



# Wastewater surveillance in non-sewered settings

16 MAY 2024

# AGENDA

- **Wastewater-based Epidemiological Surveillance & the IWA Cluster**  
*Gertjan Medema, KWR Water Research Institute, Netherlands*
- **Non-sewered settings**  
*Sudhir Pillay, Water Research Commission, South Africa*
- **SARS-CoV-2 surveillance in non-sewered settings in South Africa**  
*Gina Pocock & Fiona Els, Waterlab & National Center for Infectious Diseases, South Africa*
- **Ensuring Health Equity: Infectious Disease Surveillance in South East Asia through Non-sewered Wastewater**  
*Leshan Wannigama, Chulalongkorn University and King Chulalongkorn Memorial Hospital, Thailand*
- **Wastewater-Based Multi-Pathogen Surveillance in Malawi: Lesson and possible application to One Health**  
*Petros Chigwechokha, Malawi University of Science and Technology, Malawi*
- **Wastewater surveillance from Vellore, India - utilities, insights and challenges**  
*Dilip Abraham, Christian Medical College Vellore, India*
- **Q&A Panel Discussion**  
*Speakers & Moderators*
- **Close**  
*Gertjan Medama & Sudhir Pillay*

## WEBINAR INFORMATION

- This webinar will be **recorded and made available “on-demand”** on the [IWA Connect Plus](#) platform and IWA Network website, with presentation slides, and other information.
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## WEBINAR INFORMATION



- **‘Chat’ box:** please use this for general requests and for interactive activities.
- **‘Q&A’ box:** please use this to send questions to the panelists. (We will answer these during the discussions)

***Please Note:** Attendees’ microphones are muted. We cannot respond to ‘Raise Hand’.*



# SPEAKERS & MODERATORS



**Gertjan Medema**  
Netherlands



**Petros Chigwechokha**  
Malawi



**Fiona Els**  
South Africa



**Sudhir Pillay**  
South Africa



**Gina Pocock**  
South Africa



**Dilip Abraham**  
India



**Leshan Wannigama**  
Thailand

# Wastewater Surveillance in Non-Sewered Settings

GERTJAN MEDEMA, SPECIALIST GROUP: HEALTH-RELATED WATER MICROBIOLOGY



# IWA CLUSTER WASTEWATER-BASED EPIDEMIOLOGICAL SURVEILLANCE

## Vision

Global network of professionals and institutions  
for rapid info exchange and dialogue within  
water and between water and public health

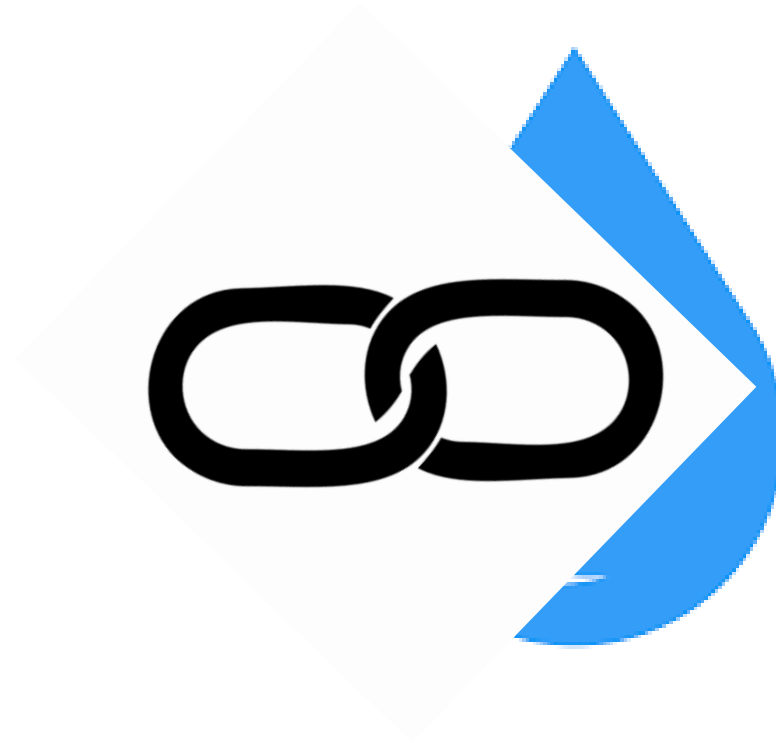


# MULTIDISCIPLINARY IN WATER





# LINKING WATER TO PUBLIC HEALTH



Deutsches Ärzteblatt





Gertjan Medema  
KWR  
Netherlands



Sudhir Pillay  
WRC  
South Africa



Gina Pocock  
Waterlab  
South Africa



Fiona Els  
NICD  
South Africa



Petros Chigwechokha  
University of Science  
& Technology  
Malawi



Leshan Wannigama  
Chulalongkorn  
University  
Thailand



Dilip Abraham  
Christian Medical  
College Vellore  
India

## GETTING TO KNOW YOU

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# IWA CLUSTER WASTEWATER-BASED EPIDEMIOLOGICAL SURVEILLANCE



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The Source

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wastewater surveillance for  
public health,  
with WHO, US CDC, MoH  
Thailand, Ottawa Public Health

# Surveillance in Non-Sewered Settlements: Non-Sewered Settings: The Why, What (and later How?)

SUDHIR PILLAY, SPECIALIST GROUP: NON-SEWERED SANITATION (NSS)

**inspiring change**

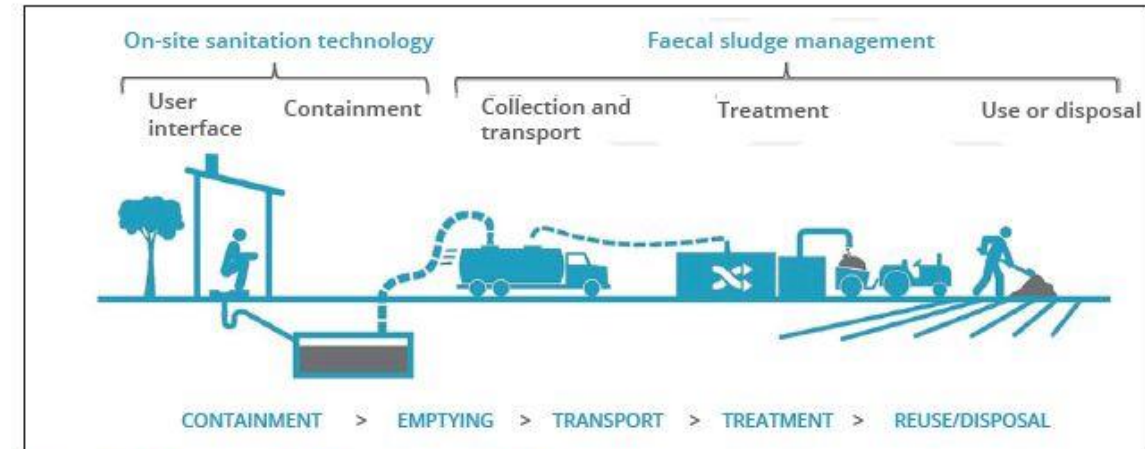
Main photo by WaterLab PTY LTD





## WHAT ARE NON-SEWERED AREAS

- Any sanitation system treating human excreta that operates **without connection to any sewer or drainage network**
- A **non-sewered sanitation (NSS)** system is a sanitation system that 1) is not connected to a networked sewer system and 2) collects, conveys, and fully treats the specific input, to allow for safe reuse or disposal of the generated output
- Where there is no sewer:
  - Greywater (washing water) discharge into toilet or environment
  - Faecal sludge can accumulate
  - On-site recycling (new generation systems)



**Figure 3.2: A typical FS Management System**

Source: Modified from Wikipedia.org, 2020

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# SANITATION VALUE CHAIN

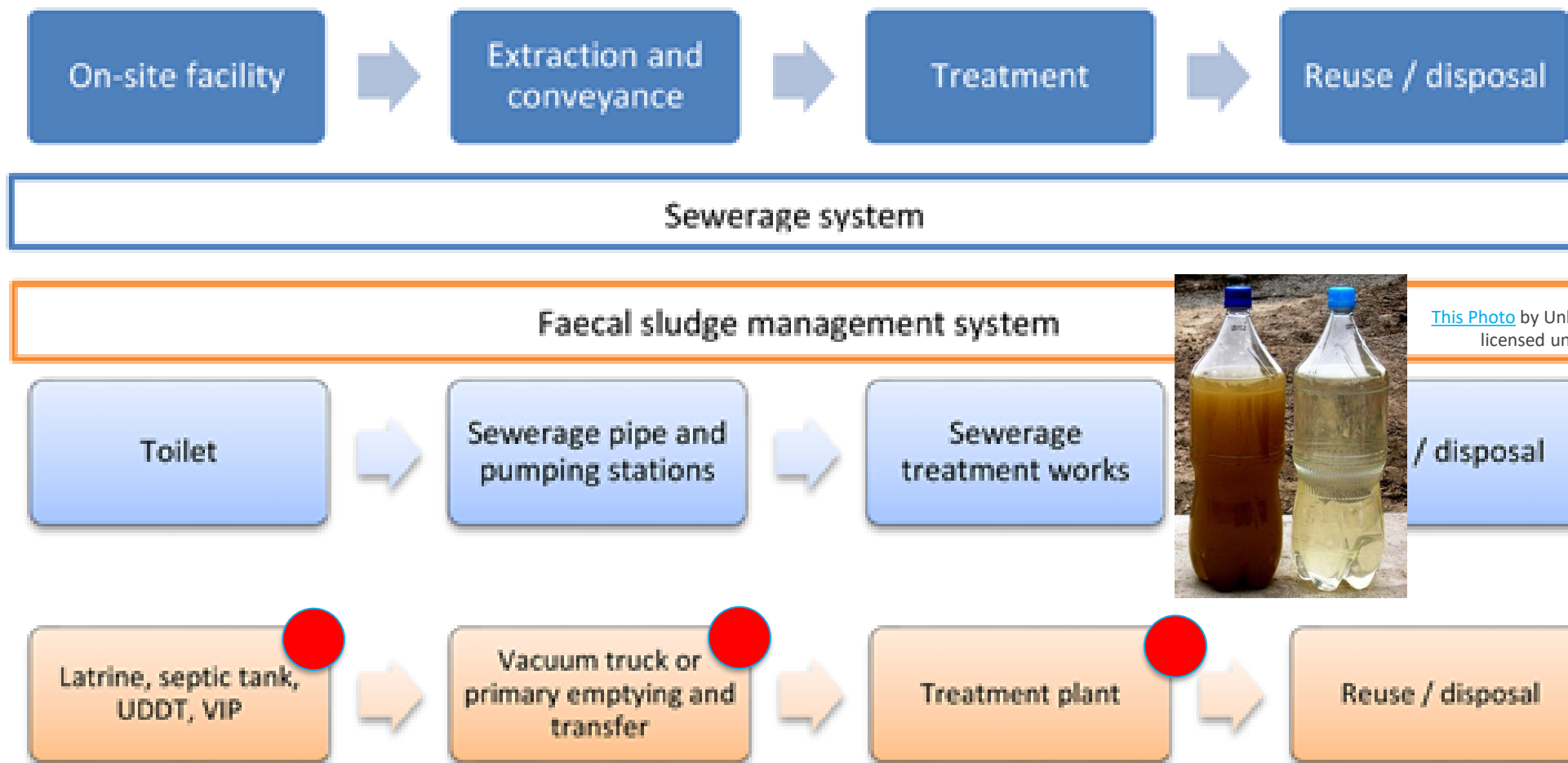
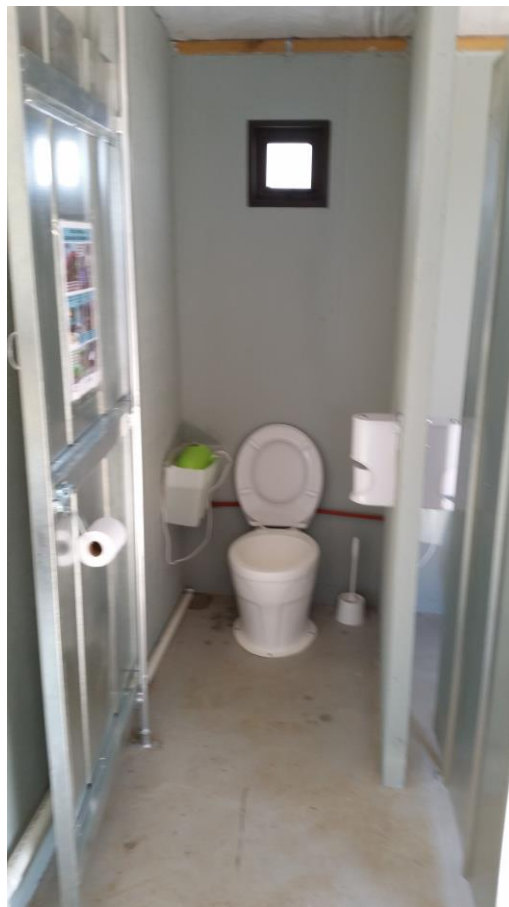


Image From: <https://iwaponline.com/washdev/article/8/2/176/39027/The-cost-of-urban-sanitation-solutions-a>

# NSS ENVIRONMENTS – HIGHLY VARIABLE EVEN IN SAME LOCALITY



Photos by WRC in same province



# SAMPLES – DIFFERENT WAYS TO COLLECT



Photos (a) Jimma University, (b) WRC, (c) University of Malawi, (d) WaterLab PTY LTD

## FUNDAMENTAL DIFFERENCES IN SAMPLE COMPOSITION

- **Heterogeneous Composition:** mixture of human excreta, water, urine, and other solid and liquid waste. Its composition can vary significantly depending on factors such as location, diet, and sanitation practices, making it challenging to establish consistent characterization parameters
- **Variable Chemical & Physical Properties:** wide range of organic and inorganic compounds, solids and water content, viscosity, density and particle size distribution
- **Pathogen Content:** usually contains a high concentration of pathogens, including bacteria, viruses, and parasites
- **Storage & Collection Variability:** wide range of systems from latrines to septic tanks, wet vs dry, different loading rates and designs
- **Regulatory & Standards Variability:** variation in designs and regulation
- **Sampling variability:** Analysis can be challenging due to its heterogeneous nature and the potential for stratification within storage systems, especially dry sanitation systems.



# HIGHLY VARIABLE WASTE IN NSS SYSTEMS

	Pit latrine sludge	High strength sludge from bucket latrines and public toilets	Low strength sludge from septic tanks	Sewage – in waterborne sewerage systems
Source	Brouckaert and Foxon, 2008	Heinss et al, 1998		
COD (mg/l wet)	90 000-225 000	20 000-50 000	< 10 000	500 – 2 000
COD (mg/g dry)	210-1230	571-1429	<333	50-200
N as NH <sub>4</sub> (mg/l wet)	9 000 (TKN)	2 000-5 000	<1 000	30-70
N as NH <sub>4</sub> (mg/g dry)	100 (TKN)	60-150	<33	3-7
Total solids (%)	20	>3.5	< 3	< 1
Soluble solids(mg/l wet)	220 000	≥30 000	≈ 7 000	200-700





# THE CHALLENGE OF NSS SURVEILLANCE

## ■ Variety of NSS systems

- Variable characteristics (not like WWTW)
- Variable sampling sites (toilets, transport like honeysuckers, treatment works)
- Not a regulated site
- Lack of formalised workers for sampling (training & SOPs required)

## ■ NSS systems are highly variable per site & country

- Variable design
- Washers vs wipers
- Dry vs flush
- Water ingress

## ■ Standards, Financing & Regulation

- Business case & modelling (vs clinical)
- Long-term financing
- Citizen-science approach? SOP for sampling, collection and analysis



## Methods for Faecal Sludge Analysis



Konstantina Velkushanova • Linda Strande • Mariska Ronteltap  
Thammarat Koottatep • Damir Brdjanovic • Chris Buckley

*Should we include into this or develop a separate technical guidance document?*

# HOW WOULD IT BENEFIT DEVELOPING COUNTRIES

- Urban slums – Social Distancing difficult
- Shared water & sanitation facilities
- Accessibility & affordability to medical care
- Availability of test kits for patient testing
- Alleviate supply chain for test kits & vaccines
- COVID-19: Inequity with vaccines



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# SARS-CoV-2 surveillance in non-sewered settings in South Africa

DR GINA POCOCK AND FIONA ELS

# BACKGROUND



- Wastewater surveillance has recently emerged as a flexible and robust method to monitor infectious diseases
- It has been used in various countries to monitor SARS-CoV-2
- Recently started to monitor for other pathogens (influenzas, cholera, typhoid etc)
- Wastewater infrastructure need to be in place for it to work effectively



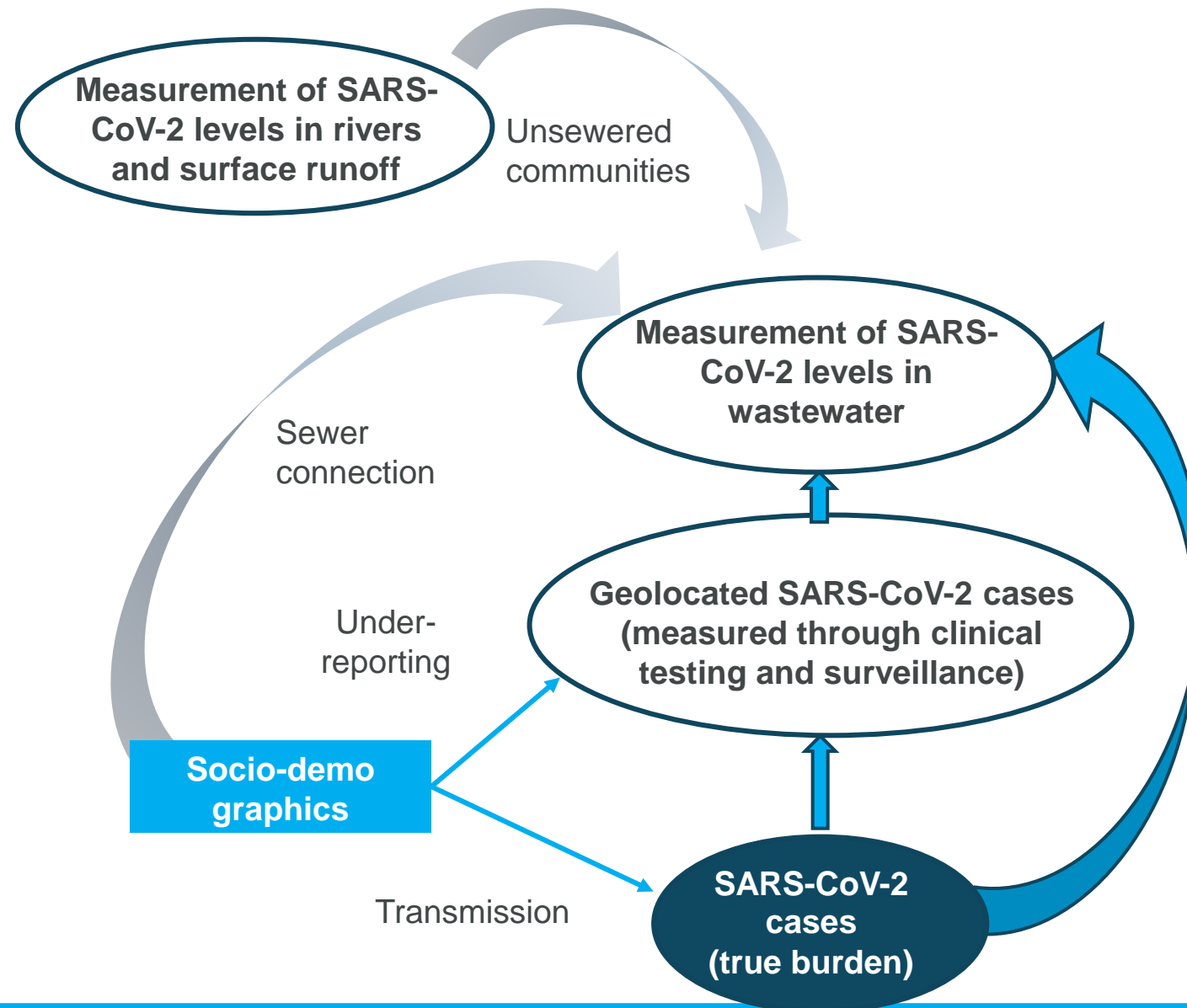
# BACKGROUND



- South Africa has a functioning and representative wastewater surveillance system for SARS-CoV-2 located at sentinel site wastewater treatment plants
  - This programme has achieved success in monitoring levels and genotypes circulating in the population
- However, 43% of the population is not connected to a sewer network (Statistics SA)
  - In South African metropolitan areas 16.8% live in informal dwellings
- Utility of environmental surveillance in unsewered areas is presently unmeasured in our context and data pertaining to wastewater surveillance will have global reference

**We developed a conceptual framework to help us understand what approach to take in evaluating environmental surveillance**







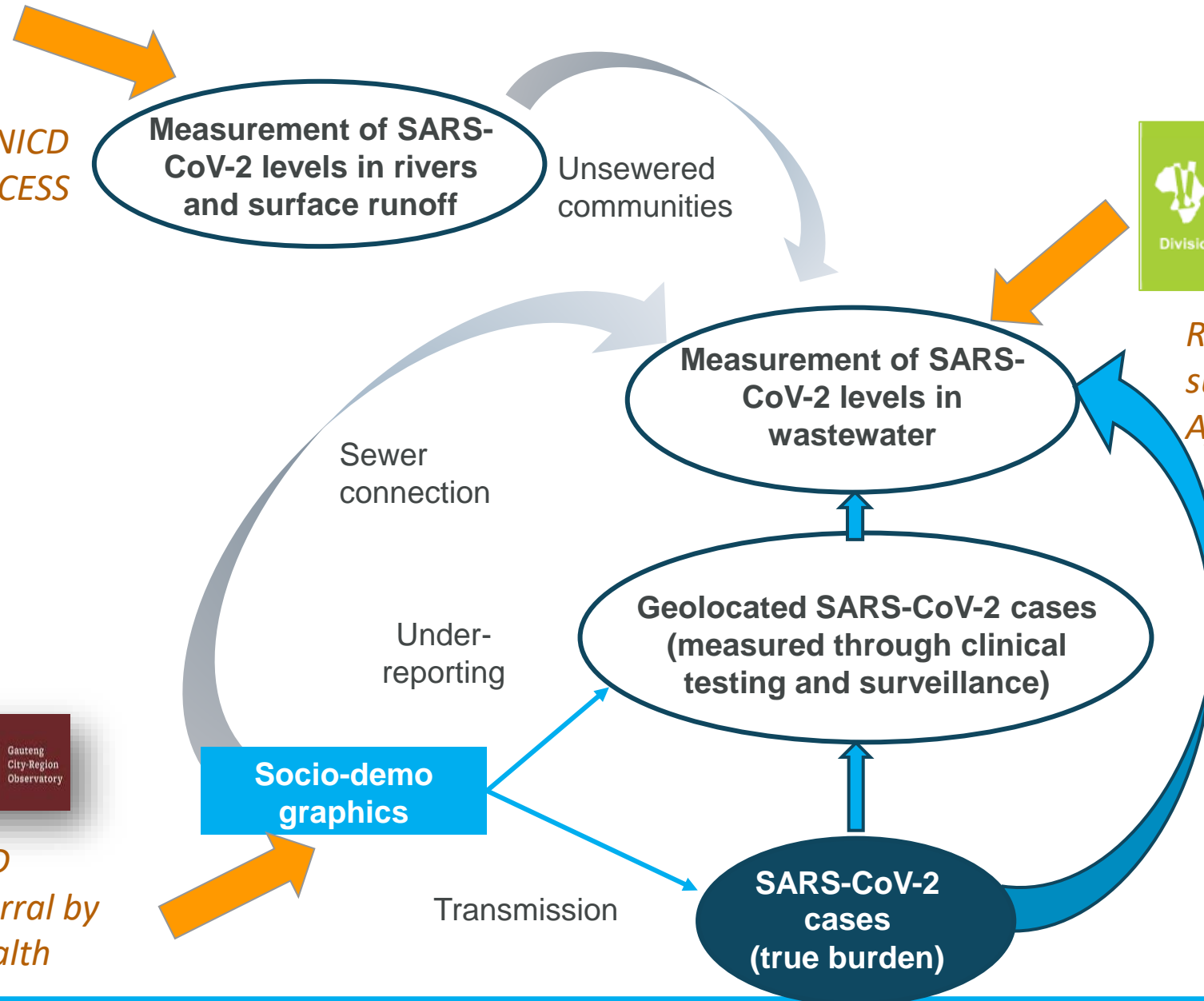
*Waterlab - identified by NICD researchers through SACCESS network participation*



*Routine wastewater surveillance in South Africa*



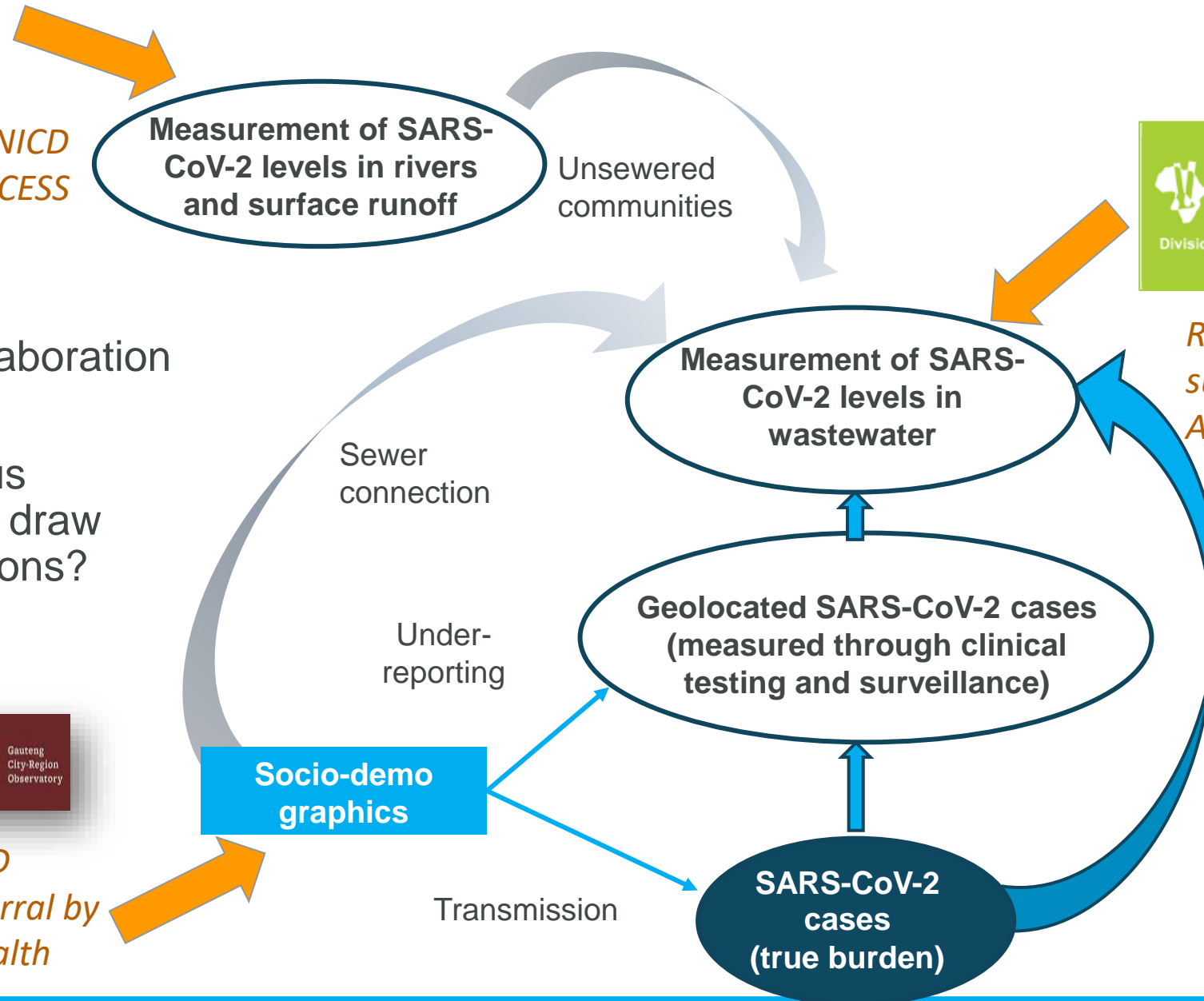
*GCRO - identified to NICD researchers through referral by Wits School of Public Health*



*Waterlab - identified by NICD  
researchers through SACCESS  
network participation*

- Interdisciplinary collaboration was needed
- How do we fit various datasets together to draw meaningful conclusions?

*GCRO - identified to NICD  
researchers through referral by  
Wits School of Public Health*



*Routine wastewater  
surveillance in South  
Africa*



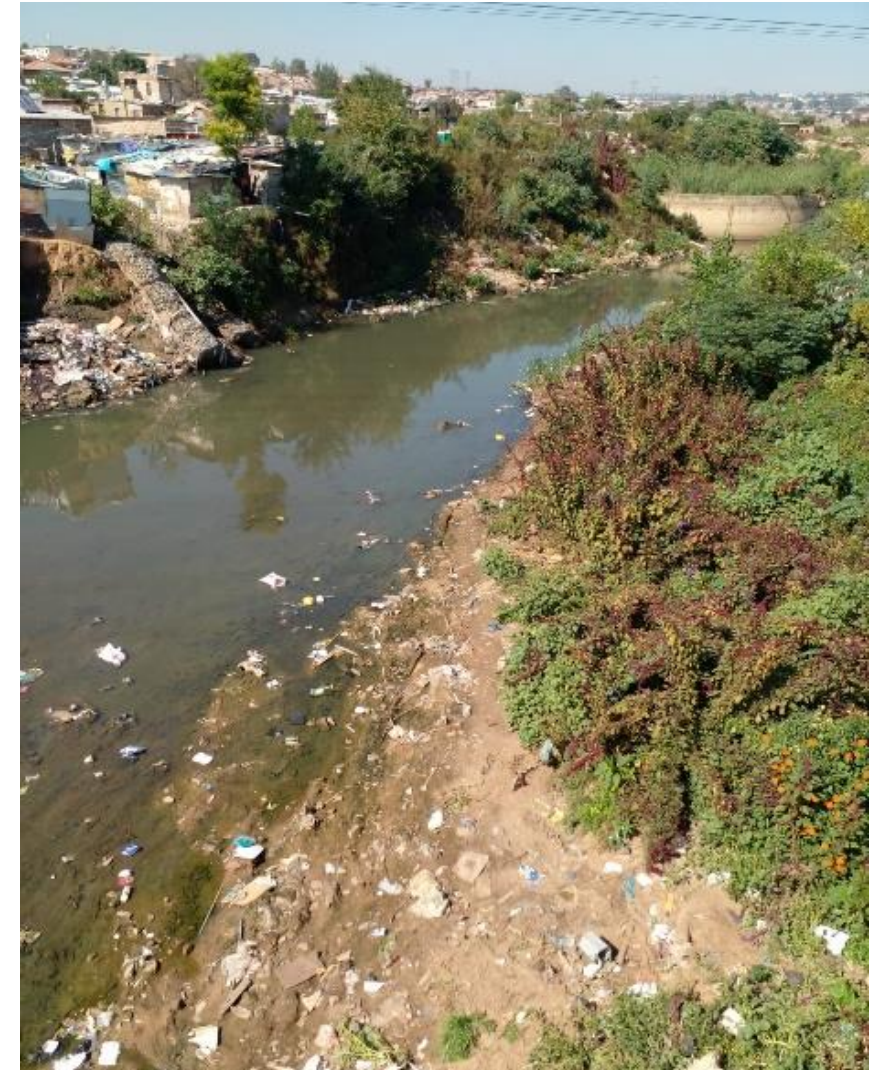
# AIMS AND OBJECTIVES

## Aim:

Characterise populations in non-sewered areas where wastewater samples were tested for SARS-CoV-2 to better understand the relationship with environmental surveillance

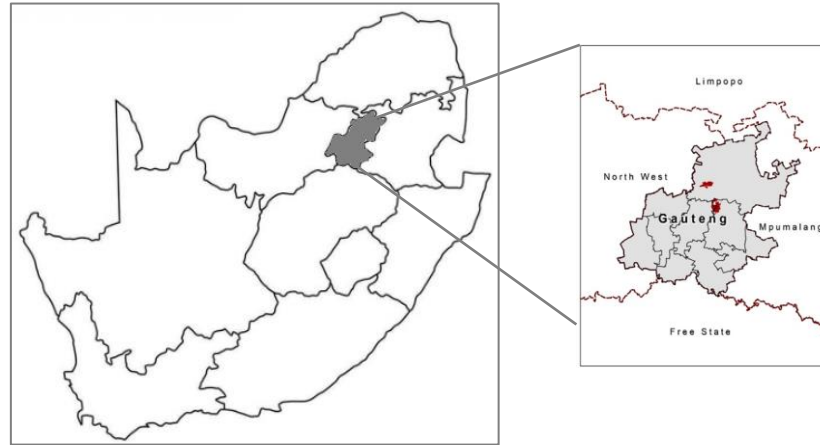
## Objectives:

1. To identify sites, collect samples and quantify SARS-CoV-2 in non-sewered settings in densely populated areas of Gauteng province
2. Characterise catchment areas (topography and drainage profile), population demographics and Quality of Life at sample collection sites
3. Geolocate laboratory-confirmed SARS-CoV-2 cases to catchment areas
4. To describe relationships between environmental SARS-CoV-2 concentration, demographics and laboratory-confirmed cases



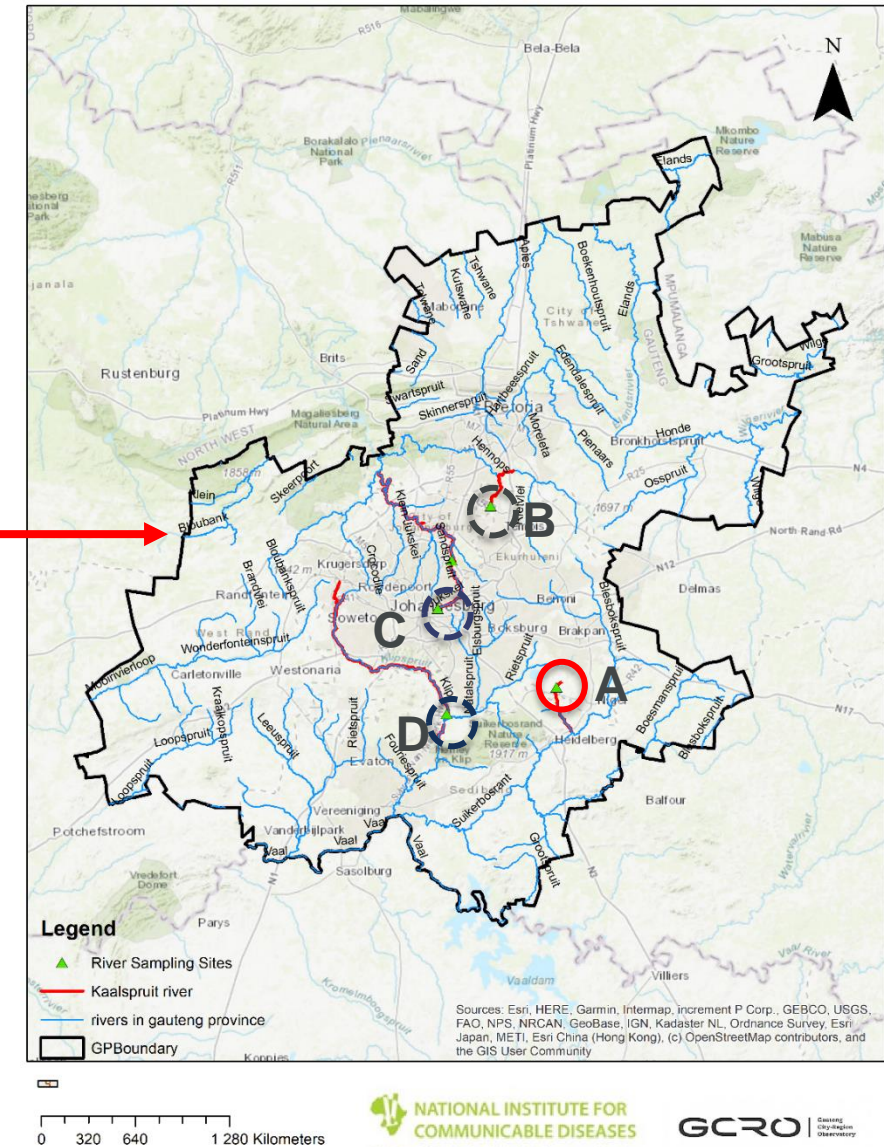
# METHODS

- Descriptive study to determine causal and contributory relationships regarding incidence of communicable disease and social determinants of health



- Gauteng is one of 9 Provinces in South Africa, and has 46 rivers
- We purposely identified 4 river sampling sites within the City of Ekurhuleni and the City of Johannesburg that are impacted by untreated surface run-off from informal and unsewered communities:
  - Blesbokspruit (A), Kaalspruit (B)
  - Jukskei River (C), Klipspruit (D)

## Hydrology of Gauteng





# METHOD: SAMPLE TYPES

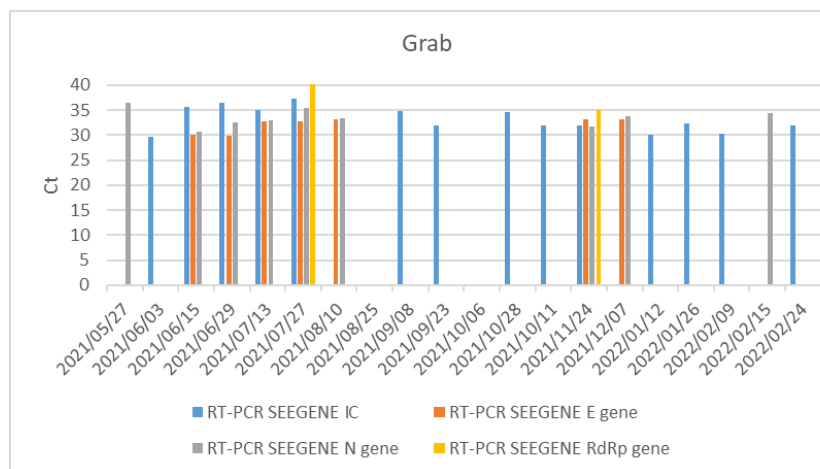
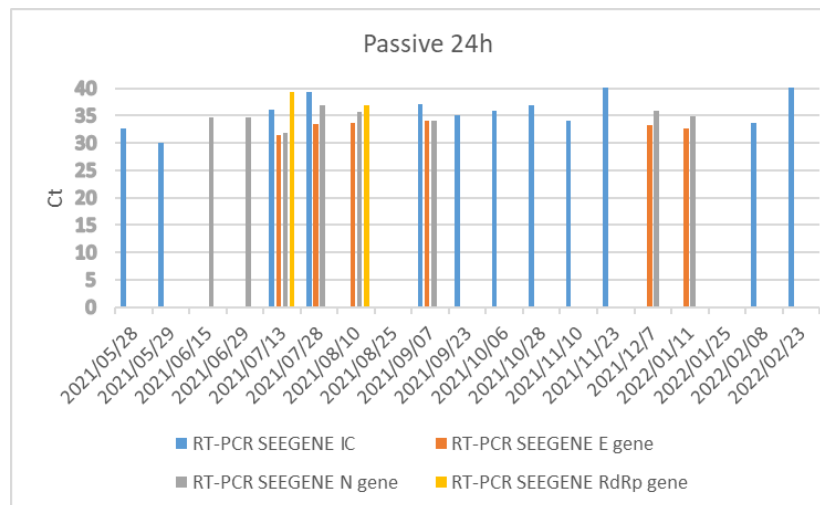
## Sample selection

### Grab Samples

- Identified informal settlements lacking access to sewerage sanitation located in close proximity to a river
- Rivers provide more stable sample points than greywater pools which are transient in nature

### Passive samples

- Device covered in shadecloth and anchored at sample point for a specific period, then viral nucleic acid eluted from gauze in the lab
- Passive sampling was conducted in parallel to grab samples at a subset of sample sites



Results: Grab vs Passive, Jukskei River downstream Alexandra



Concentration	Nucleic acid extraction	SARS-CoV-2 RT-PCR & amplification	Molecular analysis platform	Limit of quantification (genome copies/ml)
Skimmed milk flocculation	QIAamp® Ultrasens® Virus kit (1mL)	Allplex™ 2019-nCoV Assay and 2019_nCoV_N positive control plasmid	QuantStudio™ 5 Real-Time PCR System (Applied Biosystems)	16



# BLESBOKSPRUIT SAMPLING SITE

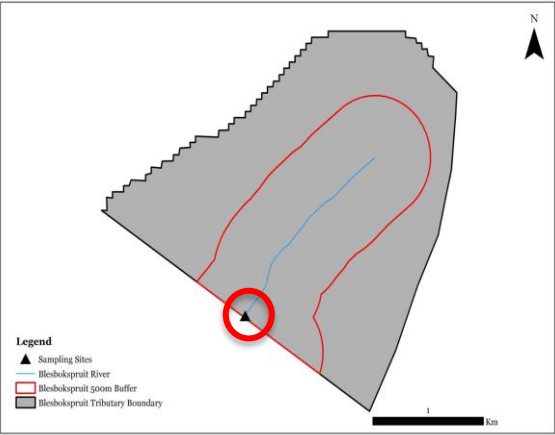


Fig 1: Buffer (red line) and catchment areas (black line) upstream of sampling site

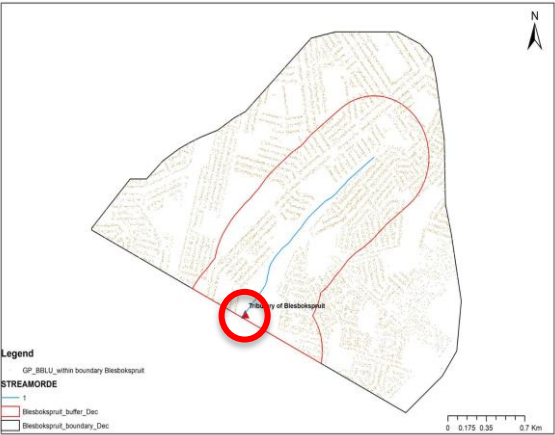


Fig 2: Building base land use upstream of sampling site

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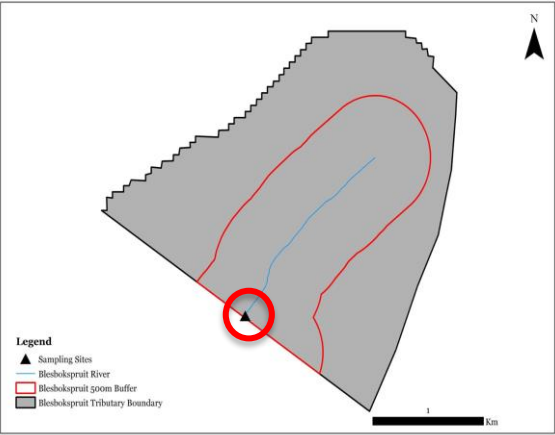


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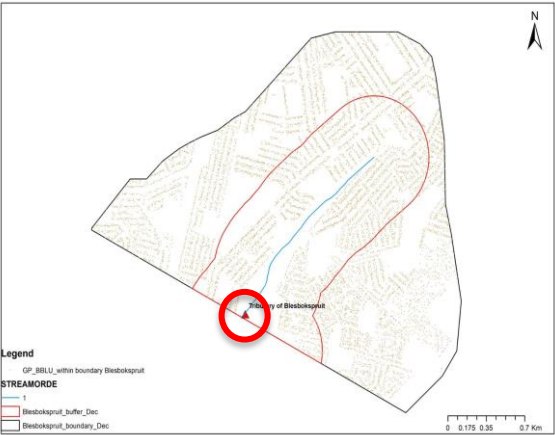


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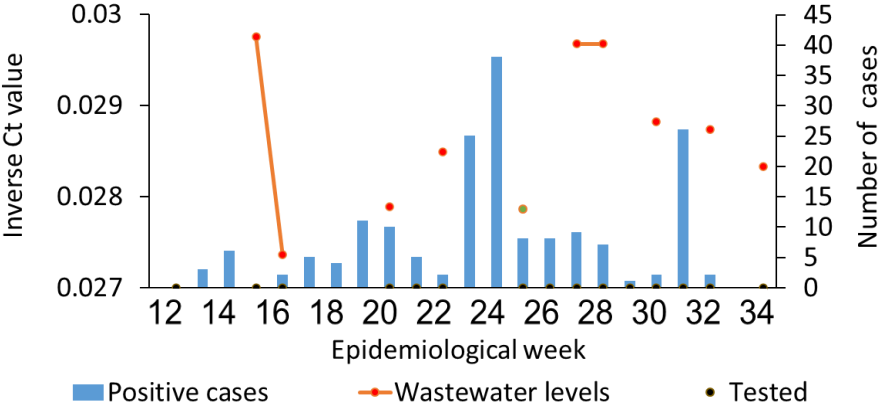


Fig 3: Geolocated SARS-CoV-2 cases in the catchment area (blue bars) and quantitative SARS-CoV-2 river results (Ct threshold), from epidemiological week 12-35, 2021.

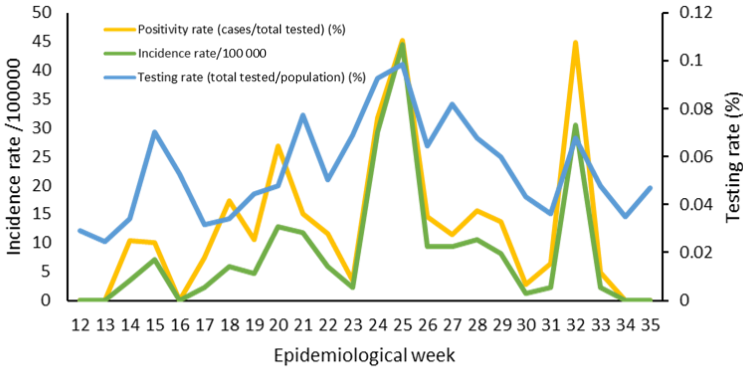


Fig 4: Positivity rate, incidence rate and testing rate of geolocated SARS-CoV-2 cases in the catchment area, from epidemiological week 12-35, 2021.

# BLESBOKSPRUIT SAMPLING SITE

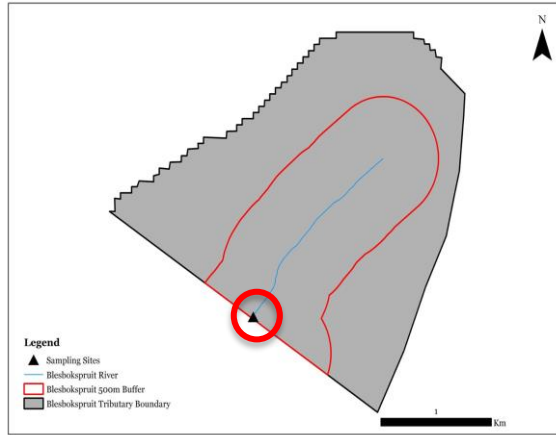


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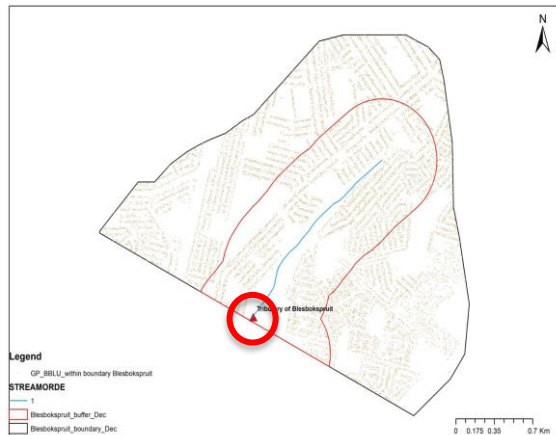


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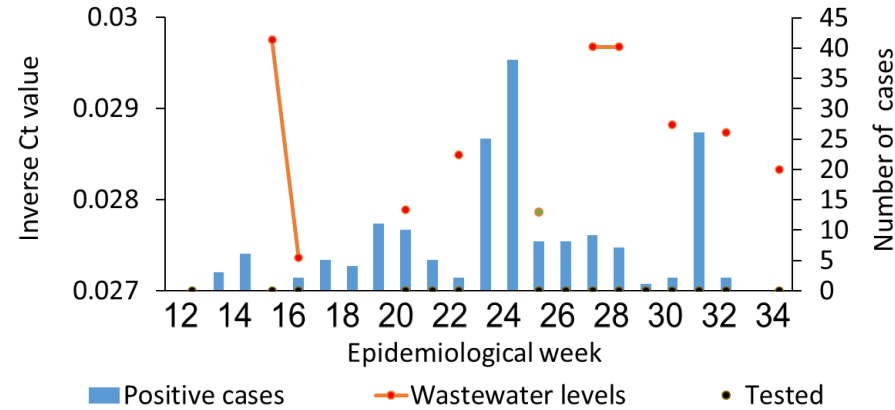


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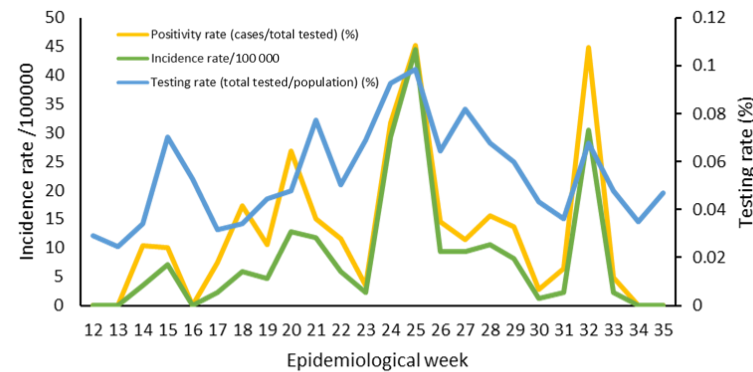


Fig 4: Positivity rate, incidence rate and testing rate of geolocated SARS-CoV-2 cases in the catchment area, from epidemiological week 12-35, 2021.

## Quality of Life Survey

	Blesbokspruit Catchment	City of Ekurhuleni
Total population	85,383	4,074,030
Population density (people/km <sup>2</sup> )	15,580	2,063
Cumulative incidence/100 000	172	2,470
Number of respondents	47	2214
<b>Economic:</b>		
Income <R1600	25 (53.0%)	830 (34.5%)
No income	1 (2.2%)	35 (1.5%)
Informal dwelling	3 (6.7%)	475 (21.5%)
% ≥ 3 persons per room	50.3	35.9
<b>Water:</b>		
Piped, into the dwelling	24 (50.8%)	2,505 (72.5%)
Piped, into the yard only	23 (49.2%)	714 (20.7%)
<b>Healthcare:</b>		
Uses public health care facilities	41 (88.7%)	2,385 (69.1%)
Tried and successfully tested for COVID-19	45 (95.9%)	3,384 (98.0%)
Struggled to access healthcare	2 (4.1%)	228 (6.6%)
COVID-19 Index (% unable to protect against COVID-19)	39.7	31.7
Quality of life (Score out of 100)	55.1	61.3

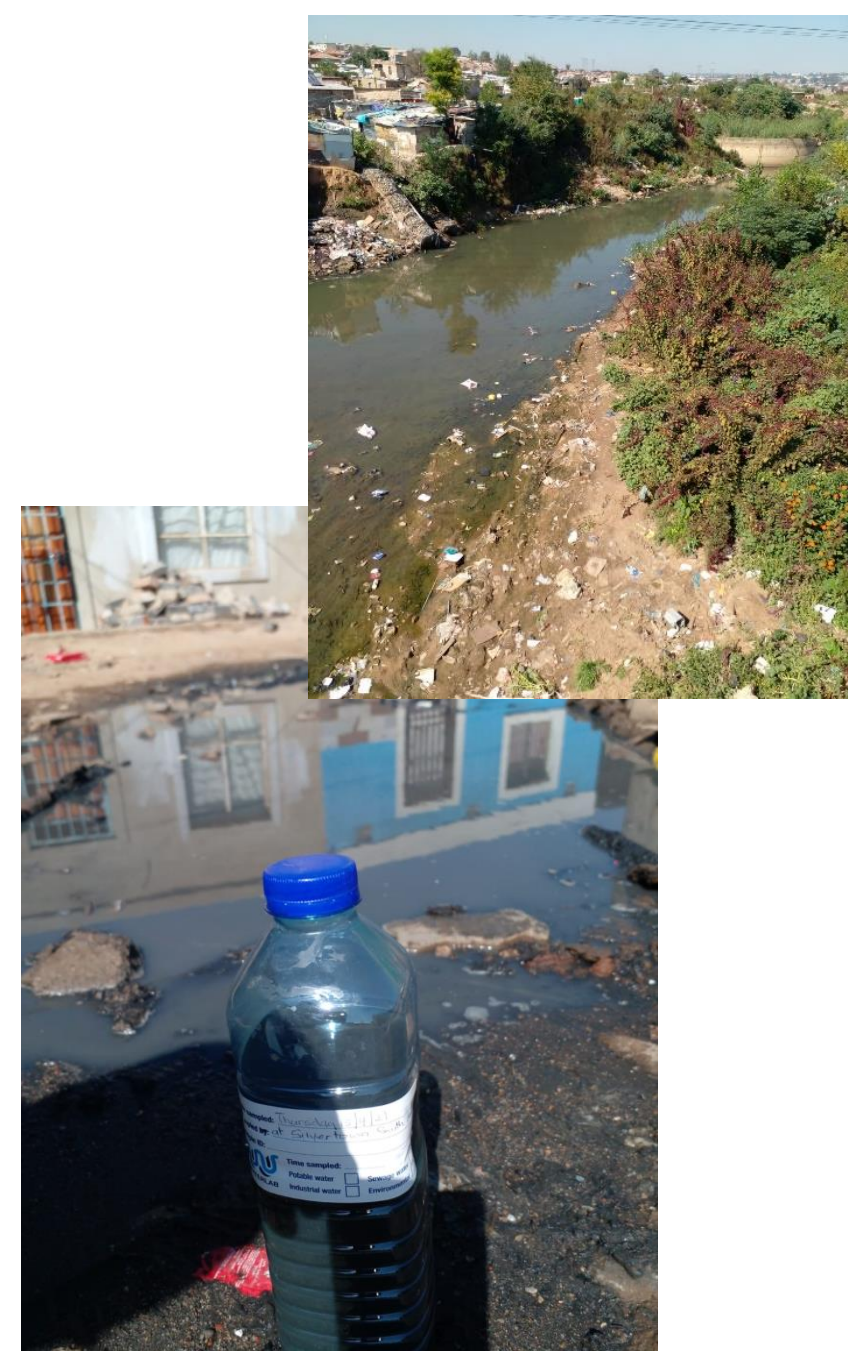


## INTERPRETATION AND IMPACT

- Positive results in rivers indicate that the river is being used for sanitary purposes, or there is broken infrastructure present
- Access to care is limited in informal settlements, and they have higher vulnerability,
- This could have impacts on disease transmission and severity, especially if patients don't get treatment on time
- Strengthening environmental surveillance in marginalised communities could bridge the gap for vulnerability
- When environmental surveillance indicate higher concentrations of disease (especially in vulnerable communities), additional support and services should be rendered

# KEY LESSONS AND CONCLUSIONS

- Sampling logistics can be difficult and time consuming, requiring extensive support from municipalities, river action groups and community leaders
  - Transport of large volumes of water is costly, and cold chain must be maintained out of rural areas
  - Dilution during rainy season may hinder detection
- Passive sampling may overcome issues of low yield during high dilution periods, allow for easier and cheaper transport of samples, and improved consistency
  - Sample processing is much quicker, although the logistics of passive sampling are more onerous and require community assistance
- Opportunity for training and capacity building, and development of community “champions” to assist with sampling



## KEY LESSONS AND CONCLUSIONS

- Urban water streams represent a rich and highly relevant source of information about exposure to pathogens
  - as well as the opportunity to monitor emerging contaminants, lifestyle indicators, and antimicrobial resistance
- Environmental surveillance can better inform clinicians or public health authorities on disease burden and health service needs in marginalised communities
- Traditional clinical surveillance can be strengthened to include environmental water testing
  - Report to policy makers in real time (district health officials)
  - Report to public health officials on a dashboard
- Samples can be used for wastewater-based epidemiology for a broader scope of pathogens, using the same methods as applied for SARS-CoV-2





## CHALLENGES

- How best to start collaborations, and enter these marginalised communities?
- How do we get buy-in and support from the community
- How do we share information with the communities?
- Public health policies; How do we inform clinicians?

# THANK YOU





# Ensuring **Health Equity**: Infectious Disease **Surveillance** in SouthEast Asia through **Non-sewered** **Wastewater**

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Department of Infectious Diseases and Infection Control, Yamagata Prefectural Central Hospital Yamagata, Japan

Centre of Excellence in Antimicrobial Resistance and Stewardship, Faculty of Medicine, King Chulalongkorn Memorial Hospital, Chulalongkorn University, Bangkok, Thailand



山形県立中央病院  
YAMAGATA PREFECTURAL  
CENTRAL HOSPITAL



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จุฬาลงกรณ์มหาวิทยาลัย  
Pillar of the Kingdom



CHARLES  
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Yamagata Prefectural University  
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The University  
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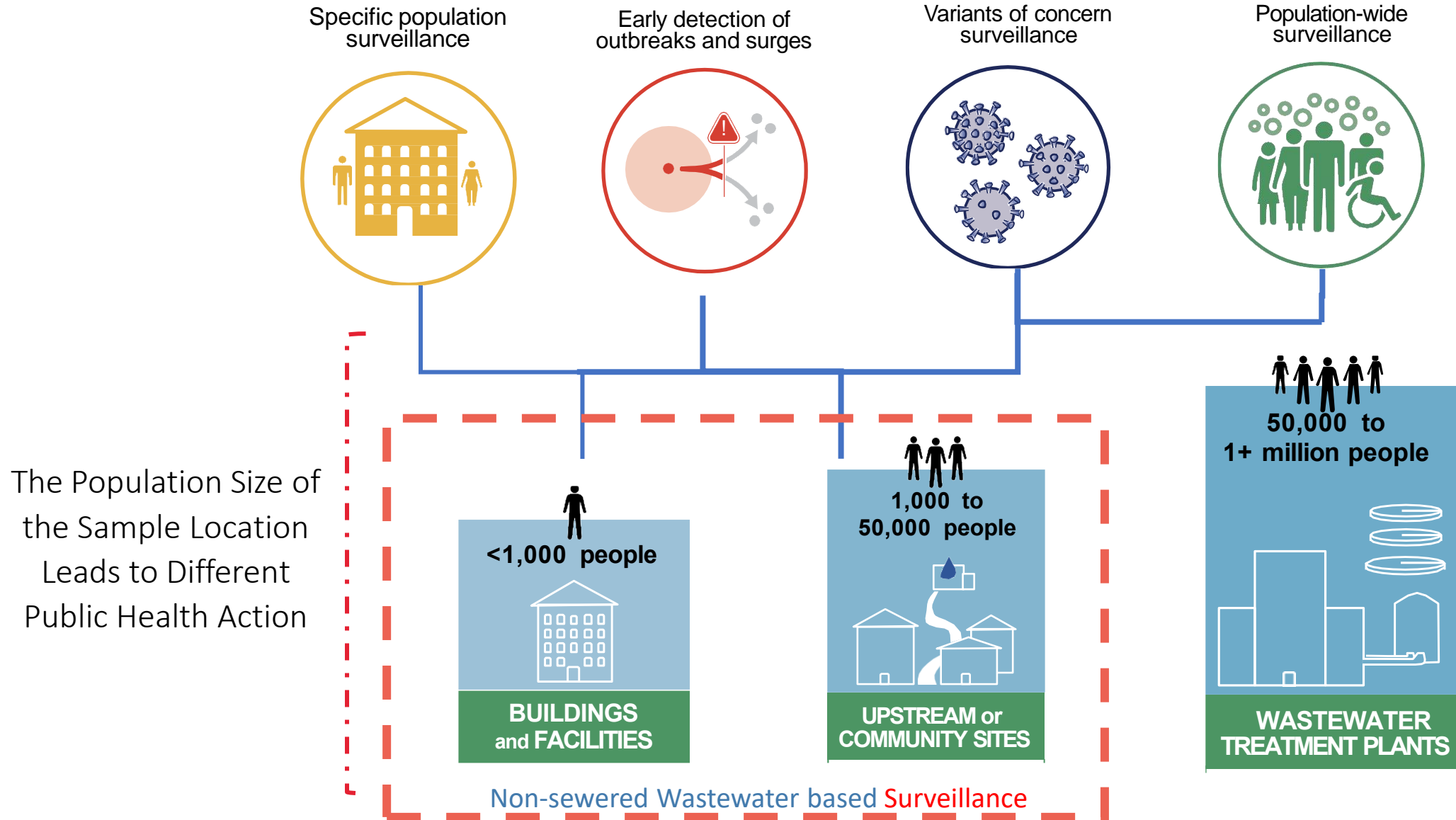
東北大学  
TOHOKU UNIVERSITY



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Discover. Prevent. Cure.

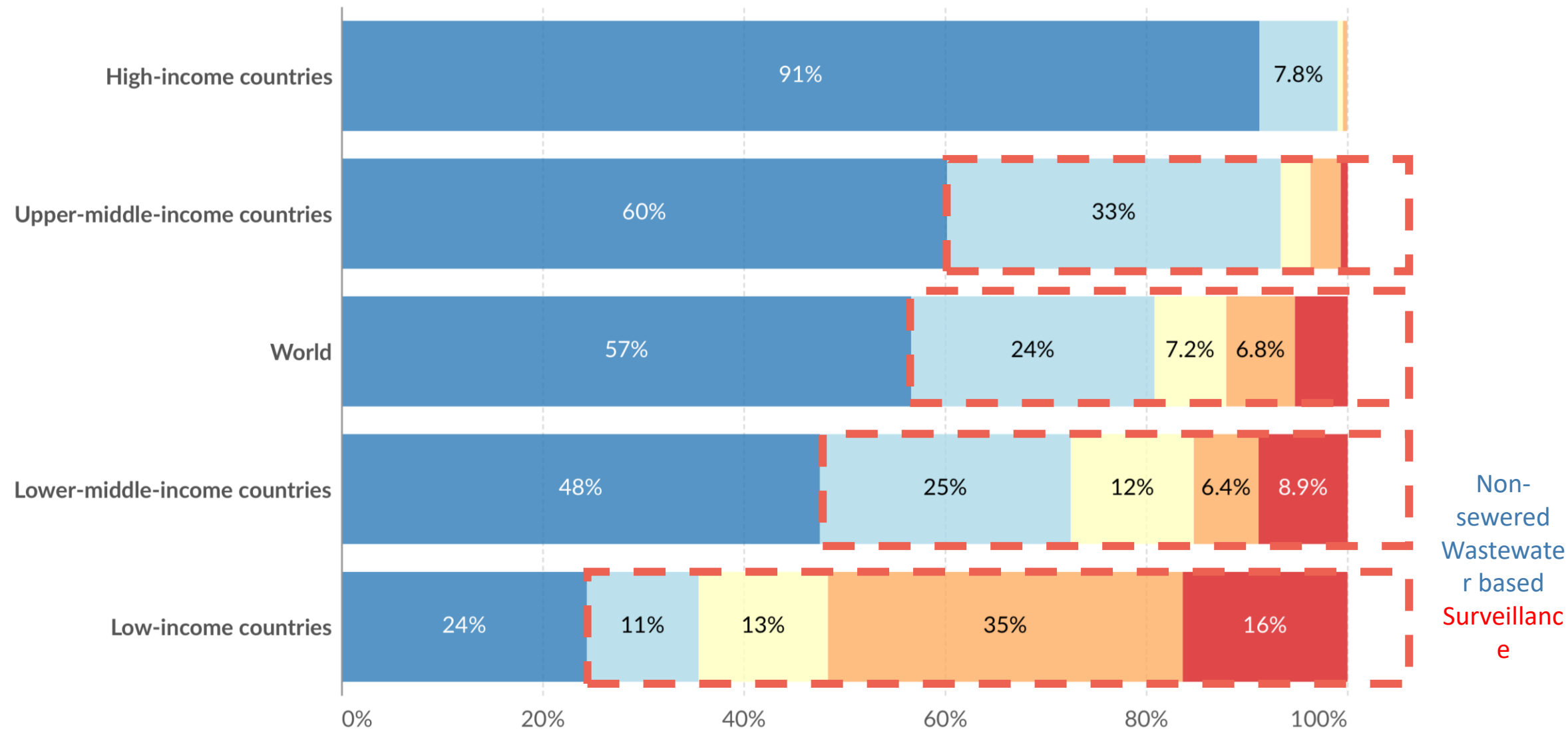


# Four Uses of Wastewater Testing results

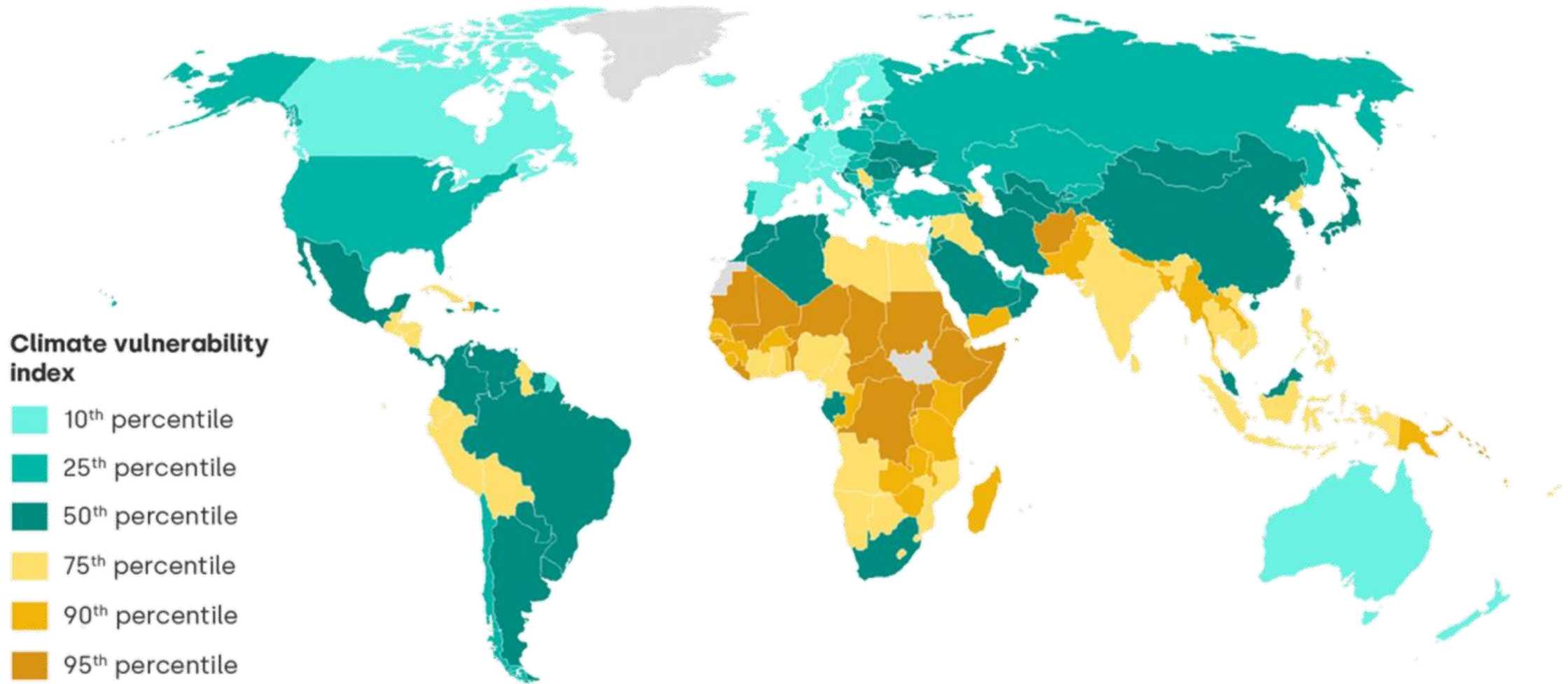


# Share of the population using sanitation facilities

■ Safely managed   ■ Basic   ■ Limited   ■ Unimproved   ■ Open defecation



## Regions in sub-Saharan Africa and South Asia are among the most vulnerable to climate change



Source: Data from ND-GAIN (<https://gain.nd.edu/our-work/country-index/>)



Despite the **potential** for wastewater testing to include **populations** that are not monitored through **individual testing** or **sewered testing**, there are health **inequalities** that must be addressed in low and middle-income countries

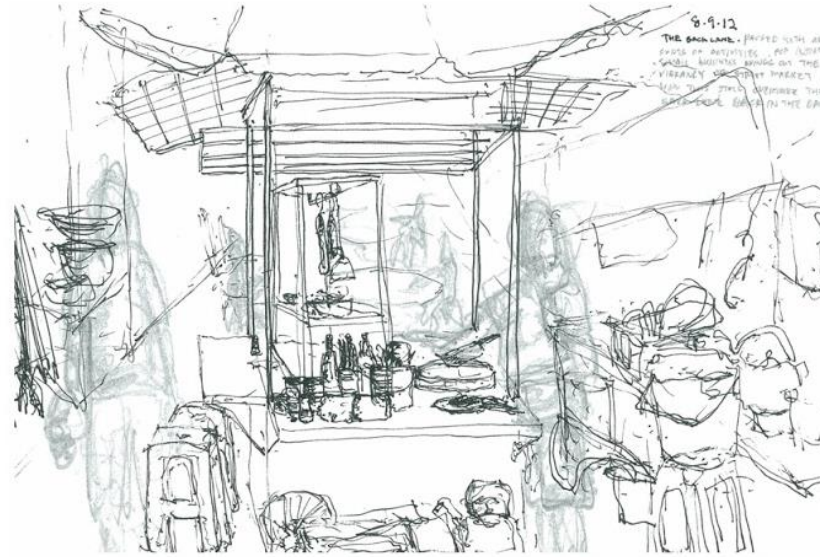




# Identifies new clusters and outbreaks, which then triggers investigation and action through **Non-sewered Wastewater based Surveillance**



Community houses

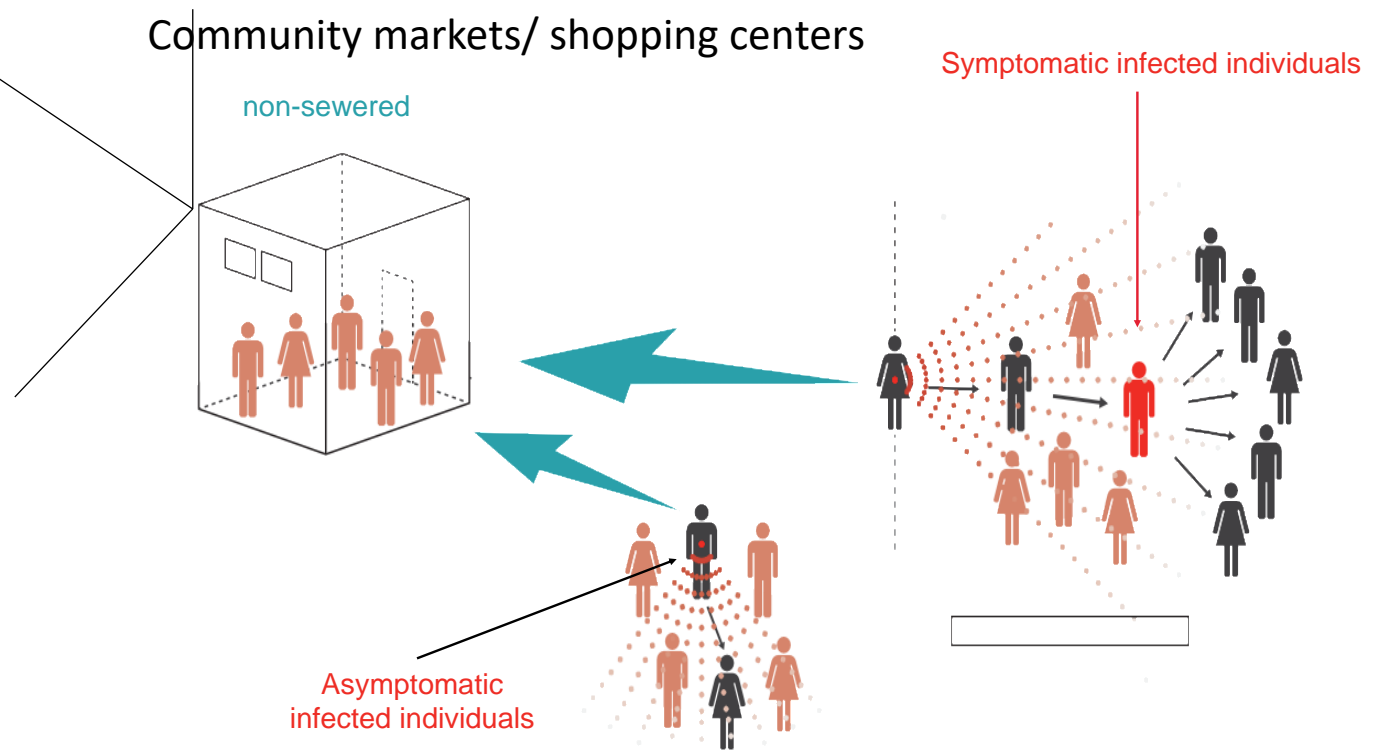


Community markets/ shopping centers

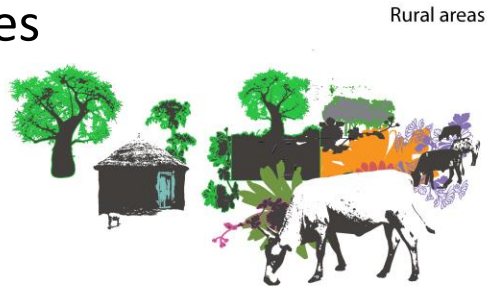


sketch by TAN SOON CHERN

Entertainment venues



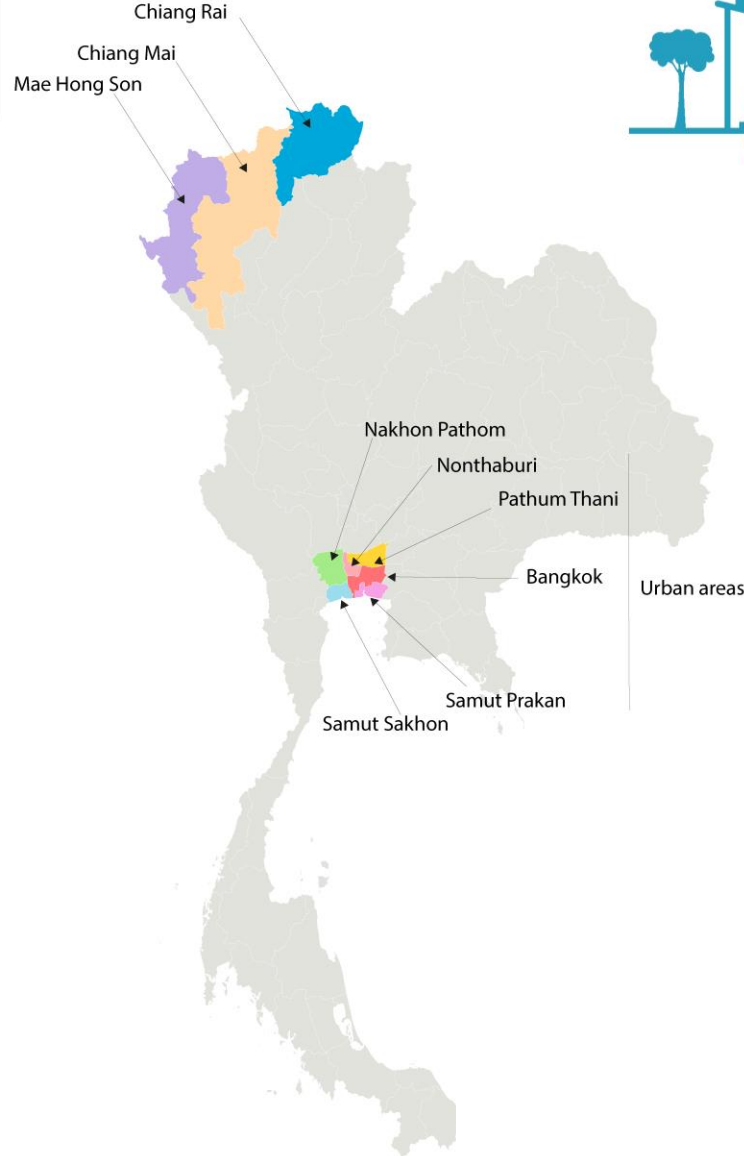
# Participatory infectious disease surveillance through **non-sewered** wastewater in low and middle -income countries



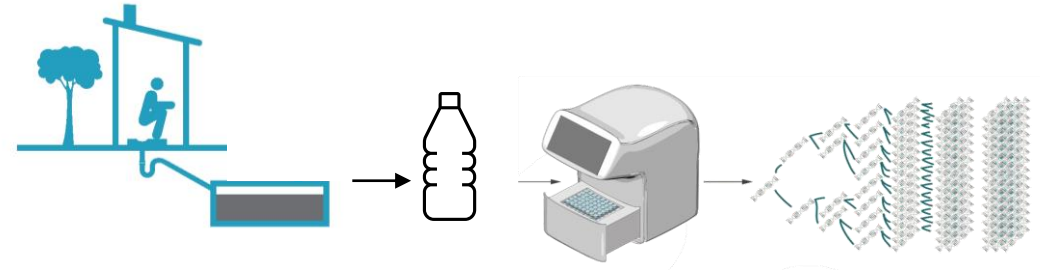
Rural areas

- Community/food markets
- Community malls

In total, 1512 samples were collected from Bangkok, 2976 from five adjacent provinces, and 744 from rural provinces between July 2020 and August 2021.



Wastewater sampling provinces in Thailand



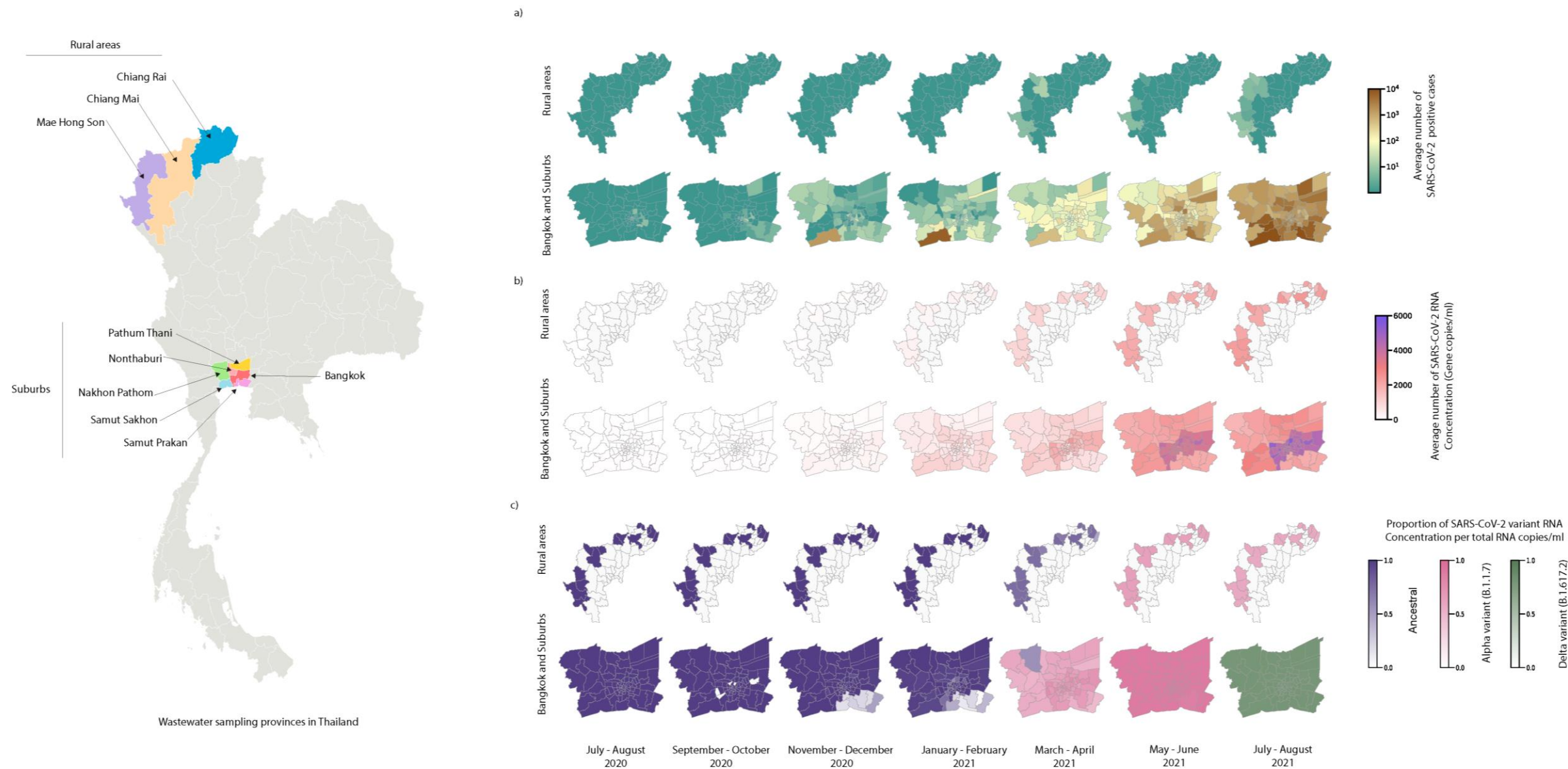
Standard grab sampling method followed by molecular detection



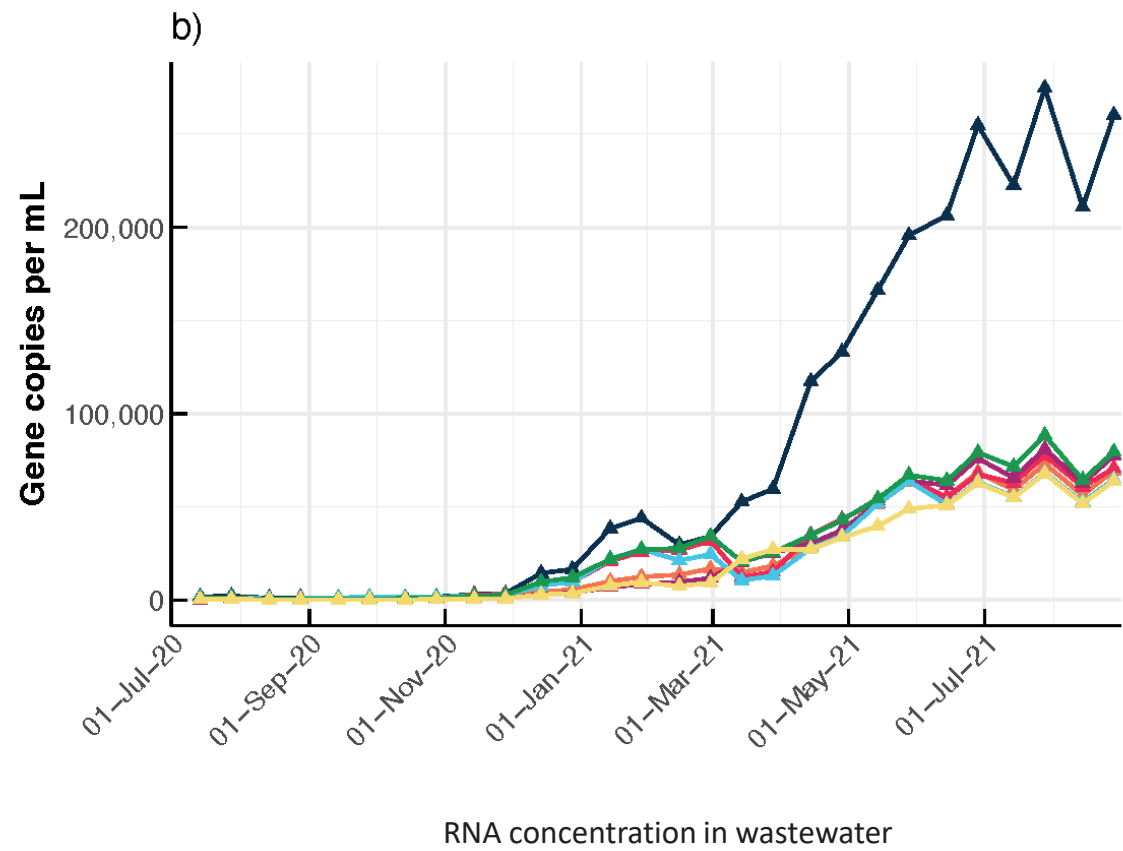
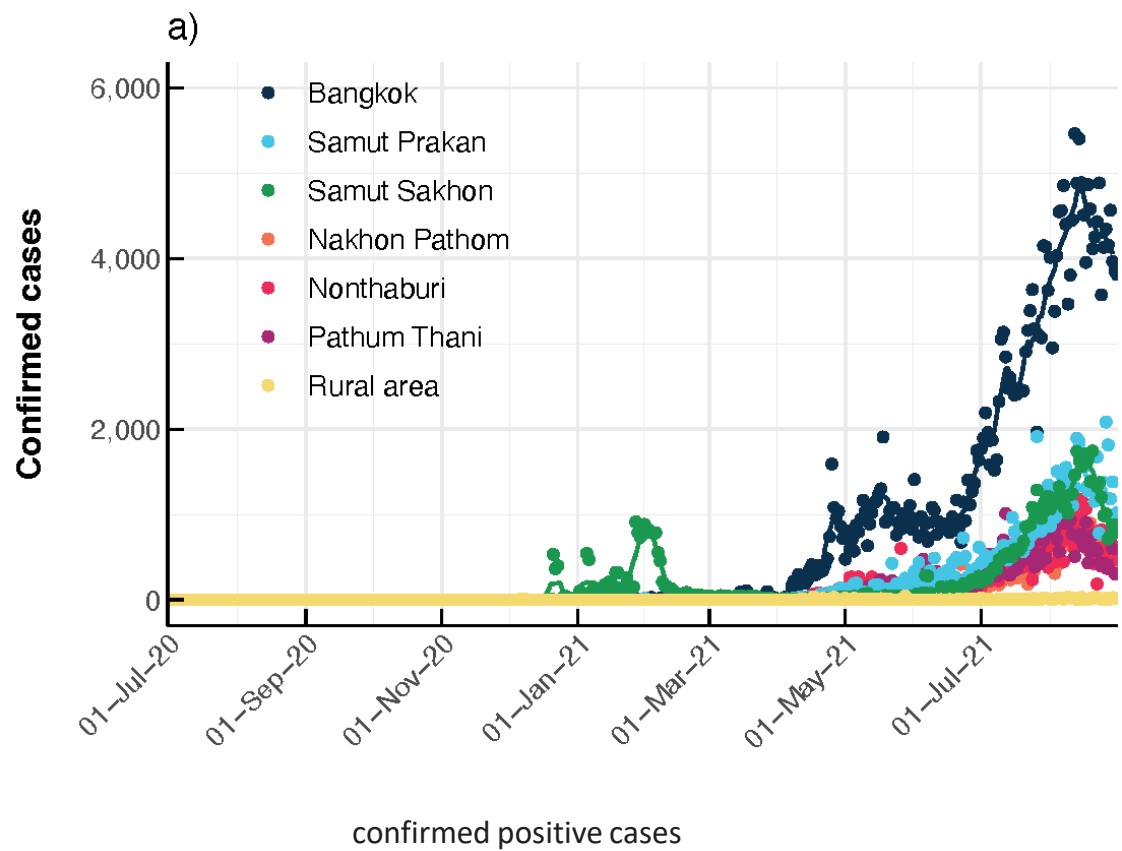
- Condominium complexes
- Cafeteria and shopping centers
- Community/food markets
- Office complexes
- Wastewater treatment plants
- Entertainment/leisure venues
- Work sites (construction camps)
- Housing complexes



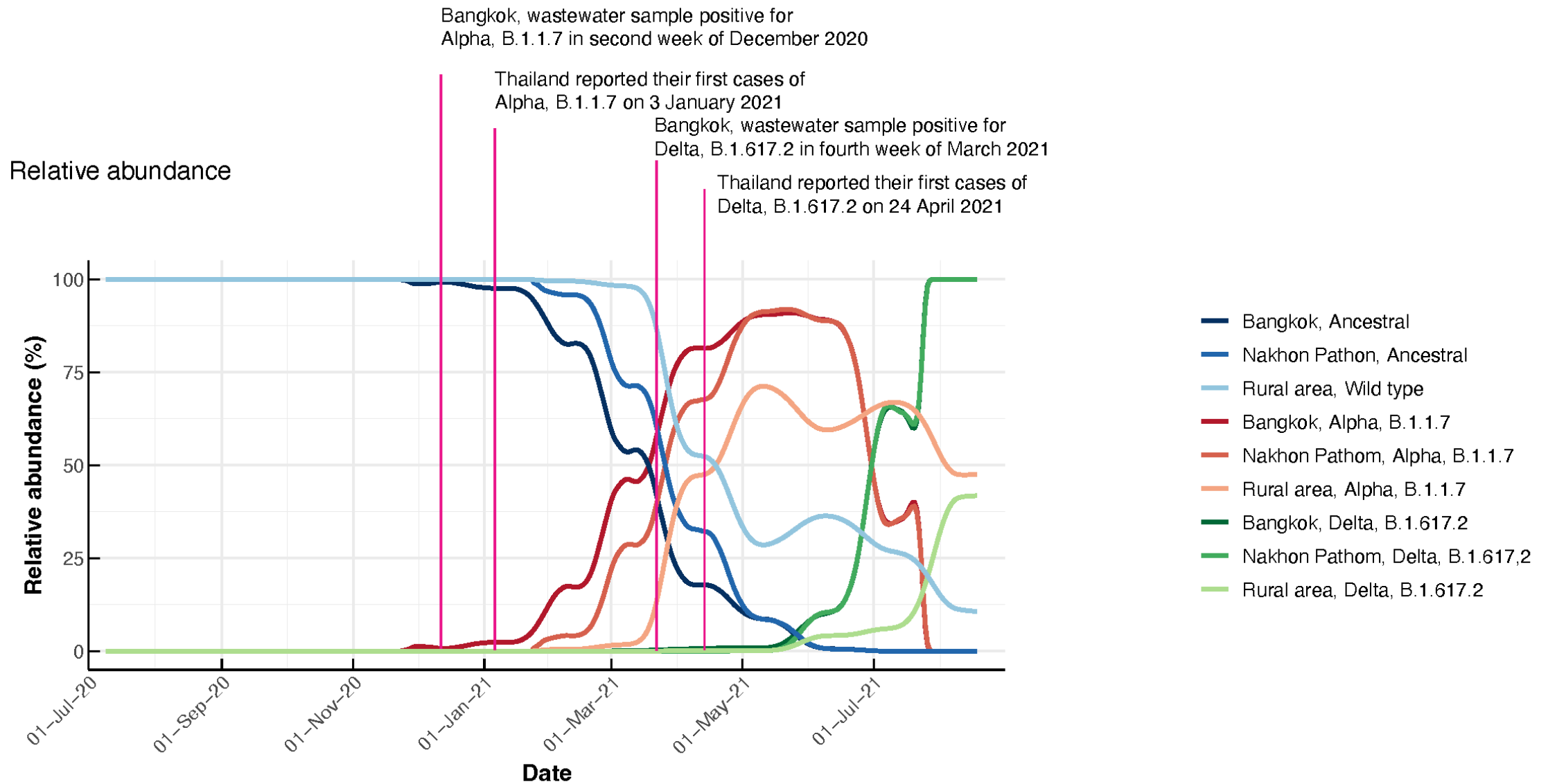
Starting in November 2020, SARS-CoV-2 RNA concentrations in wastewater increased gradually, aligned with the increasing trend in daily newly reported cases in Bangkok and five surrounding provinces. Rural areas followed the same direction, even though SARS-CoV-2 RNA concentration remained relatively low compared to Bangkok and the five surrounding provinces



An increase in the estimated incidences based on wastewater SARS-CoV-2 RNA concentrations was observed before infected individuals were officially reported

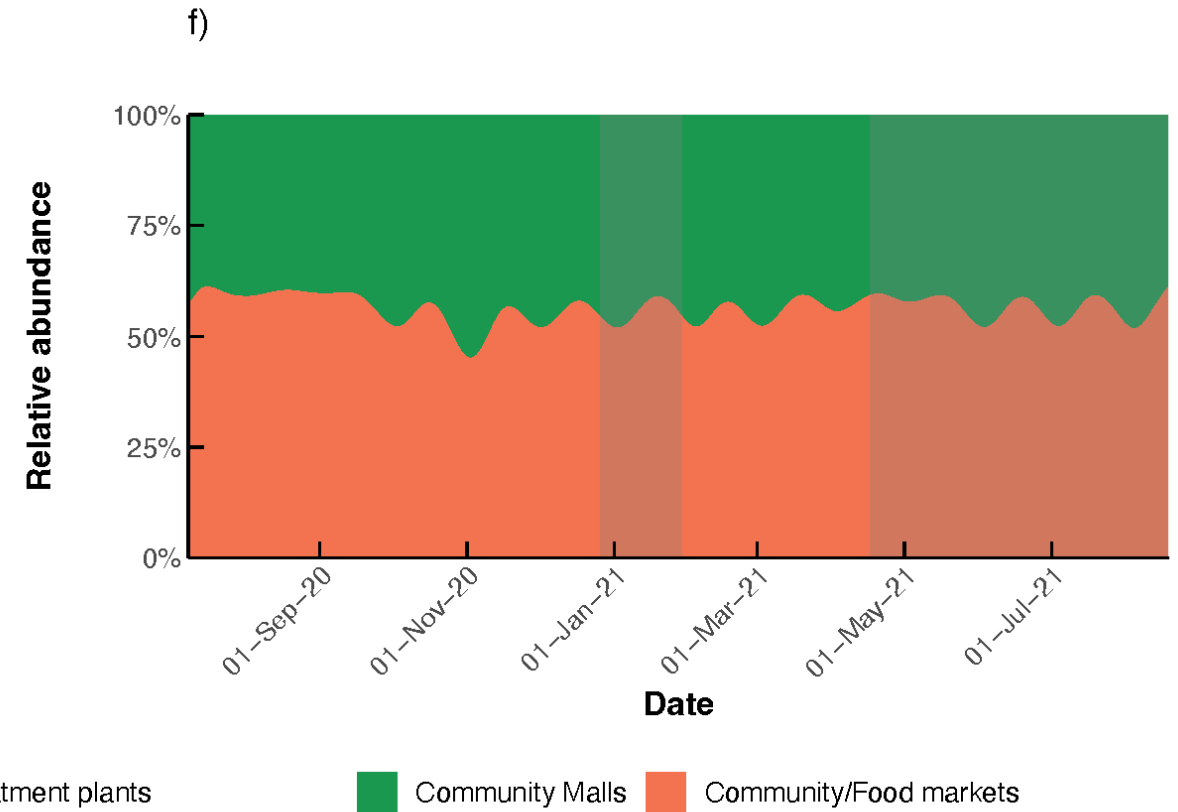
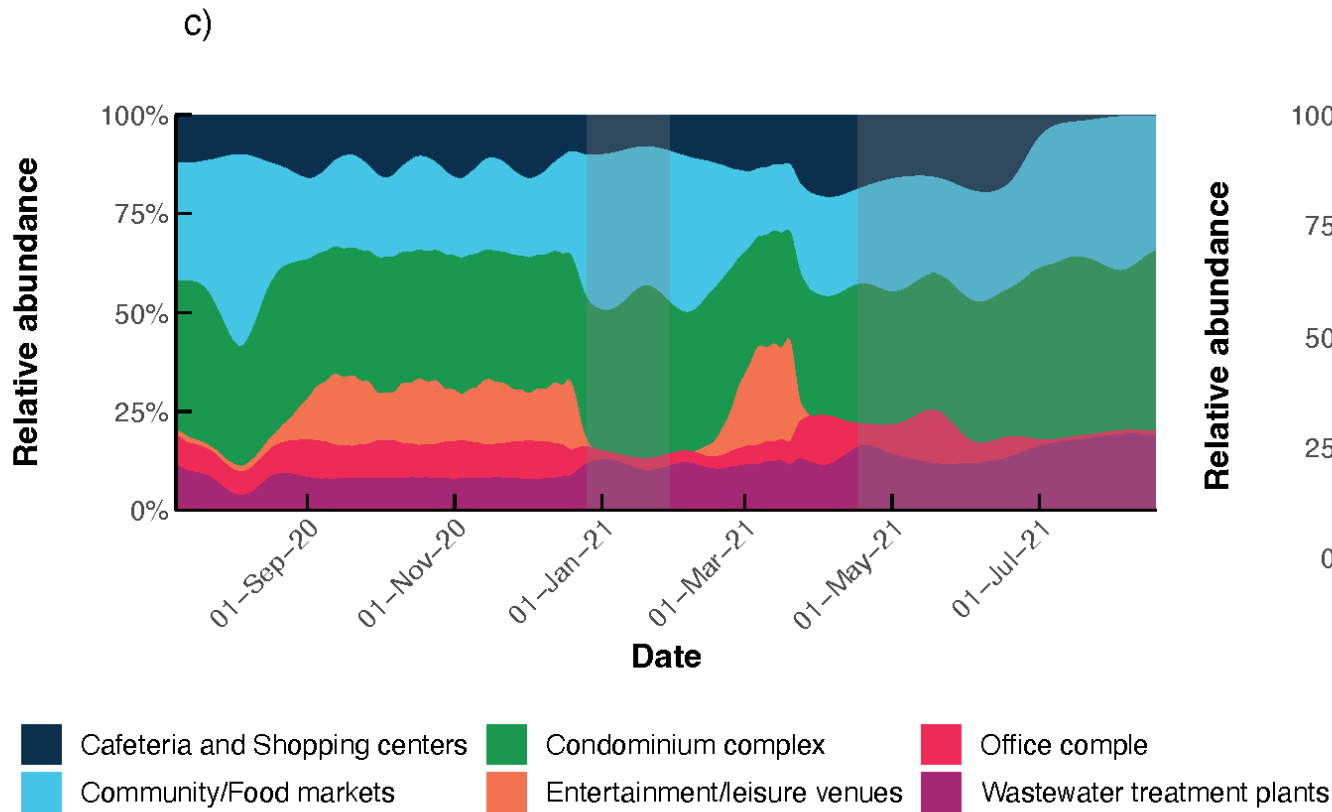


Wastewater-based estimated incidence showed a steep increase starting from the last week of November 2020. However, case-based incidences remained low until April 2021



The relative abundance of different SARS-CoV-2 variants in wastewater revealed that the **ancestral SARS-CoV-2** was dominant until early March of 2021 in Bangkok and the five surrounding provinces and remained dominant in **rural areas** until mid-April 2021





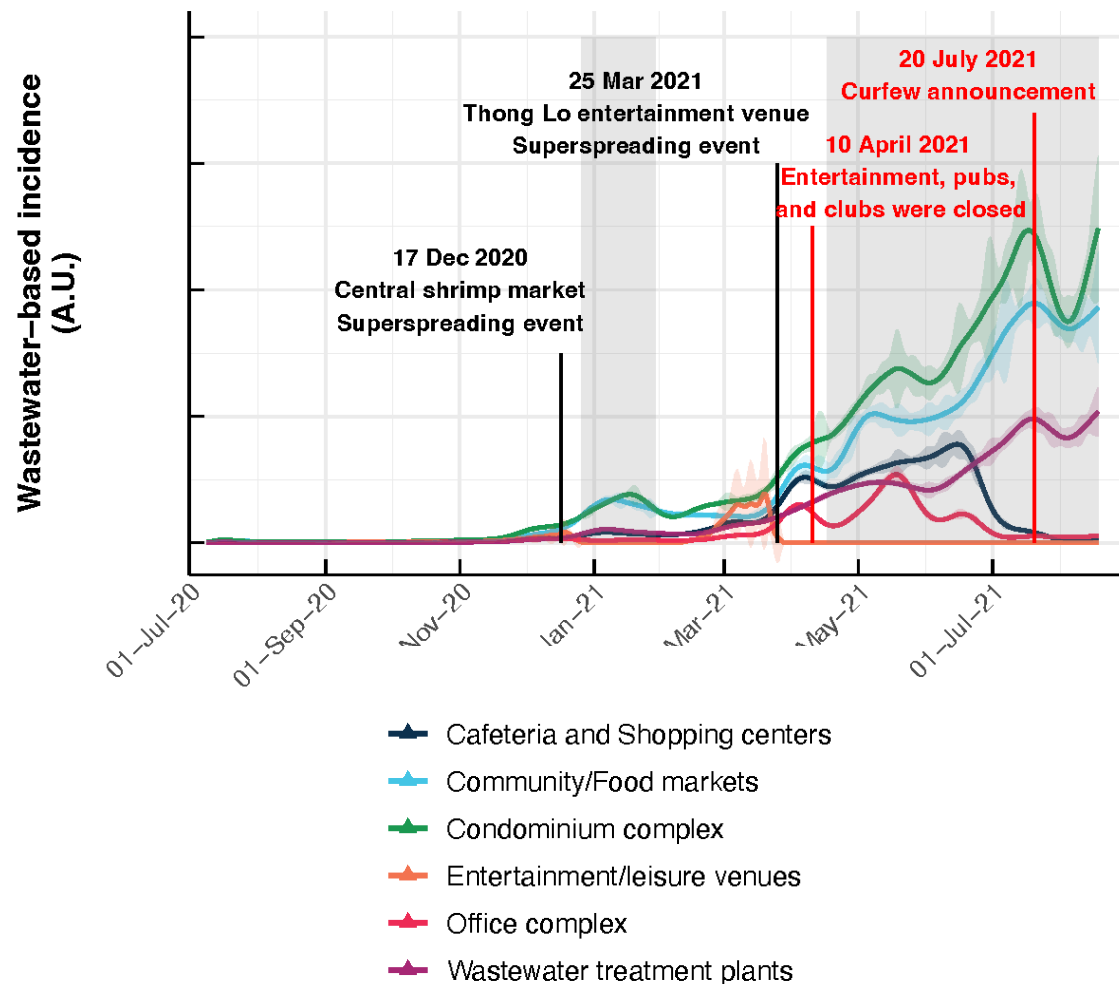
Analysis of SARS-CoV-2 viral RNA concentrations in different facilities revealed that the relative abundances of SARS-CoV-2 viral RNA concentrations in wastewater were relatively high in community/food markets and condominium complexes in Urban areas.

In rural regions, community malls and community/food markets had similar SARS-CoV-2 viral RNA concentrations in wastewater.

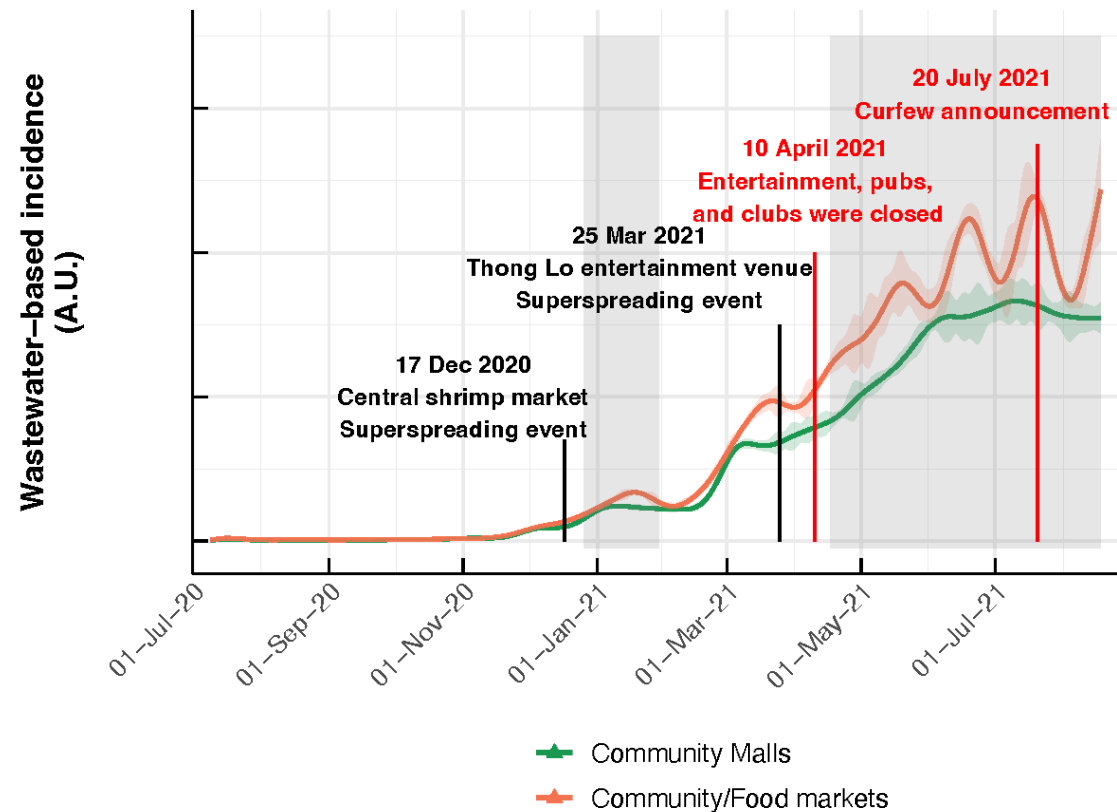
Estimation of infection incidences for each facility showed that **community/food markets and condominium complexes** tend to carry more infected people (symptomatic or asymptomatic) during the

lockdown period in Bangkok

Bangkok



Rural area

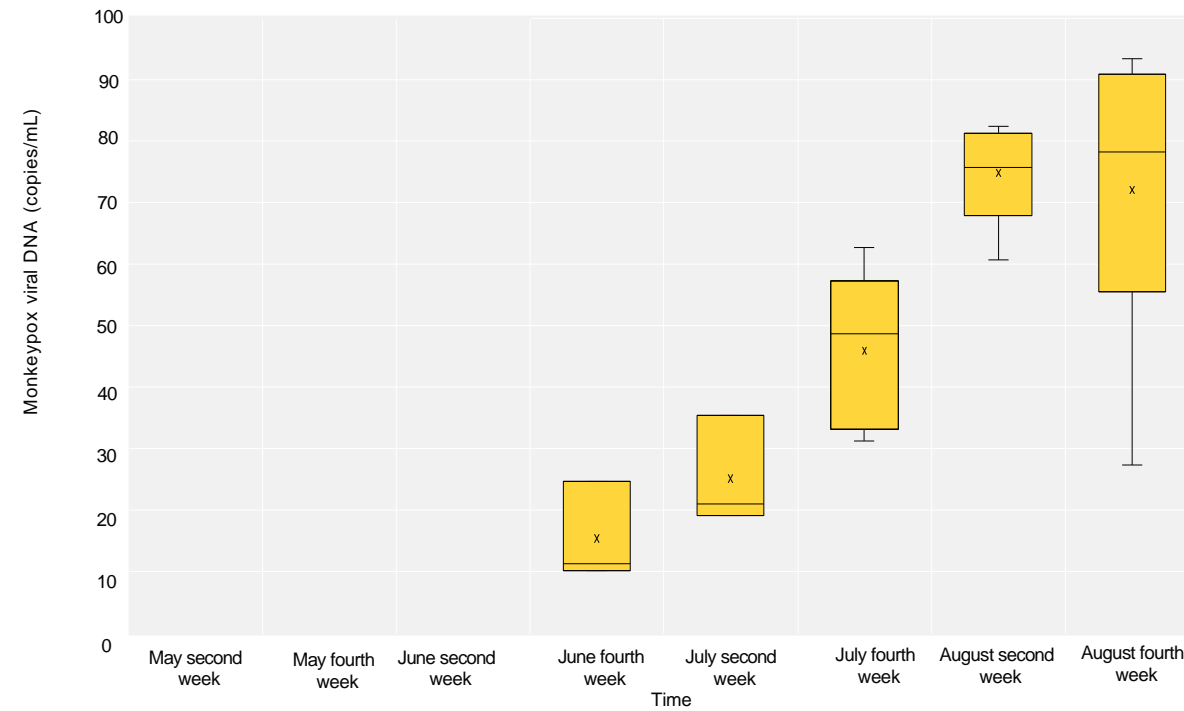


**Community/food markets** had higher estimated infection incidences in **rural regions** than community malls based on viral RNA concentrations in wastewater during the lockdown period

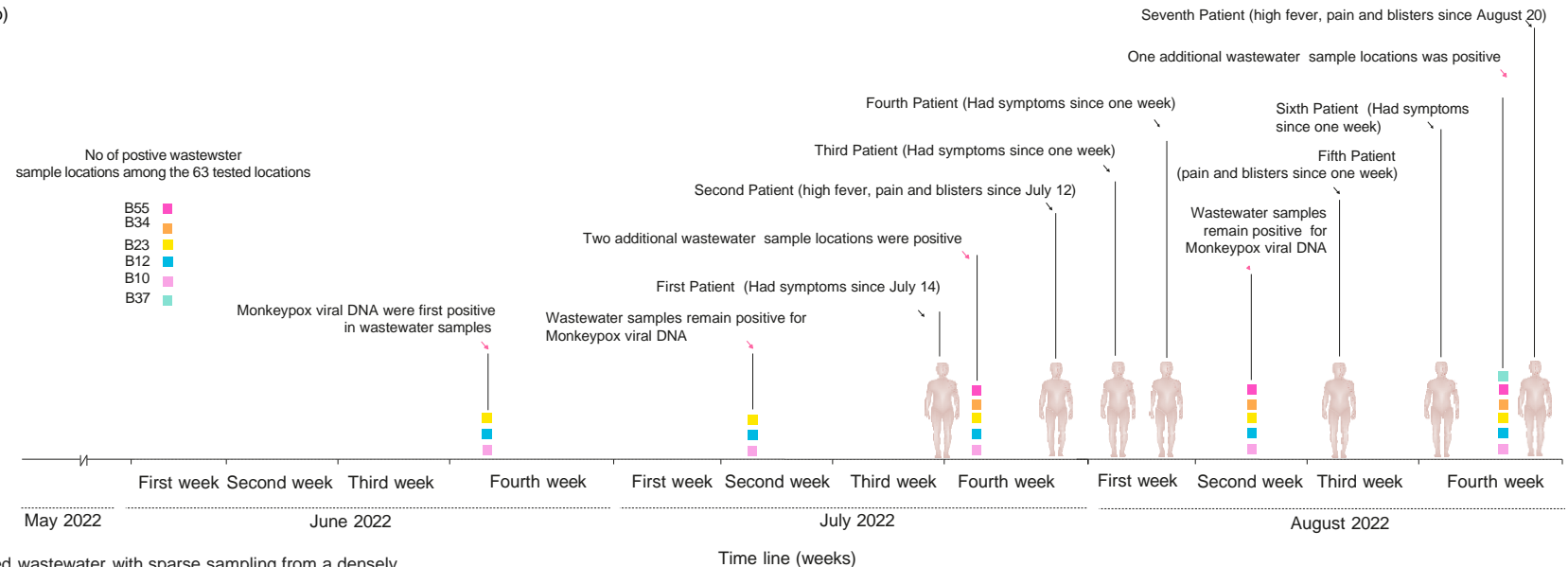
## Multiple traces of monkeypox detected in non-sewered wastewater with sparse sampling from a densely populated metropolitan area in Asia

- First dataset on monkeypox viral DNA detection from wastewater in Thailand
- Monkeypox viral DNA tracing is possible with sparse sampling events
- Non-sewered locations showed positive signals for monkeypox DNA
- Sanger sequencing confirmed the identification of the monkeypox virus.
- Findings expand the use of wastewater surveillance in resource limited countries

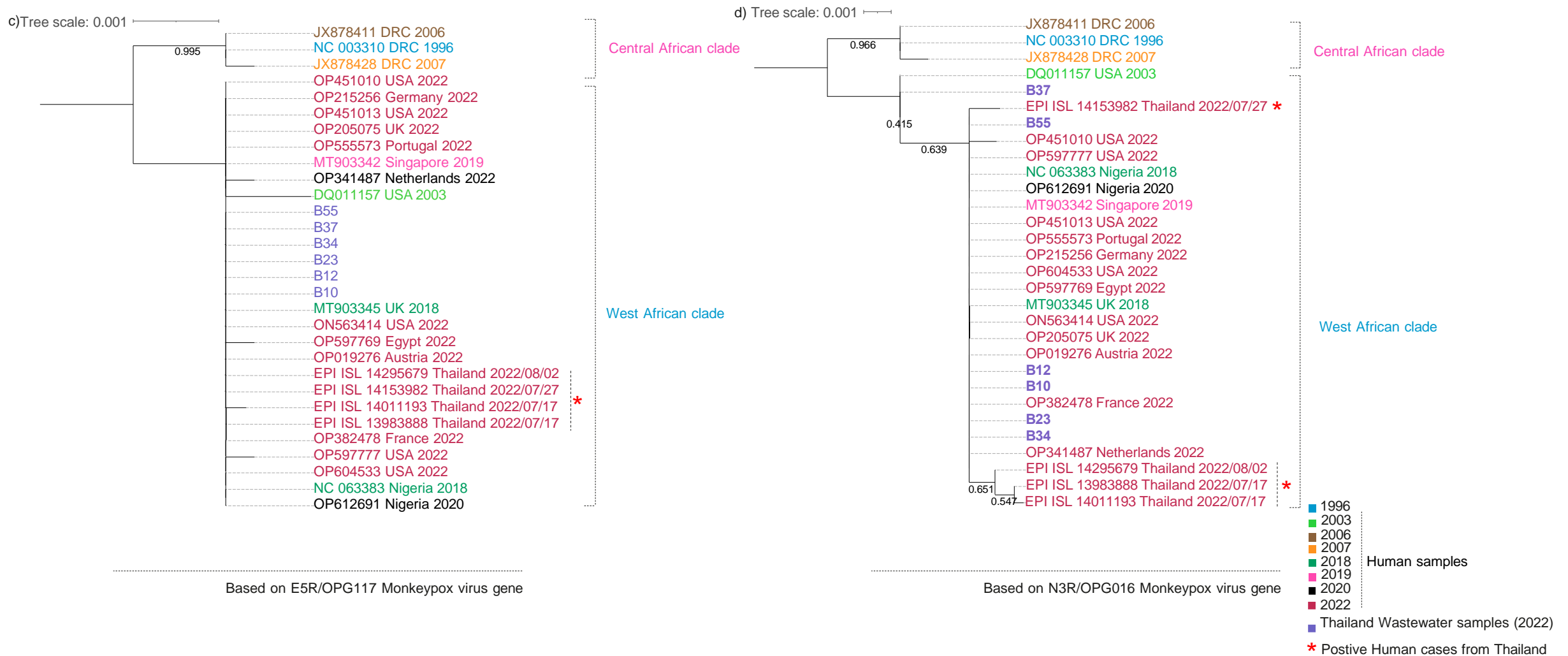
a)



b)







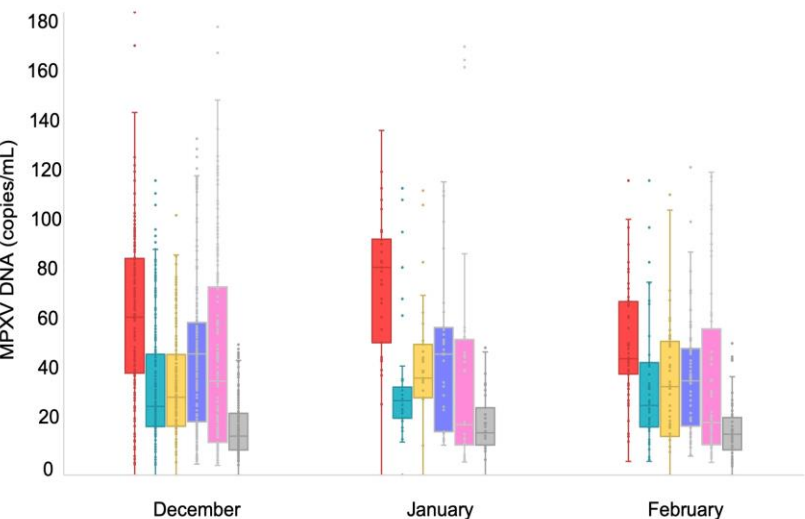
Phylogenetic analysis of the sequence of c) **E5R/ OPG117** and d) **N3R/OPG016 gene** reflect that wastewater samples are

clustering with **positive clinical cases from Thailand** and other geographical origins connected to the **monkeypox 2022 outbreak**

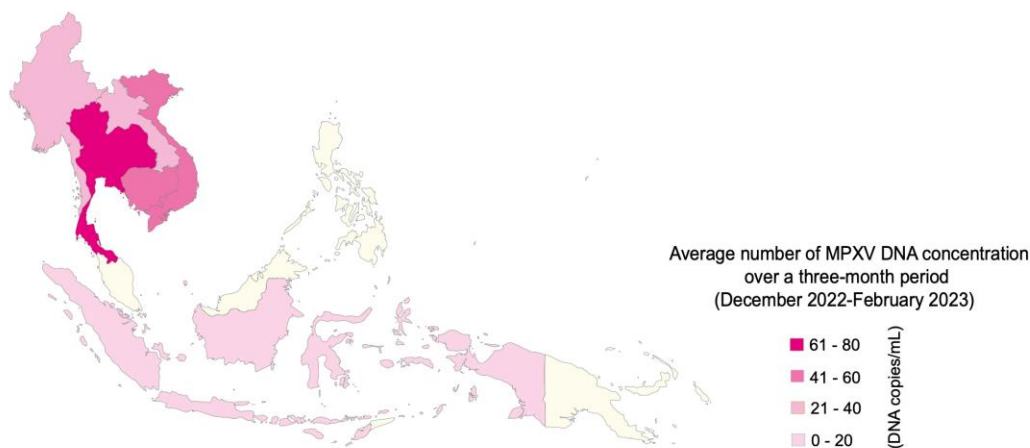
# Non-sewered wastewater surveillance in six Southeast Asian countries revealed positive signals for Monkeypox virus (MPXV)

DNA, indicating local transmission.

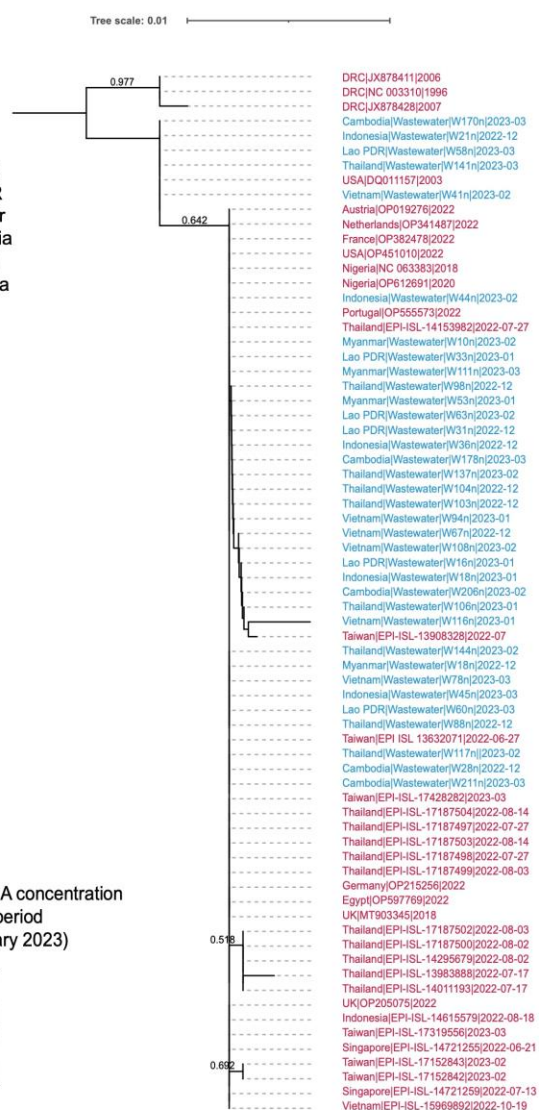
a)



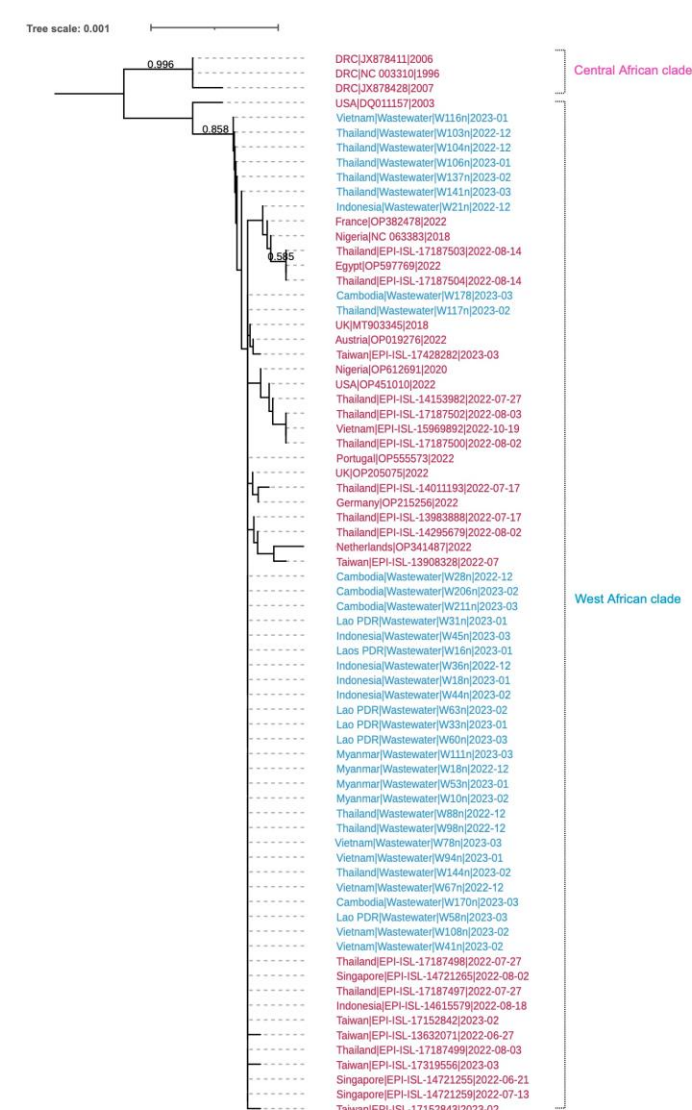
b)



c)

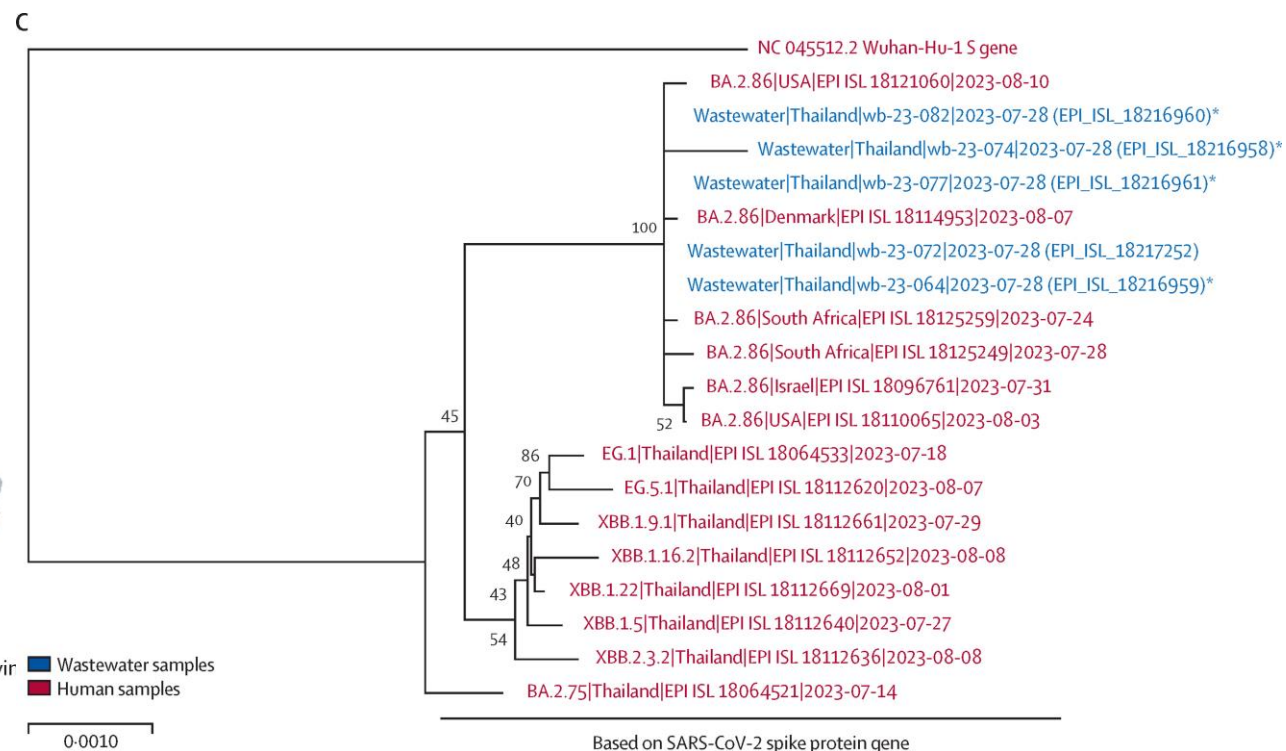
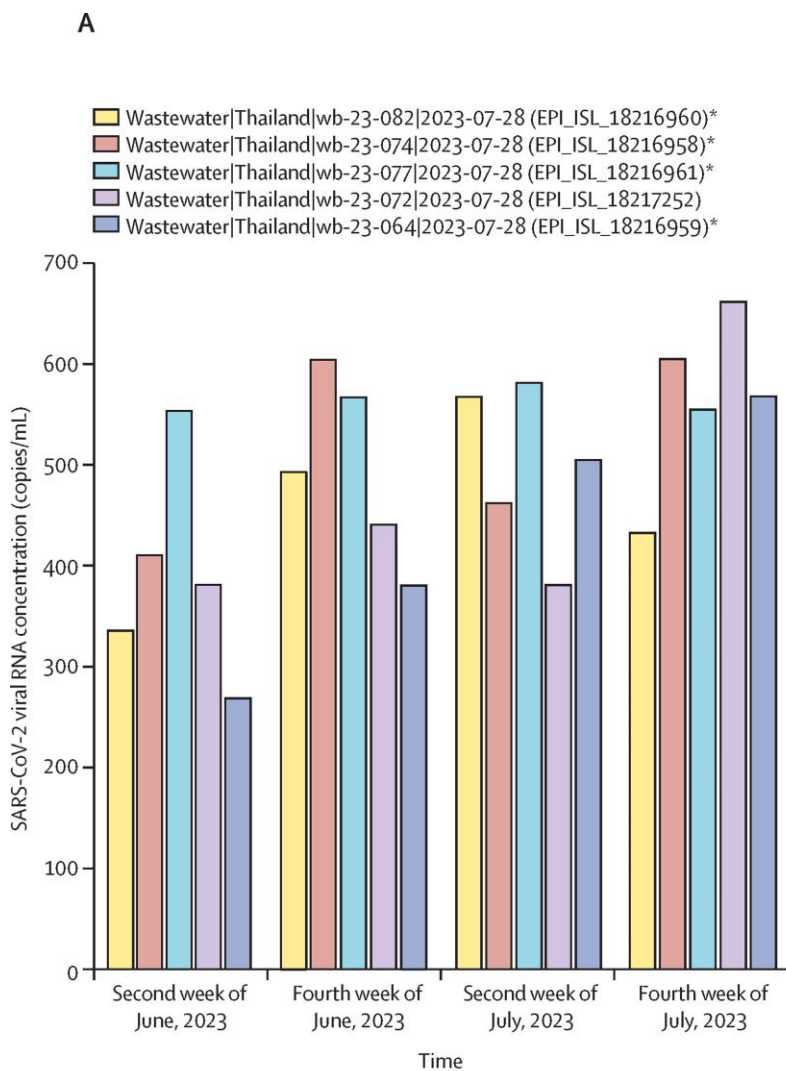


d)



■ Wastewater samples  
■ Human samples

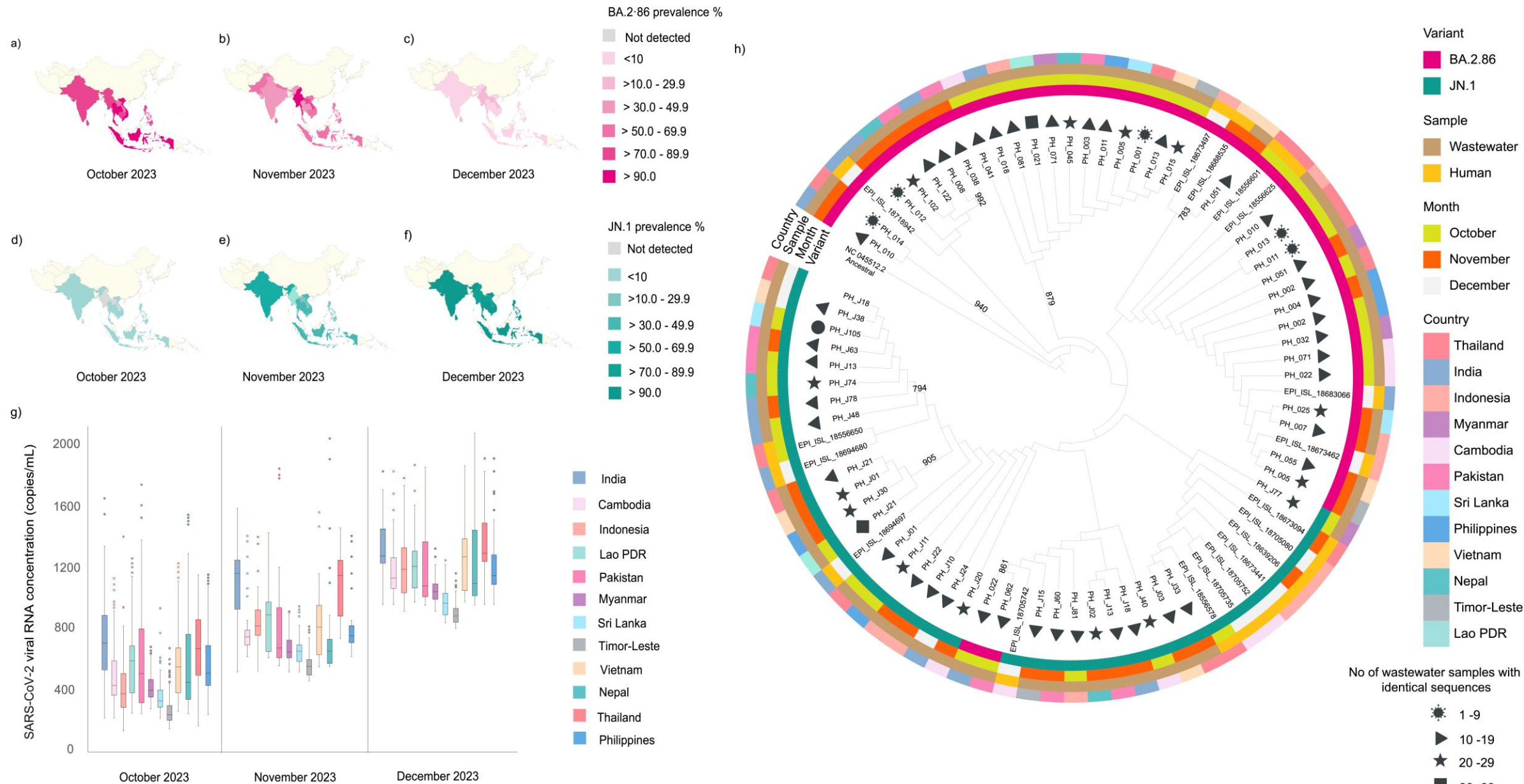
# Tracing the new SARS-CoV-2 variant BA. 2.86 in the community through wastewater surveillance in Bangkok, Thailand.



- Four identical sequences from different locations suggest parallel circulation across multiple areas.
- Furthermore, we identified two different sequences within the same short time frame and small geographical area.
- These sequences were found in different commercial or public venues with non-sewered wastewater



Wastewater-based epidemiological surveillance of SARS-CoV-2 new variants BA.2.86 and offspring JN.1 in South and Southeast Asia through non-sewered wastewater



# Summary

- Participatory infectious disease surveillance through Non-sewered Wastewater can provide actionable information in more a clinically relevant timeframe
- Non-sewered Wastewater showed the distribution of infectious disease across different facilities
- We provide an equitable approach to wastewater monitoring with sparse sampling
- Also proven useful for assessing risk factors, vaccine effectiveness, and patterns of healthcare utilization while being less expensive, more flexible, and more scalable than traditional systems.
- Participatory infectious disease surveillance through Non-sewered Wastewater provides unique disease information that is not available through traditional surveillance sources
- Findings advance health equity in low-resource countries with poor sewer system



Thanks to all the members and collaborators



山形県立中央病院

Yamagata Prefectural Central Hospital



CHULALONGKOR UNIVERSITY



THE UNIVERSITY of WESTERN AUSTRALIA



QIMR Berghofer Medical Research Institute  
THE FUTURE OF HEALTH



We thank all the volunteers, LGBTQIA+ community and marginalized, vulnerable indigenous communities who kindly supported with sample collection. Our team comprises individuals who self-identify as underrepresented ethnic minorities, gender minorities, members of the **LGBTIQOO+1 Ally** community and individuals living with disabilities.

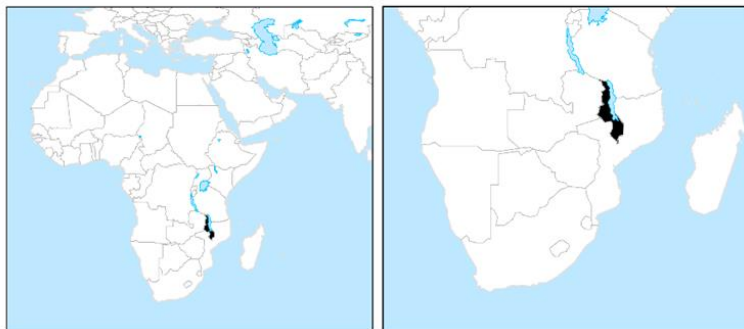
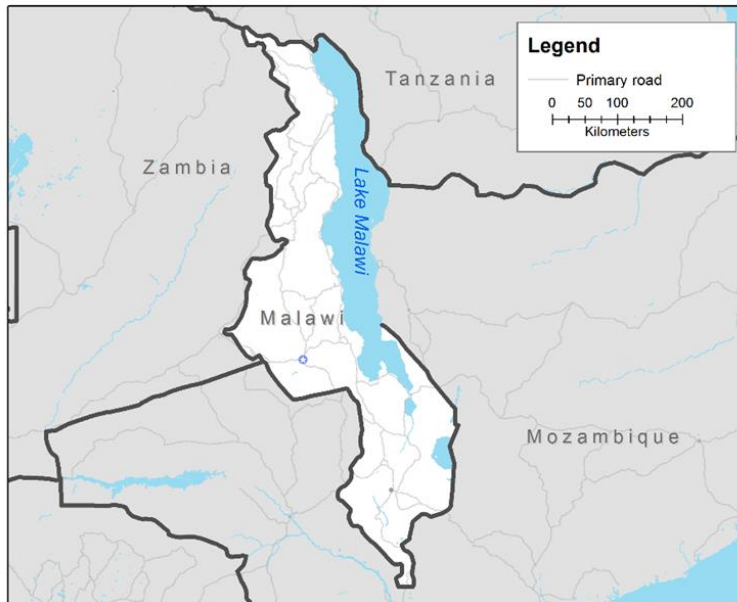


# Addressing Challenges for Establishing a Non-Sewered Wastewater Surveillance system in Malawi

PETROS CHIGWECHOKHA, MALAWI UNIVERSITY OF SCIENCE AND TECHNOLOGY (MUST)



# SANITATION PROFILE FOR MALAWI



Key Area	Stats/Burden
Diarrheal disease burden	824 foodborne DALY per 100,000
Household drinking water (basic)	70% of the population
Household sanitation facilities (safely managed)	24% of the population
Household hygiene (basic)	8% of the population
School drinking water (basic)	78% of schools
School sanitation facilities (basic)	66% of schools
Health care facilities water (basic)	78% of facilities
Health care facilities sanitation (basic)	3% of facilities
Literacy levels	67.31%
Unemployment rate	18.5%

❑ **SS Africa remains behind - 7% sewered sanitation against 43% global status.**

# CONTEXT

Low WBE  
surveillance

Most LMICs remain  
behind on  
wastewater-based  
pathogen  
surveillance and  
detection

92% of MW relies on on-site  
and NS sanitation systems.



- ☐ Fecal sludge management problems
- ☐ Pit emptying services unavailable/ expensive



- ☐ The pits are often abandoned once full

Utilization of WBE to monitor  
community health status difficult/  
impossible.

**Public Health  
monitoring  
Challenges**

Dependency on clinical data for  
community disease monitoring, often  
unavailable and unreliable



# NSS WASTEWATER SAMPLING – NOT EASY

On-site sanitation sampling is a daunting task as opposed to centralized treatment plants

- ❑ NS sampling not a luxury – requires dedication
- ❑ Let us honor the tireless efforts of those who brave these challenging environments.



# POLICY LEVEL CHALLENGES

Policy challenges are often related to the nascency of WBE

Limited commitment from governments

Lack of awareness by policy makers



WHY??

Associated with the fact that WBE is a relatively new in most LMICs

Need promote awareness among relevant government authorities

# TECHNICAL CHALLENGES IN SETTING-UP NS SURVEILLANCE



Sampling Frame and  
protocol not easy



## POSSIBLE SOLUTION

Need for fully functional  
lab for wastewater-based  
surveillance



Limited Equipment  
Lack of BSL-2 and RT-qPCR



## POSSIBLE SOLUTION

North-south partnerships  
and collaborations



Limited Human Capacity  
(Lab and Bioinformatics)



## POSSIBLE SOLUTION

Embark on Training capacity  
building programs  
Well trained workforce is  
critical



Unavailability of well  
documented clinical data



## POSSIBLE SOLUTION

Governments needs to  
strengthen data digitization



# LOGISTICAL AND PROTOCOL CHALLENGES

## Supply of Consumables

Most reagents not available in  
African countries



## Supply chain bottlenecks



## Ethical considerations

IRB approval and site  
approval takes time and  
require good planning



## Operation and maintenance of laboratory



Problems with access to foreign currency  
for purchase of supplies



# CURRENT EFFORTS

## Advancing the Use of Fecal Sludge for Timelier and Better-Quality Epidemiological Data in Low- and Middle-Income Countries for Pandemic Prevention

Petros Chigwechokha, Renée Street, and Rochelle H. Holm\*

Cite This: <https://doi.org/10.1021/acs.est.2c07788>

Read Online

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Metrics & More

Article Recommendations

**KEYWORDS:** COVID-19, fecal sludge, low- and middle-income countries, pathogen, wastewater-based epidemiology

The availability and accessibility of water, sanitation, and hygiene (WASH) services are key in preventing disease transmission.<sup>1</sup> Monitoring non-infectious severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) RNA in wastewater can serve as an early indicator of changes in coronavirus disease (COVID-19) cases in a particular catchment area.<sup>2</sup> Fecal sludge is excreta held on site (unsewered) in sanitation systems, such as at households, in shared outhouses, and in septic tanks. These on-site sanitation systems, a positive step toward improved household WASH services, are used by 43% of the global population, mainly comprising low- and middle-income countries (LMICs).<sup>1,3</sup> There is an urgent need to expand decentralized (non-sewered system) surveillance as LMICs are currently being overlooked in pandemic prevention as it is currently focused on SARS-CoV-2 wastewater from centralized (sewered) systems.

Wastewater surveillance of centralized systems presents a vastly distinct set of challenges compared with fecal sludge surveillance. For example, sewers also contain gray water and sometimes stormwater, and wastewater is commonly sampled as a composite volume over a 24 h period, whereas fecal sludge is most commonly grab sampled. Broadening the focus of non-sewered sanitation surveillance (NSSS) should accommodate fecal sludge analysis to support globally relevant public health strategies.

Even before the COVID-19 pandemic, fecal sludge was used as a valuable source of information for understanding community health.<sup>4</sup> Unfortunately, interest in fecal sludge research has been largely limited to a small group of researchers funded by a few core donors. The pandemic has renewed the focus of both science and policy. Surveillance of pathogens in fecal sludge may need to be adapted and could include *Vibrio cholerae* or poliovirus; however, this should be determined by assessing localized needs. The potential of the public health impact in LMICs can be amplified by using fecal sludge data in addition to clinical data in multipathogen disease surveillance and continuous local monitoring for early warnings of a pandemic.

Research partnerships are required to progress from individual clinical patient samples (which has been happening

at a limited scale in healthcare settings) to pooled community samples in terms of fecal sludge. The importance of research partnerships is not new,<sup>5</sup> but possibly more complex in the Southern African sanitation field with five decades of human waste research.<sup>6</sup> With no formal global network, international nonprofit institutions can fill this role by bringing together academic research partners and public health officials for interdisciplinary and collaborative research and increased awareness of NSSS. Such partnerships can enable these settings to innovatively adapt academic or healthcare laboratory operations while maintaining high-quality data and also promote peer-reviewed literature originating from resource-limited laboratory settings.

Timelier and better-quality global epidemiological data sets are required for better pandemic preparedness in the future. There is a need in LMICs for the different, but unique, value of NSSS compared to that of wastewater surveillance, built-in capacity at local laboratories that can include fecal sludge as part of continuous multipathogen disease surveillance, and long-term global research collaborations focused on creating systems with data-driven approaches. Challenges will remain, including importation and lengthy delays in supplies and equipment (such as polymerase chain reaction primers and probes that are not manufactured in many LMICs), repurposing existing LMIC laboratory spaces for fecal sludge samples, and limited LMIC access to biosafety level 3 and 4 laboratories for sample processing, thereby limiting pathogen target lists. We recommend that LMICs navigate through these barriers to ensure that fecal sludge surveillance benefits public health, generates local policy-relevant information for future pandemic prevention and control, and holistically supports global WASH services.

Special Issue: Data Science for Advancing Environmental Science, Engineering, and Technology

## Promoting Surveillance in Sub-Saharan Africa: Moving to Wastewater and Environmental Genomic Surveillance Requires More Attention

Rochelle H. Holm\*, Petros Chigwechokha, Craig Kinnear, Anna Winters, and Renée Street

Cite This: *ACS EST Water* 2023, 3, 1994–1996

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Article Recommendations

Laboratory infrastructure and local capacity are essential for surveillance programs to achieve the Sustainable Development Goals. Globally, morbidity and mortality due to unsafe household drinking water, sanitation, and hygiene remain major problems.<sup>1</sup> During the coronavirus disease 2019 (COVID-19) pandemic, more than a few water, sanitation, and hygiene (WASH) researchers pivoted to use polymerase chain reaction (PCR)-based methods to indicate community disease spread from sewer and non-sewered sanitation samples. The transition from WASH (Sustainable Development Goal 6) to wastewater-based epidemiology (WBE; Sustainable Development Goal 3) was a comfortable shift. Physical laboratory infrastructure, equipment, and workforce capacity were quickly adapted to meet public health needs during the pandemic. However, the expansion of genomic surveillance with sanitation samples requires more thought leadership around environmental health research considerations in low-resource settings as the transition is proving to be much more challenging.

In sub-Saharan Africa, there is a workforce gap in the sanitation field. In the case of Malawi, opportunities are available for approximately 17 600 sanitation experts against limited degree program options nationwide.<sup>2</sup> Existing higher-education programs in WASH may cover sample collection and PCR-based methods for sewer and non-sewered sanitation samples to some extent; however, no genomic surveillance higher-education degree programs are available in Malawi. Thus, transitioning to genomic surveillance requires training outside the country at training laboratories like those located in South Africa (such as the South African Medical Research Council Genomics Platform, the KwaZulu-Natal Research Innovation and Sequencing Platform, or the Centre for Epidemic Response and Innovation) or the Africa Centre of Excellence for Genomics of Infectious Disease in West

these surveillance projects should consider a multiyear, dedicated higher-education training period to build individual expertise (master's level and above), not just a short, one-week course prior to the initiation of core wastewater and environmental surveillance. Municipal piped water suppliers, such as in Malawi, are increasingly pivoting their services to include wastewater and often have existing laboratory infrastructure and local capacity. This may allow for a natural long-term extension of moving existing WASH capacity to WBE, which has been neglected in several capacity-building partner discussions.

With regard to WASH access, WBE surveillance activities are better set up for households with sewer connections that would typically be better off in resource-limited regions. While 43% of the world's population has a sewer sanitation system connection, 67% of households in Latin America and the Caribbean region are connected compared with just 7% in sub-Saharan Africa.<sup>3</sup> Any genomic surveillance in sub-Saharan Africa also needs to also consider fecal sludge from multiple households,<sup>4</sup> such as sampling pit latrines and septic tanks in marketplaces, transport hubs, graveyards, healthcare facilities, and surface water (Figure 1). Overall, the population excluded in this surveillance is important, namely the 6% of the world's population still practicing open defecation.<sup>3</sup>

Wastewater and environmental surveillance still require clinical data for calibration, which cannot be performed without strengthening and integrating clinical data systems. However, if broad-spectrum environmental genomic surveillance indicates, for example, enteric disease or antimicrobial resistance genes, the resources required to respond effectively are often unavailable in the existing healthcare systems of resource-limited regions. Furthermore, the opportunity to develop and capacitate sensitive and specific disease detection and response systems geo-enabled to empower targeted and

Need to optimize NSSS protocols to enhance wastewater sample collection, processing and analysis

Need to build local Human capacity in NSSS to enable detection and containment of re-emerging and novel disease threats

Need to build infrastructural capacity that will promote NSSS.

Need to promote research partnerships to unlock funding opportunities

# WASTEWATER-BASED MULTI-PATHOGEN SURVEILLANCE IN MALAWI

## GOAL:

- To build capacity for Wastewater Based surveillance as an early warning tool for various pathogens.
- To understand the relationship between sewage detection and clinical cases

## Objectives

- To determine feasibility of wastewater surveillance in a low-resource laboratory setting using available equipment and existing infrastructure.
- To determine if wastewater surveillance data could be modelled with clinical data for decision making process.
- To pilot a Timelier and better-quality in-country epidemiological data sets for better pandemic preparedness in the future.





# WASTEWATER-BASED MULTI-PATHOGEN SURVEILLANCE

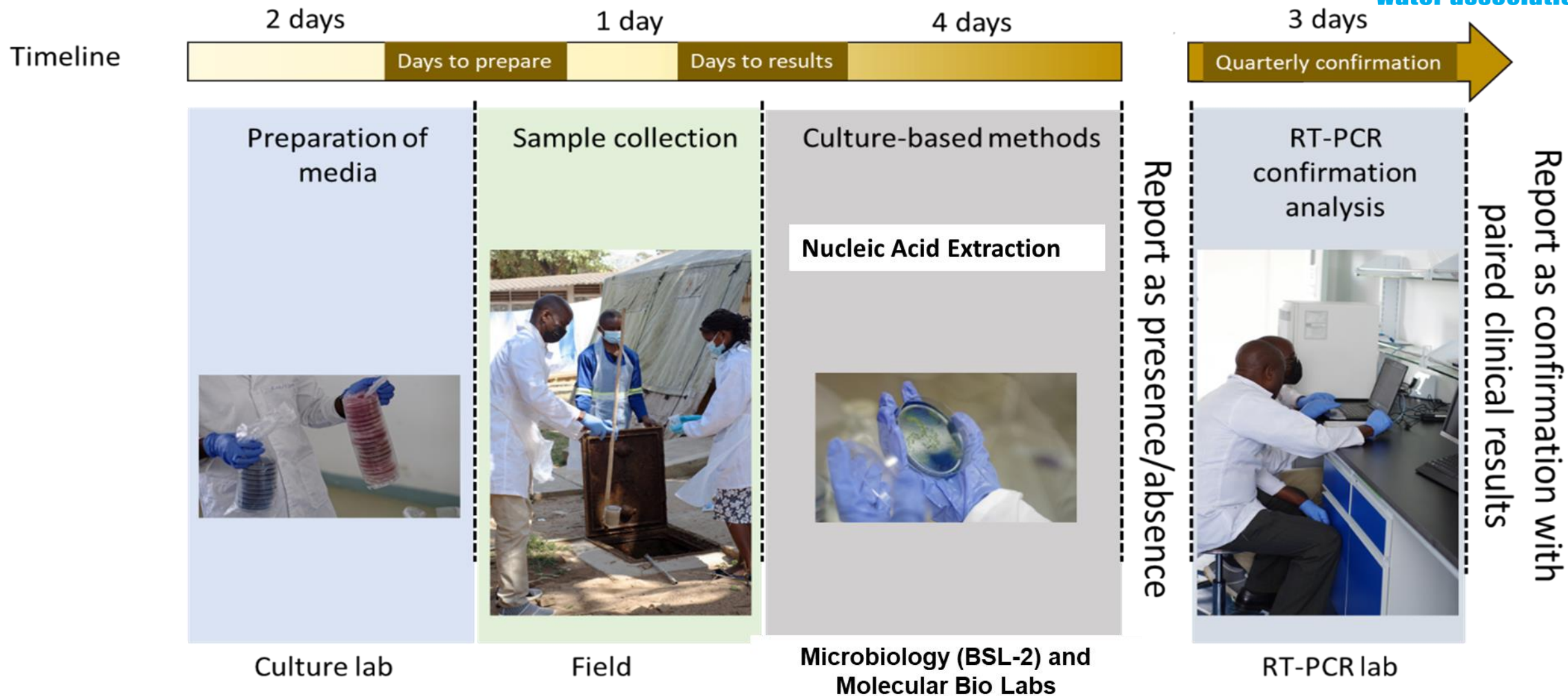
**Current focus** → Five pathogens

- *V. cholerae*
- *S. Typhi*
- Mycobacterium tuberculosis (TB)
- Measles
- SARS – CoV2

- Sample collection – August 2022
- Collection interval - Weekly
- Processing done in BSL-2
- Initial focus – PCR detection
- Switched to culture – Procurement
- PCR later done on isolated for confirmation
- Challenges in market access



# IMPLEMENTATION TIMELINE



## Timeline

1 week

Identify locally relevant fecal shedding pathogens?

1 week

What is current clinical testing approach, including BSL2, 3, or 4?

2 weeks

Obtain ethical clearance for samples, set-up sanitation lab space, define quality monitors

1-12 weeks

Start culture methods

Order PCR supplies

8 weeks

Collect and analyze sanitation samples from sites with known positive clinical cases

2 weeks

Obtain clinical data

2 weeks

Pair clinical and sanitation system data for comparison for proof of concept

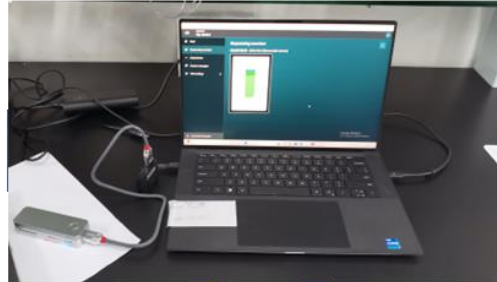
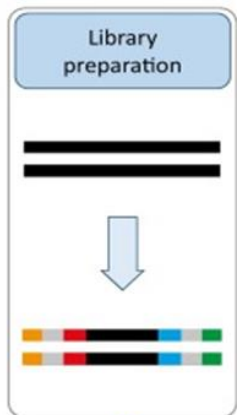


# GENOMICS ANALYSIS

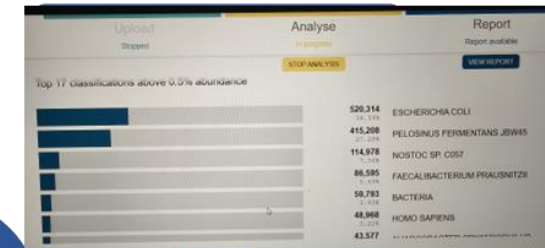
Sample filtration  
and NA extraction



NA Quantification



Sequencing



# FUTURE PLANS



To expand to more sites of sample collection as well as add other pathogens of concern to Malawi



To build a robust metagenomic sequencing capabilities for Wastewater based surveillance.



To develop a dashboard based on our weekly Lab



Train Medical microbiologists in wastewater analysis to increase the capacity in the Lab



Capacity building in Bioinformatics.



Procurement of Robust sequencing platform.

# ACKNOWLEDGEMENTS

## **Malawi University of Science and Technology**

Ruth Nyirenda

Chimwemwe Tandwe

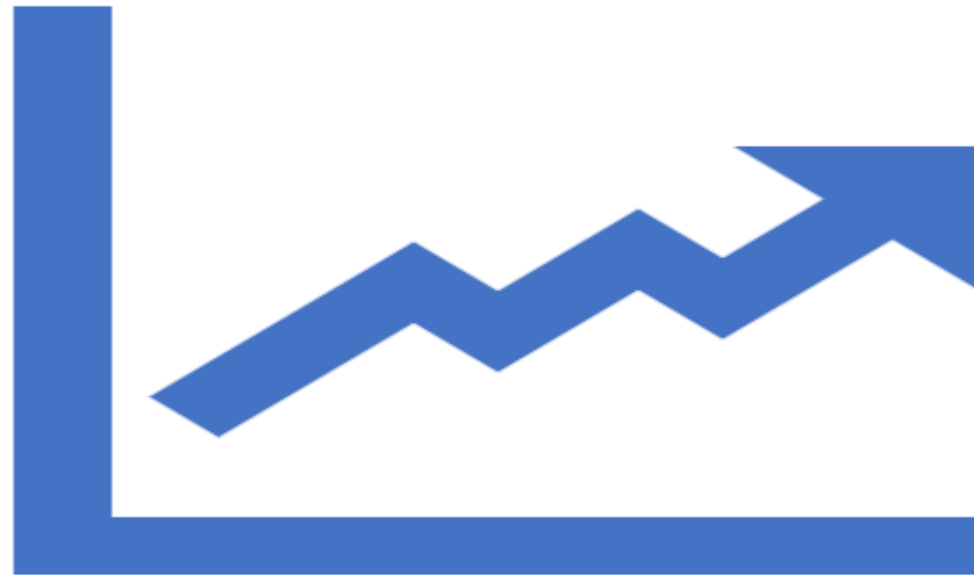
Ranken Namaumbo

## **Dr Rochelle Holm.**

University of Louisville

## **Rockefeller Foundation**

For the Financial support of the pilot





# Wastewater surveillance from Vellore, India - utilities, insights and challenges

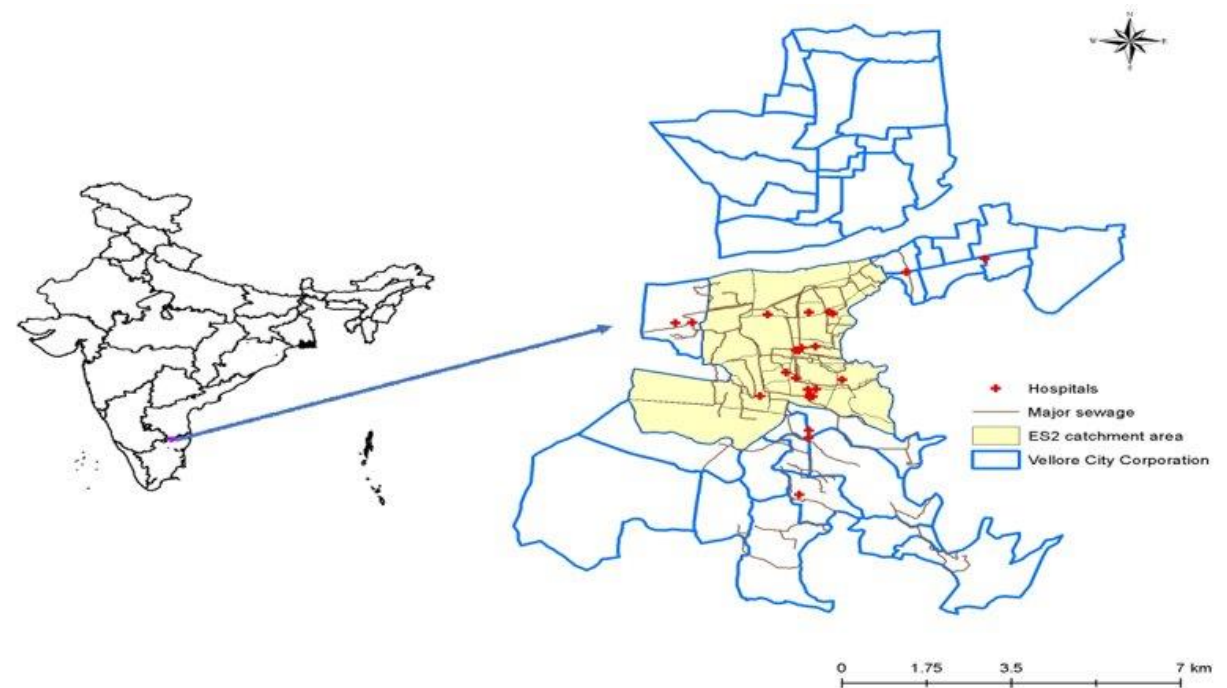
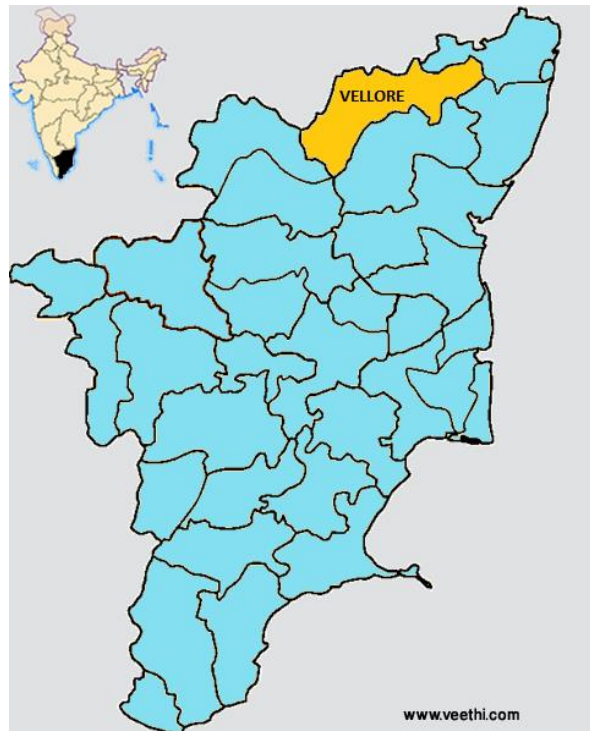
DILIP ABRAHAM, CHRISTIAN MEDICAL COLLEGE VELLORE

[dilip.abraham@cmcvellore.ac.in](mailto:dilip.abraham@cmcvellore.ac.in)

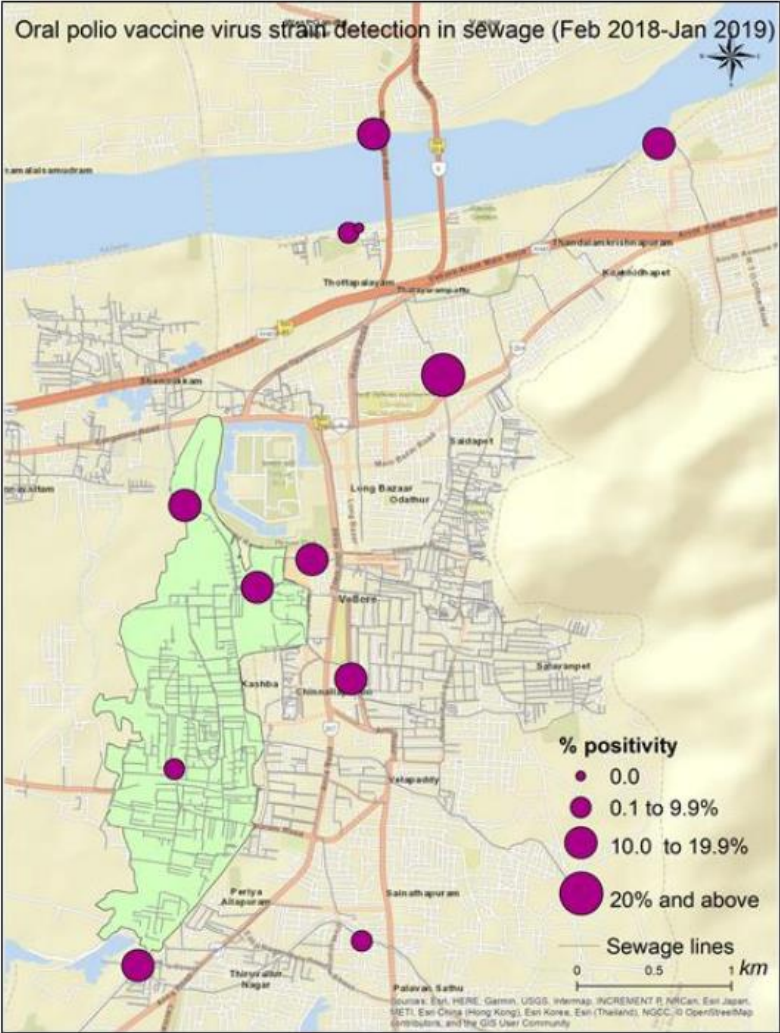


# Study area in Vellore

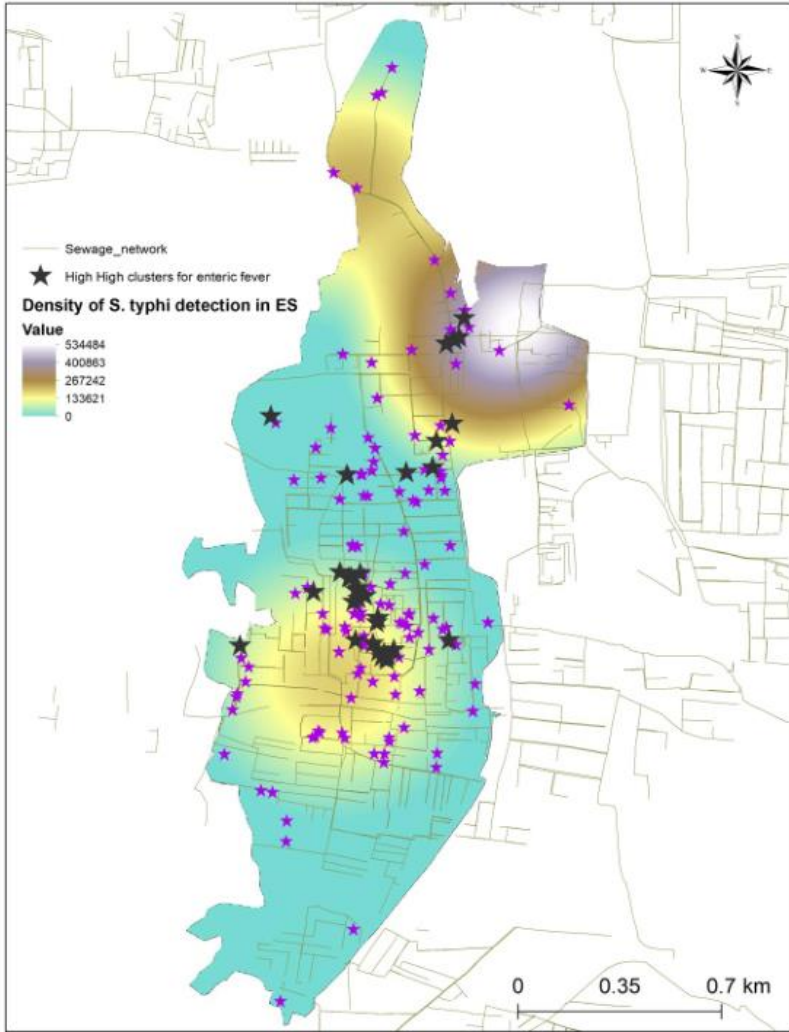
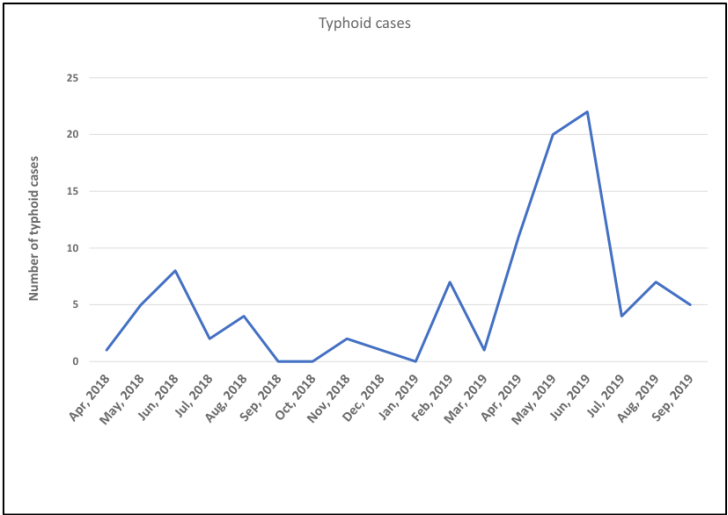
- Environmental surveillance across 24 wards of Vellore city in May 2021
- Spread over 16.25 sq.km; catchment population of 1,95,000 people
- Average population density of 26,500 / sq.km



ES – 1: ENVIRONMENTAL SURVEILLANCE FOR SABIN POLIO VIRUSES,  
SALMONELLA TYPHI AND AMR GENES IN VELLORE (JANUARY 2018 – JUNE 2020)



Distribution of typhoid cases from April, 2018 to September, 2019

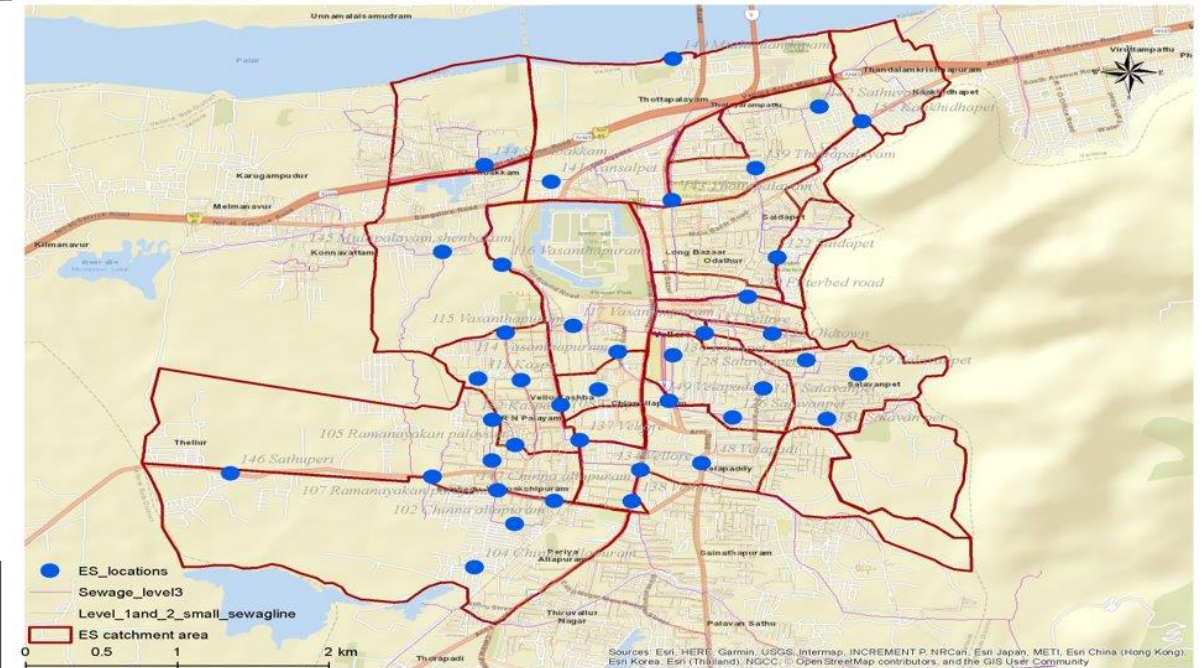
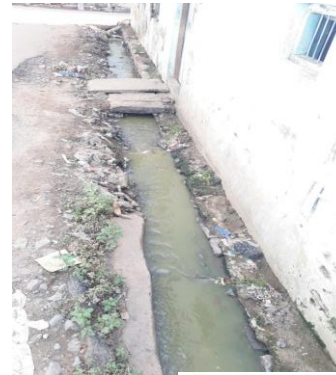
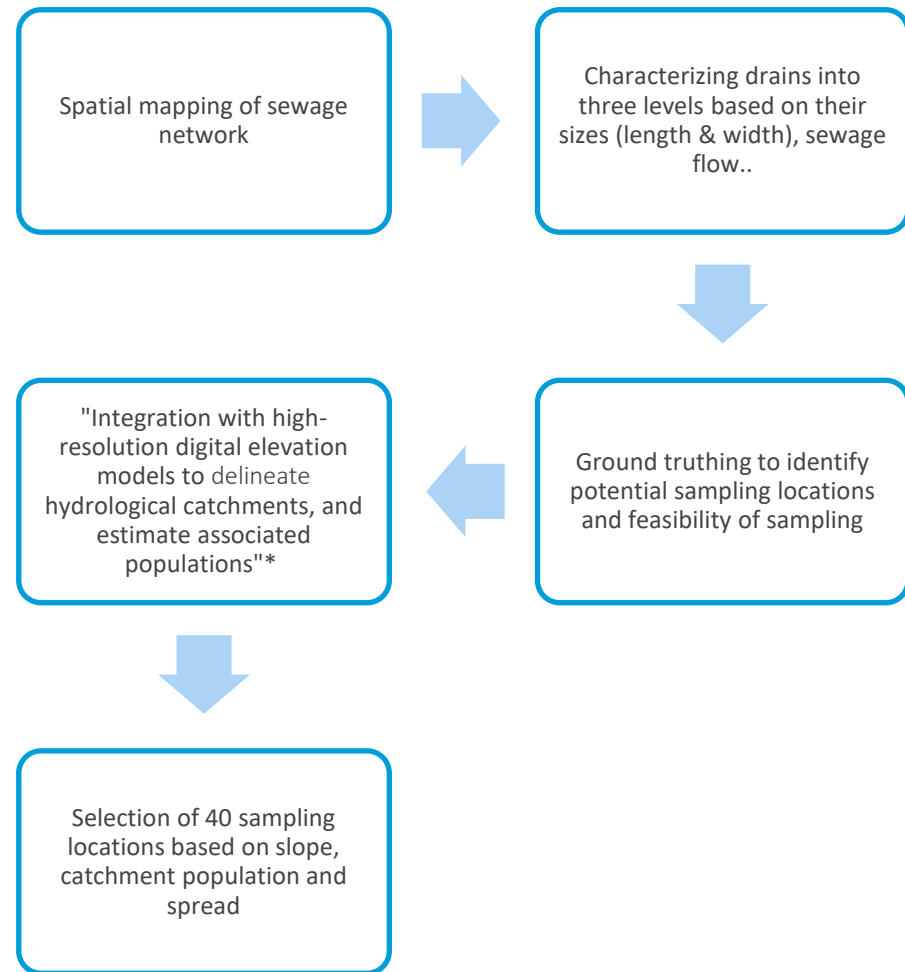




## ES2 - WASTEWATER SURVEILLANCE OF *S. TYPHI* IN INDIA AND COMPARISON WITH CLINICAL/SEROLOGICAL SURVEILLANCE (2020 – 2021)

- investigate the optimal design for a typhoid/paratyphoid environmental surveillance (ES) system that will inform vaccine introduction and impact monitoring
- 1) implement typhoid ES with partners at sites conducting blood-culture surveillance over a 12-month period (India, Malawi and potentially Ghana).
- 2) ask what ES site characteristics (catchment population, local facilities, chemical and physical water-quality parameters) correlate with detection of *S. Typhi* or human-restricted control organisms that indicate faecal contamination.
- 3) investigate whether *S. Typhi* load and genetic diversity in ES samples correlate with disease incidence rates in the local population.

## ES2 - SITE SELECTION



\*Uzzell CB, Troman CM, Rigby J *et al.* **Environmental surveillance for *Salmonella Typhi* as a tool to estimate the incidence of typhoid fever in low-income populations.** Wellcome Open Res 2023, 8:9 (<https://doi.org/10.12688/wellcomeopenres.17687.1>)

# Detection of *S. Typhi* – sample processing

## 1. Membrane filtration

- Collection – 1 L
- Filtration
- Filter processing
- Extraction
- PCR



## 2. Moore Swab

- Preparation
- Deployment
- Enrichment
- Filtration
- Filter processing
- Extraction and PCR



- 2 sample processing methods employed:
  - One grab sample and one "trap" sample
- Sampling frequency:
  - Once per month
- 3rd May 2021 – 29 April 2022
  - 520 grab samples and 517 Moore swabs



# Detection of *S. Typhi* & WW characteristics

Table 5

Salmonella strain (No. tested)	<i>ttr</i>	<i>sseJ</i>	<i>tviB</i>	<i>srfJ</i>	SPC0869	SPA2308	<i>staG</i>
<i>Salmonella Typhi</i> (556)	556	0	553	0	0	0	556
Atypical <i>Salmonella Typhi</i> (3)	3	0	0	0	0	0	3
<i>Salmonella Paratyphi A</i> (315)	315	0	0	0	0	315	0
<i>Salmonella Paratyphi B</i> (53)	53	0	0	53	0	0	0
<i>Salmonella Paratyphi C</i> (6)	6	6	6	0	6	0	0
*NTS Serovar (952)	952	938	0	380	19	50	41
Non- <i>Salmonella</i> spp. (7)	0	0	0	0	0	0	0

\*The combination of genes present were heterogeneous, please see Supplementary Table 1 for details

Nair *et al*, 2019 – A real-time PCR for the differentiation of typhoidal and non-typhoidal *Salmonella*

## Targets

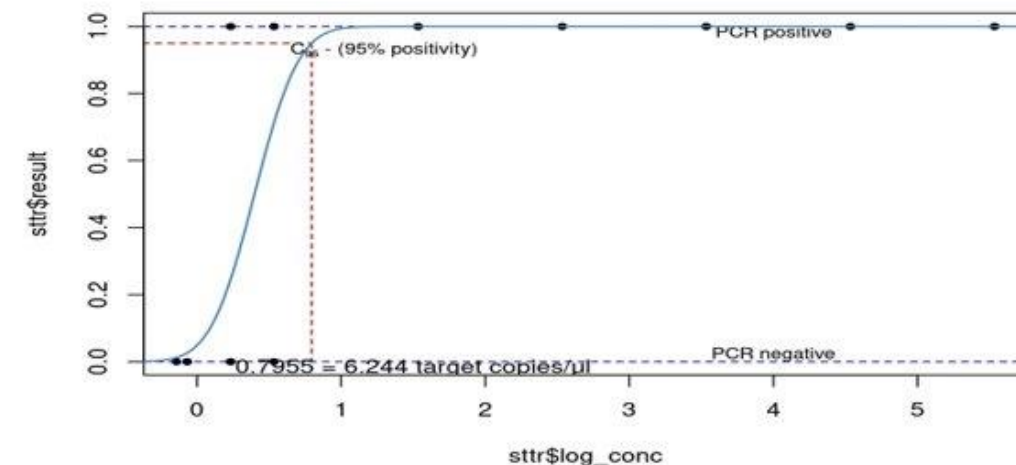
- *ttr*
- *staG*
- *tviB*
- HF-183
- IC
- Sample positive only if all Typhi targets detected
- Double positives were retested for the negative target as a singleplex PCR

## Aquaprobe AP-2000



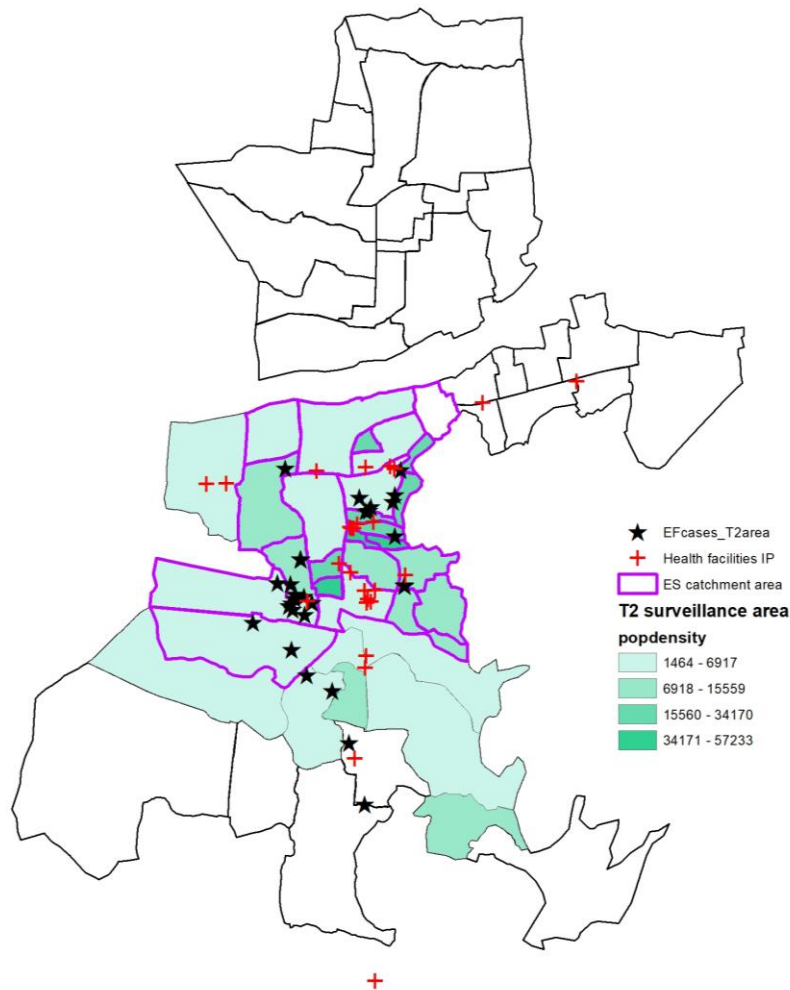
- Temperature
- Baro
- PH
- ORP
- TDS
- Salinity
- Turbidity
- ...

## qPCR standardization ttr probit plot



# Clinical incidence of Typhoid

- Hospital based sentinel surveillance since April 2021
- Recruiting fever cases aged 6 months and above from defined geographical area; 1,90,000 population across 31 wards
- Of the 1108 eligible cases, 95% recruited and blood culture done in 92% of them
- 32 cases of enteric fever over one year from study area
- 28 cases (87.5%) of EF cases were from the ES catchment area
- Of the 28 cases, 17 – *Salmonella* Paratyphi "A" & 11 *S. Typhi*

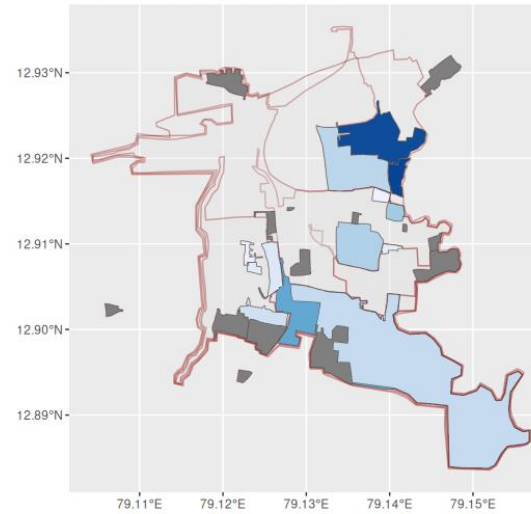


Age Group	No. of Cases	Population	Crude Incidence Per 100 000 Person Years (95% CI)
0–4	4	8082	49.5 (13.5–126.7)
5–9	3	9772	30.7 (6.3–89.7)
10–14	0	10 430	0.0 (.0–35.4)
15–29	4	35 728	11.2 (3.1–28.7)
30+	0	77 788	0.0 (.0–4.7)
Total	11	141 800	7.8 (3.9–13.9)

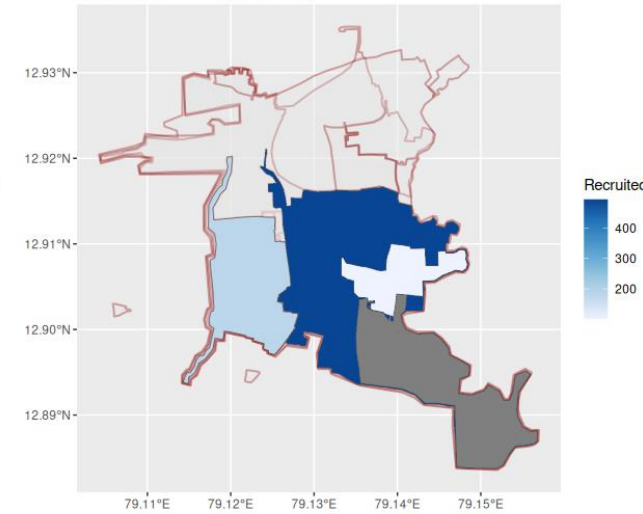
# Sero-survey for HlyE IgG

- Clinical surveillance was limited due to deviation in health-seeking behaviour during the pandemic
- Carried out HlyE IgG testing for 1200 study participants from 0–15-year-old
- Based on catchment areas (CA) for ES site
- Each CA would recruit proportionate age-stratified numbers
- Correlate ES detection with sero-incidence

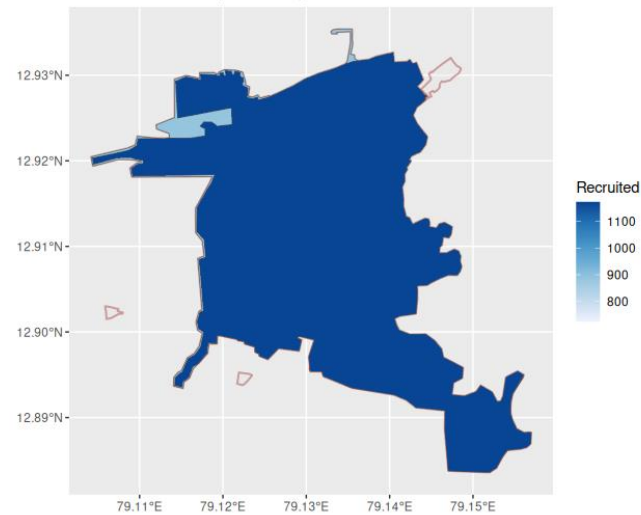
Catchment areas (independent)



Catchment areas (nested=2)



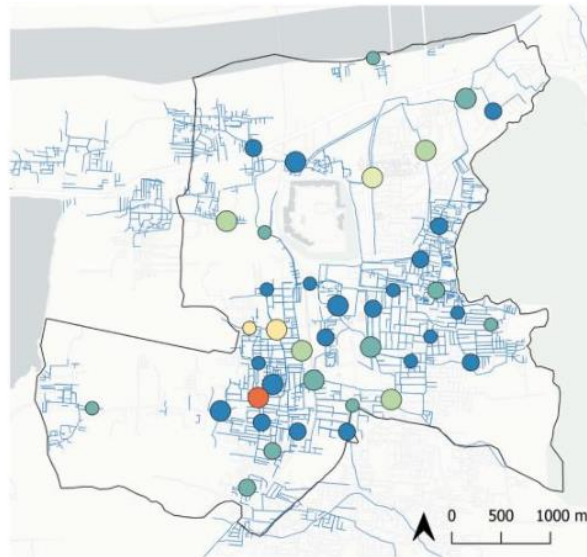
Catchment areas (nested=3)



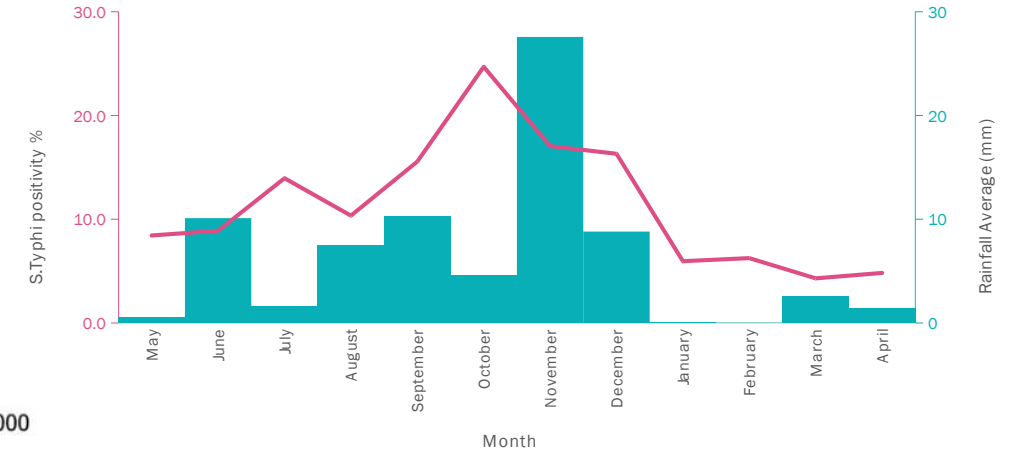
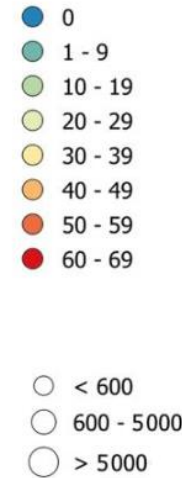
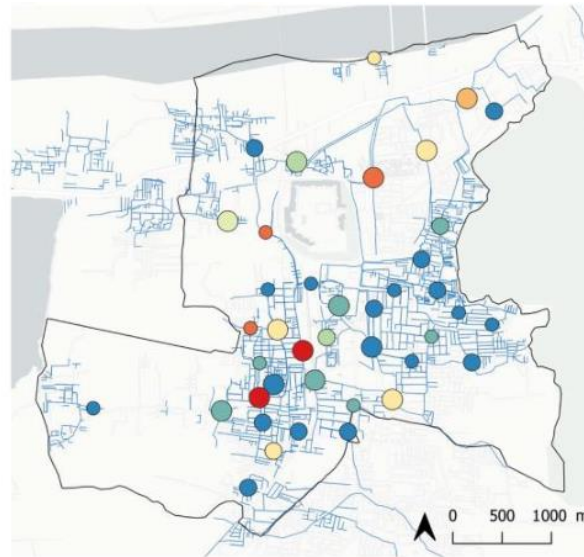


# Results - ES positivity & trends

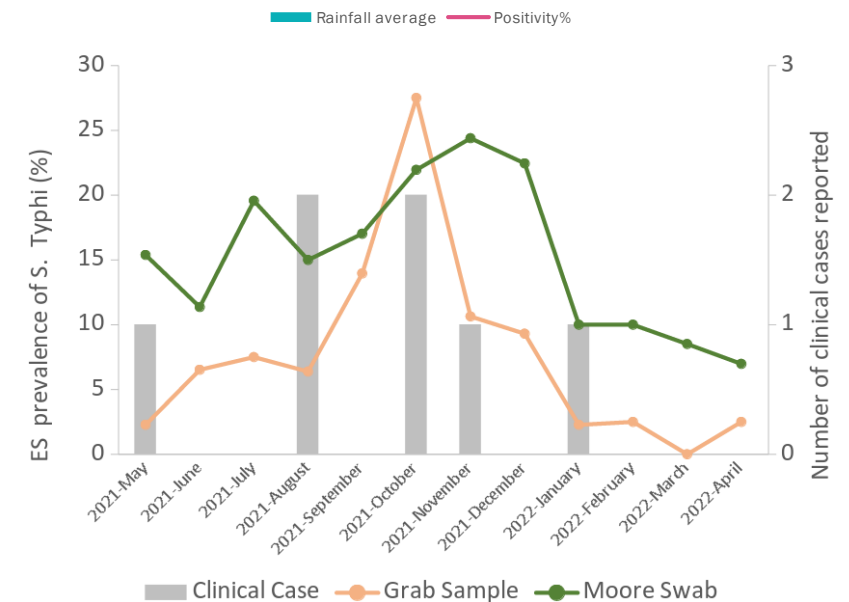
Membrane filtration



Moore swab



Type	Total samples	Positive
Membrane filtration	520	39 (7.5%)
Moore	517	79 (15.3%)
Total	1037	118 (11.4%)



# Utilities, insights and challenges?

## Challenges/Watch out for

- Validate the method at your own site and train personnel
- Loss of Moore swabs – rat bites, removal by members of the local community
- Time management
- Reagent procurement and supply chain
- Data management
- qPCR validation
- Avoid freeze-thaws of DNA extracts
- Scalability and automation
- What is a positive sample?
- Normalization – MST? Flow rate?
- Communication with public health personnel on how to use this data

## Insights

- Sampling frequency and methodology would need to fit with the defined use-case and setting of ES
- Data analysis is not standard and would need to be tailored for the pathogen and use-case
- Variability of detection is very high in LMIC settings
- Molecular targets should be chosen after thorough literature review and preferably with consensus in the field
- Delineation and description of catchment areas is essential
- Availability of some amount of clinical surveillance data or estimation of disease burden is essential for validation

## Utility

- Estimation of disease burden - typhoid
- Predicting outbreaks – SARS-CoV-2
- Detection of novel variants/variant of concern – SARS-CoV-2, polio
- Localization of source of outbreak
- Delineation and description of catchment areas is essential
- Availability of some amount of clinical surveillance data or prior estimation of disease burden is essential for validation

# Q&A Discussion

MODERATOR: GERTJAN MEDEMA & SUDHIR PILLAY



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