Day 1 Session 2: Practical approaches - specimen, genomics and data infrastructure

Aquillah Kanzi, Kareemah Suleiman, Shavanthi Rajatileka, Gerald Mboowa, Dr Dawit Wolday





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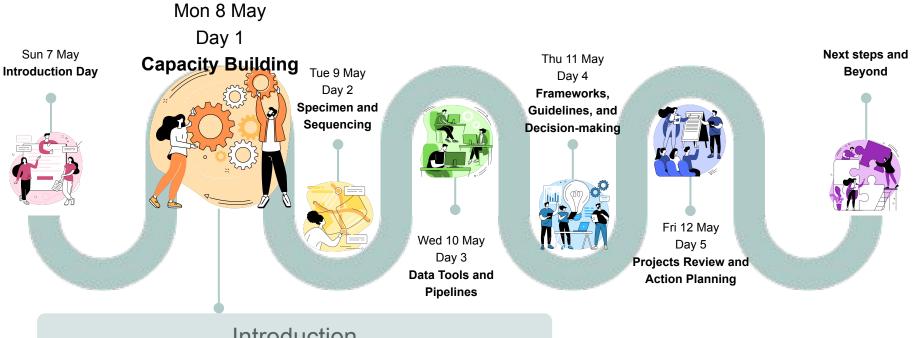
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GLOBAL TRAINING

Course roadmap



Introduction Overview of basic concepts Case studies

Session objectives

Showcase case studies about setting up infrastructure and processes - completed projects on a piece of infrastructure that has been set up for pathogen genomics Panel showcase - 5 case studies - 8 minutes each

- Biobank at IHV Kareemah
- Sequence infrastructure set up at a specific place or project network -Shavanthi
- Data infrastructure for surveillance (or network) (Africa CDC) Gerald
- Genomics infrastructure and pipelines for routine diagnostics (ASLM) -Aquillah



Institute of Human Virology Nigeria (IHVN) Biobank Kareemah Suleiman





The Biobank : Institute of Human Virology Nigeria H3Africa Biorepository (I-HAB)

Goal:

To promote population and personal Health, by facilitating cutting edge research and collaborations among African communities and beyond by providing high quality, affordable

biobanking services.





Current status of I-HAB

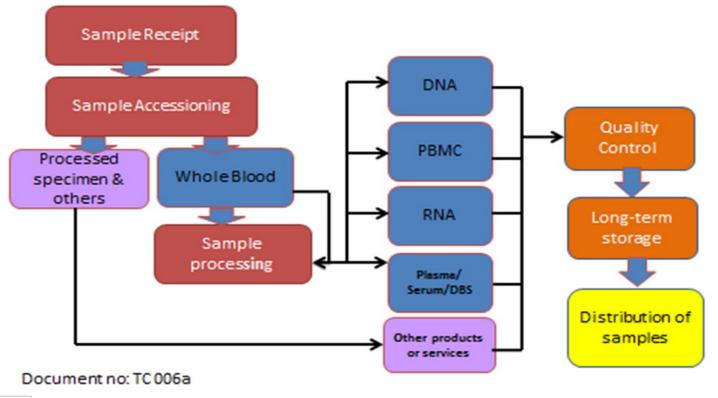
- 2009: IHVN with University of Maryland Baltimore established a biorepository network in Abuja, Zaria and Jos to support IHVN clinical and research activities.
- 2012:Abuja biorepository received funding through the H3 Africa grant supported by NIH to support African investigators in genomic research. The Abuja Biorepository became I-HAB
- 2013:I-HAB partnered with Coriell to bring the biorepository to international standards.
- 2015:H₃Africa biospecimen shipment pilot study between H₃Africa biorepository network
- Sample deposit began from 4 west African Countries-Nigeria, Benin, Ghana and Mali
 - Total biospecimen deposited at I-HAB- 32,687
- Trained 176 research staff in over 30 topics related to biobanking





How the IHVN Biobank is set up

Establishing Capacity for Pathogen Genomics Addis Ababa, Ethiopia, May 2023







Congoing initiatives and future plans for IHV Biobank Addis Ababa, Ethiopia, May 2023





Sequencing set-up Dr Shavanthi Rajatileka



National Institute for Health and Care Research







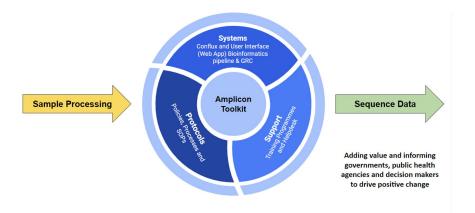
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GENOMIC SURVEILLANCE OF MALARIA IN WEST AFRICA

PROJECT OVERVIEW

- Funded by the National Institute for Health Research
- Project based at the West African Centre for Cell Biology of Infectious Pathogens (Ghana) and MRC Unit, The Gambia (The Gambia) in collaboration with the Wellcome Sanger Institute (UK).



PROJECT AIMS

- To build infrastructure (laboratory & some data systems) for genomic surveillance using amplicon sequencing in at the University of Ghana and MRC Unit, The Gambia.
- To establish a working proof of concept for genomic surveillance of malaria parasites and vectors in Ghana and The Gambia.
- To work with National Malaria Control Programmes (NMCPs) to learn to translate genomic data into actionable outputs that can be integrated into their operations.



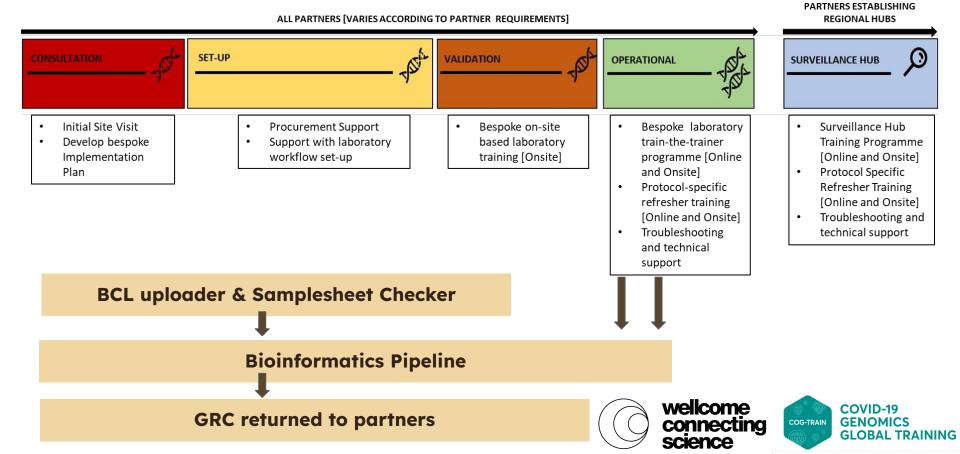
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DEPLOYMENT AND IMPLEMENTATION STRUCTURE

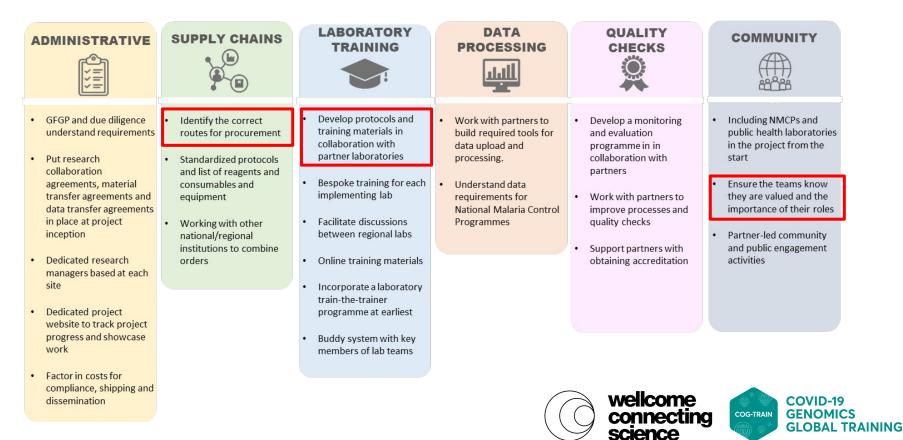


DEPLOYMENT AND IMPLEMENTATION STRUCTURE

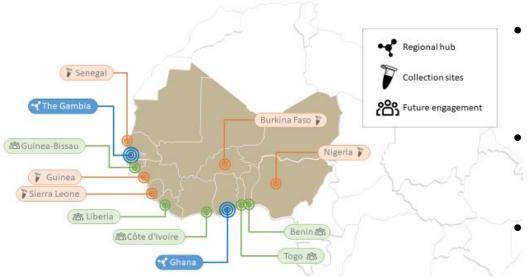
Pillar	Consultation	Laboratory Set-up	Sample Collection	Laboratory Protocol validation	Sequence Generation (Operational)	Genomic analysis	Dissemination
Major activities	 Site visits Resource review Develop bespoke implementation plan 	 SOP review Establish workflow Procurement of equipment, reagents, consumables Compliance, collaboration agreements, MTAs, DTAs 	 Collect sample Collect metadata Microscopy confirmation of species Laboratory confirmation of species 	 Optimise protocols Process test samples Sequence test samples Validation samples processed Laboratory Training Protocol QC checks 	 Process samples Sequence samples Refresher training Laboratory train-the- trainer programme Scaling up operations and improve QC 	 Bioinformatics and computational analysis Metadata linkage Generate genetic report card Visualize data 	 Report information to NMCPs/ Public Health officials/Policy makers
Implementing partner	Laboratory team Project leads Administration staff (Finance, Grants, Legal, Procurement)	Laboratory team Project leads Administration staff (Finance, Grants, Legal, Procurement)	Fieldworkers National malaria control programmes Clinic staff Laboratory staff	Laboratory staff Procurement teams	Laboratory staff Procurement teams	Bioinformatics team Data Scientists	Project PI Project Leads NMCPs Public Health Officials Policy Makers



LESSONS LEARNED



NEXT STEPS: PHASE 2



OBJECTIVES

- Expand capacity for genomic surveillance of malaria parasites and mosquito vectors in Africa and Asia, and to generate essential information required for National Malaria Control Programmes (NMCPs) to plan sustainable interventions
- **Establish regional sequencing hubs** capable of processing samples from neighbouring countries and providing NMCPs with timely, actionable genomic surveillance data
- Integrate genomic data into the routine working practices of NMCPs and to provide a working example of how such end to end genomic surveillance systems could be deployed at other locations in the world.



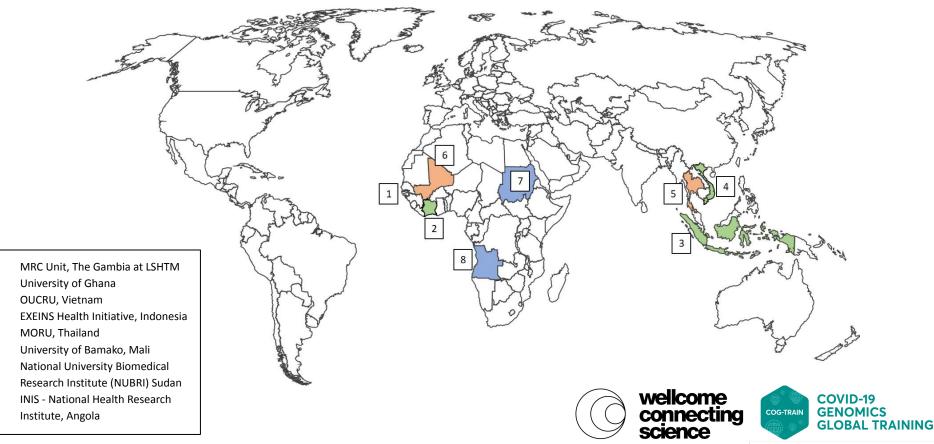
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NEXT STEPS: EXPANSION OF PARTNER SUPPORT



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AFRICA PGI DATA MANAGEMENT & EXCHANGE PLATFORM

Gerald Mboowa

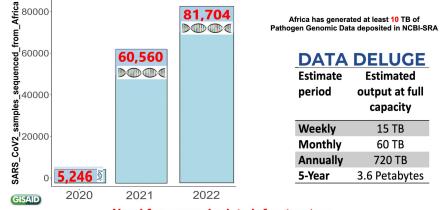


HOW THE DATA MANAGEMENT & EXCHANGE PLATFORM WAS SET UP

Data portal for surveillance of pathogens & antimicrobial resistance in near real-time

Goal of the project to develop infrastructure

- Africa PGI has equipped over 40 national public health institutions (NPHIs) with NGS platforms
- Many NPHIs lack capacity to analyse sequence data
- Federated data analysis, management, sharing and archiving



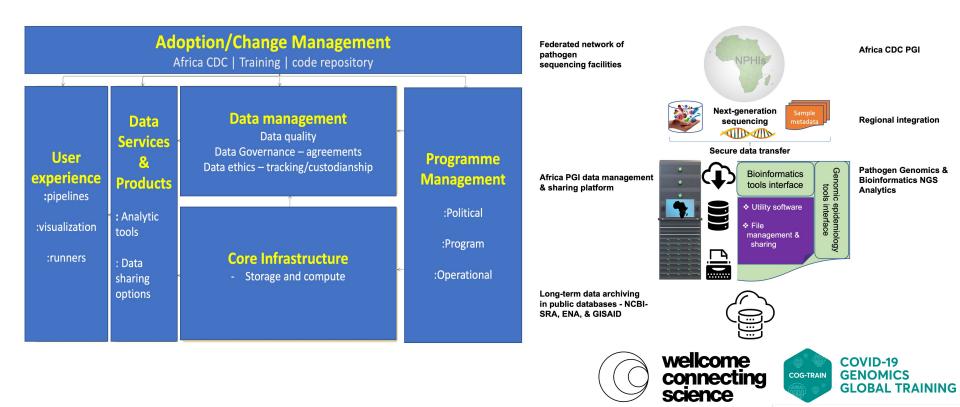
Need for genomic data infrastructure



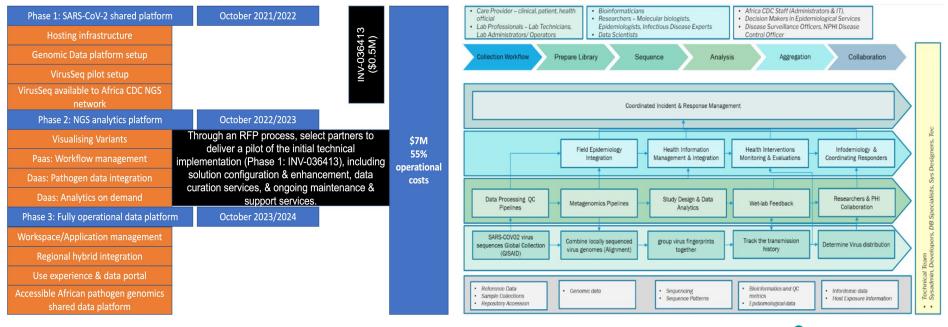
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HOW THE DATA INFRASTRUCTURE WAS SET UP



CURRENT STATUS OF THE DATA INFRASTRUCTURE CAPACITY PROJECT





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ONGOING INITIATIVES FOR DATA INFRASTRUCTURE PROJECT

- Pathogen genomics use-cases projects
 - 1. Vibrio cholerae genomic surveillance

AFRICA CDC

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Safeguarding Africa's Health

- 2. Klebsiella pneumoniae genomic surveillance
- 3. Respiratory pathogen panel (RPIP) pilot project
- Developing national multi-pathogen genomic surveillance strategies



TECHNICAL SUPPORT FOR DEVELOPMENT OF NATIONAL MULTI-PATHOGEN GENOMIC SURVEILLANCE STRATEGY



ADDIS ABABA, ETHIOPIA 2-3 DECEMBER, 2022



WWW.AFRICACDC.ORG

Genomics infrastructure and pipelines for routine diagnostics Dr Aquillah Kanzi





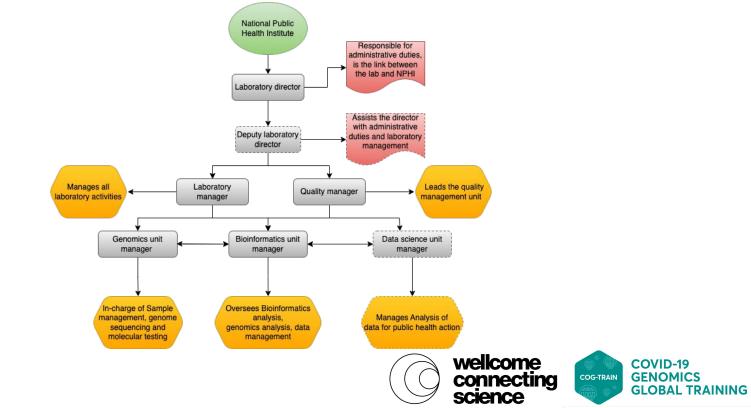
Setting up genomics infrastructure, quality management and capacity development: CARES Pathogen genomics

- ASLM CARES Pathogen Genomics project is funded by the US CDC CARES laboratory strengthening program
- Supports Africa CDC Africa PGI activities through the following:
 - Equipment and infrastructure upgrades
 - Sample collection and referral
 - Staffing and workforce development
 - Quality assurance quality management systems (QMS) and EQA
- CARES Pathogen Genomics supports nine African member states in the Africa PGI laboratory network



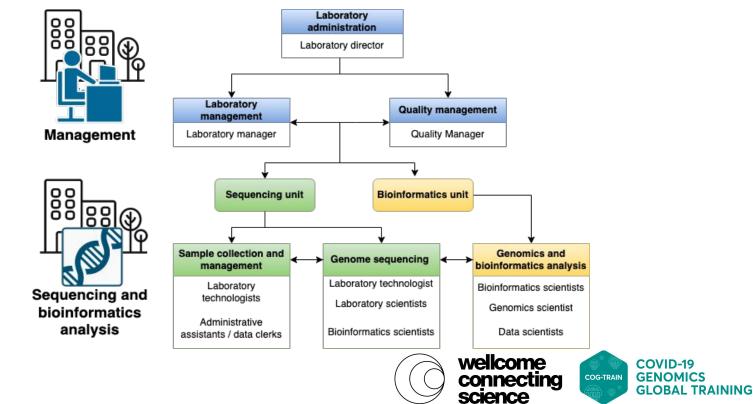


General organization of CARES supported genomics laboratories



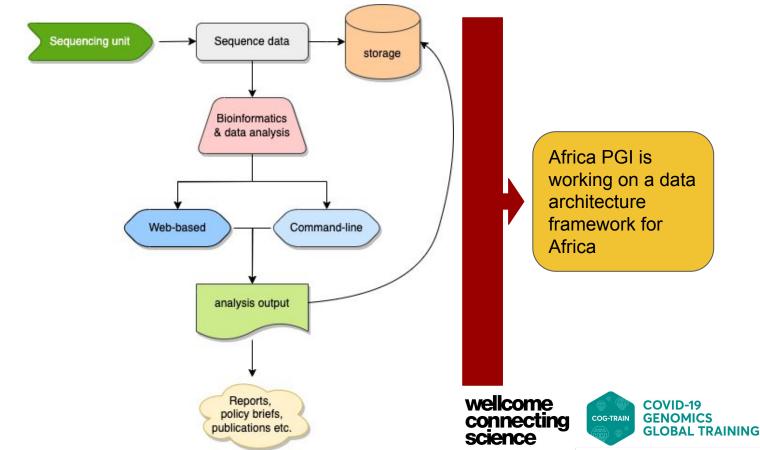


General organization of CARES supported genomics laboratories



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Implementation of bioinformatics analysis workflows





Current status of the genomics infrastructure capacity of CARES supported laboratories

Genome sequencing

- The laboratories have varying sequencing capacity depending on scope and needs
- Examples of sequencing equipment available in these laboratories include
 - Miseq
 - MiniSeq
 - Nextseq 2000
 - Oxford Nanopore MinION, GridION
 - Accessory equipment e.g, liquid handlers, DNA quantitators and quality analyzers, DNA size selection equipment, DNA shearing equipment etc.

Bioinformatics analysis

- Varying capacity of bioinformatics analysis infrastructure including,
 - desktop computers, laptops, and servers





Ongoing initiatives and future plans for strengthening capacity for CARES supported laboratories

Genome sequencing – SARS-CoV-2, other priority pathogens for surveillance and outbreak investigation

Training staff on genome sequencing and bioinformatics analysis of pathogens other than SARS-CoV-2

Strengthening quality through e.g., EQA participation and implementation of QMS

Establishment of regional genomics and EQA programme

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Establishing Pathogen Genomics Excellence at Ethiopian Public Health Institute Dr Dawit Wolday



EPHI Pathogen Genomics Initiative

<u>Overall aim:</u>

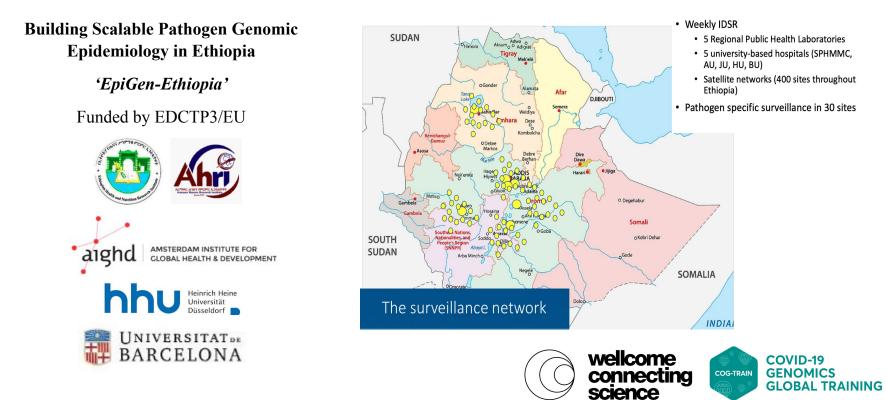
 Building capacity for integrated pathogen genomic surveillance for informed public health decision process

Overarching *specific objectives* include:

- Strengthen collection and analysis of clinical and epidemiological data and clinical samples; perform translational clinical research demonstrating the application of genomic epidemiology to inform public health decision-making
- Enhance capacity pathogen genomic sequencing Ethiopia, including strengthening lab infrastructure, human work force, pathogen genomic data analysis, integration with metadata
- Develop and implement **innovative digital diagnostic platforms** and create **semi-real time mobile phone applications** for policy decisions
- Promote communities of practice and knowledge exchange through fostering African collaboration and networking on pathogen genomic surveillance



Pathogen Genomics Centre of Excellence @EPHI Overview of the project



Current status of projects

- Current active Pathogen Genomic studies at EPHI:
 - EpiGen (recently funded by EDCTP3/EU BMGF)
 - DESTINE project on Hepatitis-C molecular epidemiology (fund: NIHR-UK)#
 - SUPER (Africa CDC/BMGF)
 - FUO-MetaGenomics (NIH-funded)
- Infrastructure already available at EPHI will be strengthened



Future plans

- **EpiGen** and other projects in collaboration with Africa CDC to serve as a platform
- Establishing 'Center of Excellence' for Pathogen Genomics at the EPHI
- Foster collaborations with international and regional bodies



Small group activity

You will be split into small groups.

What are the key focus capacity development areas needed in your country?

On your worksheet, there are areas which are important for sequencing. You
need to identify what exists in your country now (as much as you can) and
then discuss and decide what would need to be improved/put in place to run
successful sequencing.

