

# SURVEY TO ASSESS Next-Generation Sequencing (NGS) Capacity of the National Public Health Laboratories (NPHLs) in the East African Community (EAC)

## 1. NPHLs Informations

### 1.1. EAC Partner State

### 1.2. Name of the National Public Health Laboratory

### 1.3. Phone Number

Country Code

Phone Number

### 1.4. E-Mail \*

example@example.com

### 1.5. Contact/Focal Person

First Name

Last Name

## 2. NGS Equipments in the NPHLs

Which of the following NGS instruments are available in your laboratory?

### 2.1. Illumina NGS Platforms \*

iSeq100

MiniSeq

- MiSeq
- NextSeq 550
- NextSeq 1000/2000
- NovaSeq 6000
- NovaSeq X
- HiSeq
- None of the above

## 2.2. Oxford Nanopore Technologies (ONT) NGS Platforms \*

- MinION
- PromethION
- SmidgION (Smartphone Sequencer)
- GridION
- None of the above

## 2.3. Pacific Biosciences (PacBio) NGS Platforms \*

- Sequel system
- Sequel II system
- Sequel IIe system
- Revio system
- Onso system
- None of the above

# 3. Bacterial Samples Processing Capabilities

## 3.1. Which of the following can be performed by your Lab? \*

- Bacterial genomic DNA isolation
- Library preparation
- Bacterial whole genome sequencing
- Bacterial genome assembly and annotation
- Antimicrobial resistance (AMR) genotyping
- Data submission (e.g. raw reads, genome sequences, ...) in public repositories, including GenBank, ENA, SRA.

## 3.2. How many of these pathogens isolates have been sequenced yearly in your lab?

2019      2020      2021      2022      2023

Acinetobacter spp.

Escherichia coli

Klebsiella pneumoniae

Neisseria gonorrhoeae

Salmonella spp.

Shigella spp.

Staphylococcus aureus

Streptococcus pneumoniae

Other non-GLASS-priority pathogens

### 3.3. How many of these pathogens isolates have been analyzed yearly in your lab?

	2019	2020	2021	2022	2023
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Acinetobacter spp.

Escherichia coli i

Klebsiella pneumoniae

Neisseria gonorrhoeae

Salmonella spp.

Shigella spp.

Staphylococcus aureus

Streptococcus pneumoniae

Other non-GLASS-priority pathogens

## 4. Bioinformatics Computing Capacity Assessment

### 4.1. Do you have a dedicated bioinformatics computing facility or workspace in the laboratory? \*

Yes

No

### 4.2. What computational resources are available in your laboratory? (Check all that apply)

High-performance computing (HPC) cluster

Standalone servers

Workstations

Cloud computing services

None

**4.3. Do you have access to a reliable internet connection for data transfer and online analysis? \***

Yes

No

**4.4. Do you have staff with bioinformatics expertise and training in your laboratory? \***

Yes

No

**4.5. Do you have access to a reliable internet connection for data transfer and online analysis? \***

Yes

No

**4.6. If yes, please rate the level of bioinformatics expertise among your staff:**

Novice (Limited or no experience)

Intermediate (Basic skills and experience)

Advanced (Proficient with substantial experience)

Expert (Highly skilled with extensive experience)

**4.7. Please specify the roles of staff with bioinformatics expertise (check all that apply):**

NGS Data analysis

Sequence analysis

Pipelines/Workflows development

Database management

Training and capacity building

**4.8. Are there any collaborations with external organizations or institutions for bioinformatics support? \***

Yes

No

**You are almost done...**