CLIMADE AFRICA WORKING GROUP MEETING MINUTES – WEEK 11

Host: Centre for Epidemic Response and Innovation (CERI)

Date: May 9, 2023

Time: 12:00 – 13:00 p.m. (SAST)

Facilitators: Dr Eduan Wilkinson, Dr Monika Moir and Dr Yeshnee Naidoo

Attendance/ No. of Participants: 83

Start time: 12:03p.m. (SAST)

Purpose of the meeting

Sequencing methods for Cholera

Agenda Items

- 1. Welcome and update by Dr Monika Moir.
- Introduction to the presentation by Dr Yeshnee Naidoo and Presentation by Lucious
 Chabuka from the Public Health Institute of Malawi and an MSc fellow at CERI Tygerberg
 laboratories.
- 3. Questions and Discussion

Discussion points and questions

- Mr. Chabuka discussed the adaptive protocol for Cholera whole genome sequencing during the Malawi epidemic which included:
 - The history and comparison of cholera outbreaks in Malawi
 - The geographical range of outbreaks and specimen collection.
 - Sample preparation
 - sub-culture samples suspended in phosphate buffer.
 - DNA isolation using Qiagen QIAmp DNA mini kit

- The protocol was adapted for the Nextseq 1000 platform.
- Whole genome sequencing using Illuminias DNA Prep kit.
- Detailed explanation on the DNA tagmentation, amplification and library preparation steps.
- DNA quantification using Qubit or Pico green.
- Fragment size selection workflow.
- Normalization steps
- 2. The sequencing protocol can be adapted to other viruses and bacterium.
- 3. Illumina DNA Prep Kit formerly known as the Nextera DNA prep kit.
- 4. Stored samples without sub-culturing have not been tired for this protocol.
- 5. Cholera and Arbovirus sequencing protocols are available on the CLIMADE GitHub link: https://github.com/CERI-KRISP/CLIMADE/tree/master/protocols
- 6. Session recording will be available on the GitHub repository.

Adjournment and Closing points.

1. Dr Eduan Wilkinson adjourned the meeting at 12:52 p.m. (SAST).

Next Meeting

Tuesday, May 16, 2023, at 12:00 – 13:00 p.m. (SAST)

Submitted by: Yajna Ramphal

Approved By: Monika Moir