Investigating whether a new test can be used to differentiate between viruses, malaria and bacteria as causes of fever among children in Ghana

Submission date 06/03/2024	Recruitment status No longer recruiting	Prospectively registered
		☐ Protocol
Registration date	Overall study status	Statistical analysis plan
25/04/2024	Ongoing	Results
Last Edited	Condition category	Individual participant data
11/03/2025	Infections and Infestations	[X] Record updated in last year

Plain English summary of protocol

Background and study aims

This study will evaluate a new test for the cause of fever in African children. There is an unmet need for diagnostic tests that can be used in African countries where resources are scarce. Most of the time, when children have a fever, the only test available is a malaria test. If this test is negative, they often receive antibiotic treatment because tests to distinguish between other causes of infection (e.g., bacteria or viruses) are unavailable. The aim of this study is to use how the human immune system responds to infection to tell which type of infection someone has. The researchers will detect genes that are turned on and off in the blood which are specific to either bacterial, viral, or malaria infection.

Who can participate?

Children between the ages of 1 month and 15 years presenting to the recruiting facilities with fever

What does the study involve?

The study will record participant data, some of which include the clinician's notes on the signs /symptoms, medications, and preexisting conditions, and demographic information such as age, sex, location, and ethnicity. In addition to this, the researchers will request the results of laboratory tests run for routine healthcare. They will also take about a spoonful of blood for laboratory testing for the study.

What are the possible benefits and risks of participating?

There are no direct benefits to taking part in this study. However, it is hoped that the information gained from this study may someday make it easier and faster for doctors to diagnose individuals infected with germs that cause illness and fever. This study will benefit the scientific community by providing information for future use in the evaluation of this new test. There is minimal risk associated with the collection of a blood specimen. The participant may experience some mild pain, discomfort, and/or bruising. These risks are no greater for this study

than for routine testing from blood samples. Well-trained laboratory personnel will perform this according to standard guidelines to minimize the possible risks.

Where is the study run from?

The West Africa Centre for Cell Biology of Infectious Pathogens at the University of Ghana (Ghana)

When is the study starting and how long is it expected to run for? August 2022 to August 2026

Who is funding the study?

The study is funded by the NIHR (NIHR-Global Health Research Group's Digital Diagnostics for African Health Systems, project reference NIHR134694) using UK International Development funding from the UK Government to support global health research. The views expressed are those of the authors and not necessarily those of the NIHR or the UK government.

Who is the main contact? Flavia Kaduni Bawa, fkbawa@st.ug.edu.gh

Contact information

Type(s)

Public, Scientific

Contact name

Mrs Flavia Bawa

ORCID ID

http://orcid.org/0000-0002-0027-2917

Contact details

West Africa Centre for Cell Biology of Infectious Pathogens Department of Biochemistry, Cell and Molecular Biology University of Ghana Accra Ghana LG54 +233 (0)540295710 fkbawa@st.ug.edu.gh

Type(s)

Principal Investigator

Contact name

Prof Aubrey Cunnington

ORCID ID

http://orcid.org/0000-0002-1305-3529

Contact details

Section of Paediatric Infectious Disease and Centre for Paediatrics and Child Health Imperial College London
London
United Kingdom
W2 1PG
+44 (0)2075943695
a.cunnington@imperial.ac.uk

Type(s)

Scientific

Contact name

Dr Samuel Duodu

ORCID ID

http://orcid.org/0000-0003-2791-8949

Contact details

West Africa Centre for Cell Biology of Infectious Pathogens Department of Biochemistry, Cell and Molecular Biology University of Ghana Accra Ghana LG54 +233 (0)243279353 saduodu@ug.edu.gh

Type(s)

Scientific

Contact name

Dr Effua Usuf

ORCID ID

http://orcid.org/0000-0002-6515-7186

Contact details

Medical Research Council Unit The Gambia at the London School of Hygiene & Tropical Medicine Atlantic Boulevard

Fajara Banjul

Gambia

PO Box 273

+220 (0)4495835

eusuf@mrc.gm

Type(s)

Scientific

Contact name

Dr Jethro Herberg

ORCID ID

http://orcid.org/0000-0001-6941-6491

Contact details

Section of Paediatric Infectious Diseases
Department of Infectious Diseases
Imperial College London
Wright-Fleming Institute
Norfolk Place
London
United Kingdom
W2 1PG
+44 (0)20 7594 3990
j.herberg@imperial.ac.uk

Additional identifiers

EudraCT/CTIS number

Nil known

IRAS number

ClinicalTrials.gov number

Nil known

Secondary identifying numbers

NIHR134694

Study information

Scientific Title

Evaluating the potential of a Lacewing host-response assay to distinguish between malaria, bacterial and viral infection as causes of acute childhood febrile illness in The Gambia and Ghana

Acronym

LacewingAFI

Study objectives

The measurement of the expression of a small number of genes in blood using a point-of-care molecular diagnostic test can be used to distinguish between viruses, bacteria and malaria infections

Ethics approval required

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Ethics approval(s)

1. Approved 09/10/2023, Ghana Health Service Ethics Review Committee (Research and Development Division, Ghana Health Service, Accra, MB190, Ghana; +233 (0)302681109; ethics. research@ghs.gov.gh), ref: GHS-ERC: 017/06/23

2. Approved 29/06/2023, Ethics Committee for Basic and Applied Sciences (Ethics Committee for Basic and Applied Sciences, University of Ghana, Accra, LG1195, Ghana; +233 (0)302213820; ethicscbas@ug.edu.gh), ref: ECBAS 038/22-23

Study design

Multicenter cross-sectional study

Primary study design

Observational

Secondary study design

Cross sectional study

Study setting(s)

Hospital

Study type(s)

Diagnostic

Participant information sheet

Not available in web format. please use contact details to request a participant information

Health condition(s) or problem(s) studied

Diagnosis of malarial, viral and bacterial infections among children with fever

Interventions

Clinical samples will be obtained from participants recruited from the hospitals and stored samples from previous ethically approved studies. These will be tested for host signature transcripts using a benchtop reverse transcription loop-mediated isothermal amplification (RT-LAMP) assay and the Lacewing device. The diagnostic performance, sensitivity and specificity of the device will be assessed by comparing results with gold standards such as microscopy, bacterial cultures, and molecular diagnostics (for example TaqMan array card).

Intervention Type

Device

Pharmaceutical study type(s)

Not Applicable

Phase

Not Applicable

Drug/device/biological/vaccine name(s)

Lacewing

Primary outcome measure

Diagnostic accuracy will be measured using the area under the receiver operator curve (AUC) for the results of the Lacewing device compared against reference standard (blood film microscopy, sterile site culture, and molecular pathogen detection), assessed at the time of clinical presentation.

Secondary outcome measures

- 1. The prevalence of different types of infection classified by gene expression test will be assessed for samples collected at time of clinical presentation
- 2. The usability of the test will be evaluated through stakeholder engagements on the day that the primary outcome is measured

Overall study start date

01/08/2022

Completion date

31/08/2026

Eligibility

Key inclusion criteria

- 1. The participant is between the ages of 1 month and 15 years
- 2. Participant has a reported fever or recorded fever of \geq 37.5°C in the past 24 hours and is suspected by a clinician to have an infection
- 3. Parental permission and child assent have been obtained

Participant type(s)

Patient

Age group

Child

Lower age limit

1 Months

Upper age limit

15 Years

Sex

Both

Target number of participants

230

Total final enrolment

353

Key exclusion criteria

- 1. Children aged 16 years or older
- 2. Children without recorded/reported fever.
- 3. Inability to provide informed consent.
- 4. Children with fever from a suspected hospital-acquired infection
- 5. Children suspected by clinicians to have a high-consequence infectious disease such as viral haemorrhagic fever
- 6. Children with known HIV or other immunodeficiency

Date of first enrolment

15/03/2024

Date of final enrolment

10/03/2025

Locations

Countries of recruitment

Ghana

Study participating centre Ledzokuku-Krowor Municipal Hospital

Ghana Health Service Teshie-Tsuibleoo Accra Ghana GZ-103-2962

Study participating centre Oda Government Hospital

Ghana Health Service Akyem-Oda Oda Ghana P O Box 16

Study participating centre Kintampo Municipal Hospital

Ghana Health Service Kintampo Ghana PO Box 192

Study participating centre War Memorial Hospital

Ghana Health Service Navrongo Ghana PO Box 34

Sponsor information

Organisation

University of Ghana

Sponsor details

PO Box LG 25

Legon

Ассга

Ghana

LG25

+233 (0)302 213850

pad@ug.edu.gh

Sponsor type

University/education

Website

https://www.ug.edu.gh/

ROR

https://ror.org/01r22mr83

Funder(s)

Funder type

Government

Funder Name

NIHR Global Health Research Group on Digital Diagnostics for Africa

Results and Publications

Publication and dissemination plan

Planned publication in a high-impact peer-reviewed journal

Intention to publish date

20/06/2026

Individual participant data (IPD) sharing plan

The datasets generated during and/or analysed during the current study will be available upon request from Prof. Aubrey Cunnington (a.cunnington@imperial.ac.uk) or Dr. Samuel Duodu (Sa. duodu@ug.edu.gh)

IPD sharing plan summary Available on request