakabiri, MSc in Bioinformatics, Makerere University, 2022 – 2024

### **MENTORSHIP**

Rebecca Nakitandwe, MSc in Health Informatics, Makerere University, 2023 – 2025 Henry Mutegeki, MSc in Computer Science (Google DeepMind Scholar), Makerere University 2022 – 2024 Sandra Babirye, MSc in Bioinformatics (MakDarta Fellow), Makerere University, 2022 – 2024

## SERVICE, OUTREACH AND LEADERSHIP

Work done under the DataFace, a UK schools-based programme designed by the Cheltenham Festivals that empowers students to explore real-world issues through the lens of data.

2024 - 2025 DataFace Programme - Cheltenham Festivals

Participated in the creation of training video content for the DataFace programme, a UK-wide schools initiative run by Cheltenham Festivals that empowers students to explore real-world issues through data. Featured in video segments designed to build their confidence in understanding, interpreting and communicating effectively through visual storytelling. Also served as a judge at the programme's final competition -

https://www.cheltenhamfestivals.org/our-projects/dataface

Work done under Supporting African Maths Initiatives (SAMI), a UK-based charity dedicated to improving access to and quality of mathematics education in Africa.

2020 – 2020 Software Developer

SAMI Math Club App (A collection of mathematical problems and puzzles to support mathematical thinking, problem solving and love of mathematics, used in 5 African

countries during Maths Camp) - <a href="https://mathsclub.samicharity.co.uk/">https://mathsclub.samicharity.co.uk/</a>

2020 – 2021 Lead Software Developer

Card Deck – A web and mobile application together with a printed card deck featuring unique activities, engaging participants in games, puzzles or fun mathematical facts –

https://cards.virtualmathscamp.com/

2020 – 2021 Junior Developer

UNICEF Parenting for Lifelong Health – An open source and evidence-based parenting app to support parents and caregivers during the COVID-19 pandemic and beyond

Work done under PICSA, a method developed by researchers at University of Reading, UK that combines climate data with African farmers' insights to enhance informed agricultural decision-making through participatory methods.

2022 – 2023 **Technical Lead** 

Conducted field digital training and workshops of approaches for farmers in Malawi and Zambia on implementation of the PICSA approach - <a href="https://picsa.app/">https://picsa.app/</a>

## **OTHER PREVIOUS PROJECTS**

2023 – 2023 Web Developer

Al for Health Equity: Transforming Pandemic Preparedness in Uganda (HEAL) using

Large Language Models- Funded by Bill & Melinda Gates Foundation -

https://heal.aceuganda.org/

2021 – 2023 Lead Software Developer

An Epilepsy Self-Management and Resilience Technical application for Adolescents and

their community – Funded by Epilepsy Foundation in collaboration with Duke University

https://github.com/aceuganda/epilepsy-smart-app

2021 – 2022 Junior Data Scientist

Al-based COVID-19 chatbot for Uganda – Funded by International Development Research Centre (IDRC)

Lead Developer

Sickle Pan- African Research Consortium (SPARCO) Uganda: Strengthening Capacity for Clinical Care, Research and Training in Sickle Cell Disease—Funded by NIH/NHLBI in collaboration with UCT

2021 - 2025