



## SOP Title: qPCR for *Rickettsia* spp. (Typhus and Spotted Fever Group)

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### 1. Purpose

To ensure a standardized and consistent approach to the PCR-based detection of *Rickettsia* spp. DNA in **whole blood samples** collected as part of the ZAFI project.

### 2. Scope

This SOP applies to all laboratory personnel involved in the processing, analysis, and interpretation of samples for *Rickettsia* spp. PCR testing under the ZAFI project across Kenya, Uganda, and Ethiopia.

### 3. Abbreviations

- **qPCR:** Quantitative Polymerase Chain Reaction

### 4. Responsibilities

| Role                              | Responsibility                                                                                                                                                                                                                                                         |
|-----------------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>Laboratory technician</b>      | Perform laboratory testing according to this SOP; handle and process samples appropriately; adhere to biosafety and quality control procedures; document results accurately in logbooks and KOBO tools; report deviations or issues to the laboratory lead/supervisor. |
| <b>Laboratory lead/supervisor</b> | Oversee laboratory testing; ensure staff competency and adherence to quality and biosafety standards; review and validate results; escalate issues as appropriate.                                                                                                     |
| <b>Site PI</b>                    | Ensure relevant staff are trained on this SOP.                                                                                                                                                                                                                         |

### 5. Safety and Precautions

- Preparation of mastermix should be conducted in a “Clean room” or “Clean hood” that is separate to template DNA and any amplification procedures to prevent contamination
- Template (DNA) and samples should be stored separately to mastermix and assay reagents
- When calculating the volume of mastermix to make, include enough volume for a single positive control per target and a negative control

## 6. Consumables and Reagents

| Item                                          |                                         |
|-----------------------------------------------|-----------------------------------------|
| Reagents                                      | Equipment                               |
| TaqMan Fast Advanced Master Mix               | Microcentrifuge                         |
| Primers and Probes                            | Pipettes (10µl, 20 µl, 200 µl, 1000 µl) |
| Nuclease Free Water                           | Vortex                                  |
| Rickettsia Standard Positive Control (gBlock) | <b>Tubes/Plates/other</b>               |
|                                               | Plastics consumables                    |
|                                               | Pipette tips                            |
|                                               | Disposable gloves                       |

## 7. qPCR Procedure

### 7.1 Preparing the PCR Reaction Mix

- Retrieve the working (10uM) stocks of primers, probes and master mix for the Rickettsia qPCR and fully thaw reagents on ice. (Details of preparation of primers is detailed in **SOP 325 Preparation of primers and probes for qPCR**).
- When completely thawed vortex each tube briefly and spin in microcentrifuge to collect liquid at the bottom.
- Calculate the number of reactions needed for the samples (one reaction per sample) **including a positive control for each target and a negative control**. Multiply the volume per reaction of each component by the total number of reactions.
- Combine these volumes into a 1.5ml Eppendorf. **\*This should be done in a separate room/hood to template addition.\***

| Table of Reactions                   |                     |                        |                                                                        |
|--------------------------------------|---------------------|------------------------|------------------------------------------------------------------------|
| Component                            | Final concentration | Volume per Reaction    | Total volume to add to the 1.5ml Eppendorf assuming <i>n</i> reactions |
|                                      |                     | 96-well (0.1ml) plates |                                                                        |
| <b>Reaction mix</b>                  |                     |                        |                                                                        |
| TaqMan Fast Advanced Master Mix (2x) | 1x                  | 5.50                   | 5.5 x <i>n</i>                                                         |
| Forward Primer (10uM)                | 200nM               | 0.28                   | 0.28 x <i>n</i>                                                        |
| Reverse Primer (10uM)                | 200nM               | 0.28                   | 0.28 x <i>n</i>                                                        |
| Probe (10uM)                         | 200nM               | 0.28                   | 0.28 x <i>n</i>                                                        |
| NFW                                  | -                   | 4.68                   | 4.68 x <i>n</i>                                                        |
| <b>Aliquot per well</b>              |                     |                        |                                                                        |
| Reaction mix                         |                     | <b>10ul</b>            |                                                                        |
| DNA/Template volume                  |                     | 2.5ul                  |                                                                        |
| <b>Total volume</b>                  |                     | <b>12.5ul</b>          |                                                                        |

- Vortex briefly to mix and spin to ensure all the liquid is at the bottom
- Transfer 10µl of PCR Reaction Mix to each well of the plate
- Retrieve DNA eluates immediately prior to template addition and thaw on ice
- Transfer plate to template addition room/space and add 2.5µl DNA (Extracted from sample), or nuclease-free water for Non-Template Control, to each well. Include Nuclease Free Water as a Non-Template Control and positive DNA/RNA as a positive control.

- i) Seal the reaction plate with optical adhesive film or strip-caps and then centrifuge briefly to bring the PCR reaction mix to the bottom of the well.

### 7.2. Setting up the machine

- a) Set up the following run profile on the qPCR machine.  
 b) Ensure each well is selected to collect data in the FAM channel

| Thermal Profile for Rickettsia qPCR |             |      |        |                                    |
|-------------------------------------|-------------|------|--------|------------------------------------|
| Step                                | Temperature | Time | Cycles | Fluorescence acquisition (channel) |
| Hold                                | 50          | 2:00 | 1      |                                    |
| Hold                                | 95          | 0:20 | 1      |                                    |
| Denature                            | 95          | 0:03 | 40     |                                    |
| Anneal/Extend                       | 60          | 0:30 |        | FAM                                |

### 7.3 Interpretation

- a) The negative control must have no amplification present in any channel  
 b) The positive controls must have amplification in their respective channels (FAM)  
 c) Results with a Ct value below 38 can be considered positive.

### 8. Record Keeping

- Enter results into the results sheet
- Enter results into the dedicated **KOBO form**

### 9. Primer Sequences and Further Details

| Primer/probe ID | Sequence                                            |
|-----------------|-----------------------------------------------------|
| Rickettsia-F    | TCG CAA ATG TTC ACG GTA CTT T                       |
| Rickettsia-R    | TCG TGC ATT TCT TTC CAT TGT G                       |
| Rickettsia_CS-P | 6-FAM-TGC AAT AGC AAG AAC CGT AGG CTG GAT G-BHQ-1-3 |

### 10. Review and Approval

Approved By: *Prof Siobhan Mor*

Title: *Chief Investigator, ZAFI*

Date: *3 March 2026*