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# Method optimisation for detection of *Salmonella* Typhi from the environment

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# Malawi background & rationale



Malawi is an typhoid endemic country

- Outbreak of unknown origin in Blantyre, Malawi started 2012
- >16,000 cases per year, ~200 deaths associated with typhoid

Methods Development Project

Primary aim: culture & detect *S. Typhi* from the environment

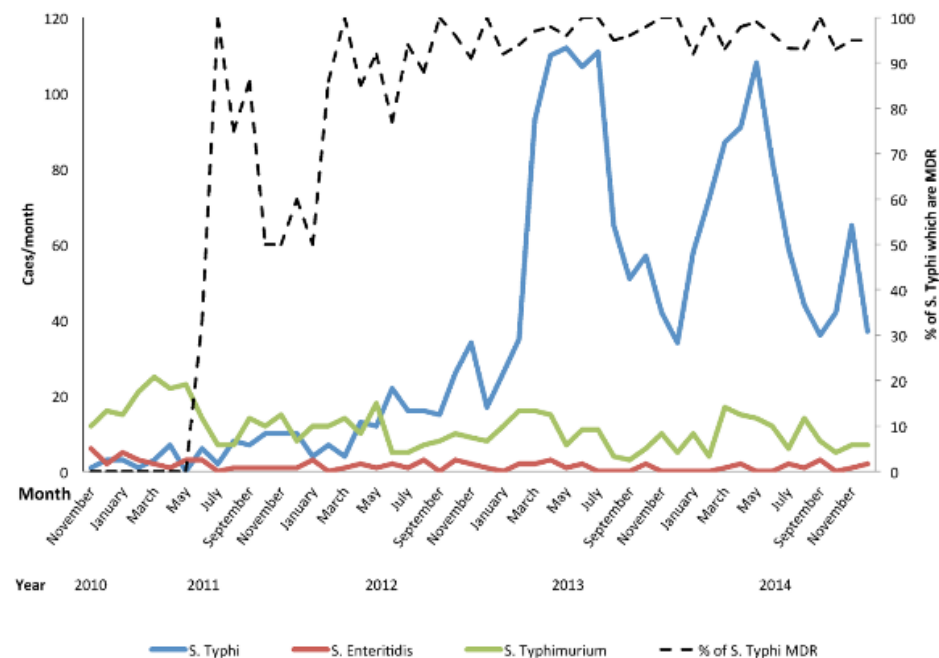


Figure 1: Feasey et al., 2015, Monthly trends in bloodstream invasive Salmonella diagnosed at QECH from November 2010-October 2014.

# Isolation of *Salmonella* Typhi from the environment is difficult but not impossible

## Sampling from sewerage proved effective

- ≤1950s
  - Moore's swabs
  - Specific required culture methods
- 1980s
  - Reinforced the use of Moore's swabs
  - Selenite-F is the most effective for *S. Typhi*
- Viable but non-culturable (VBNC)

## New approaches use molecular methods

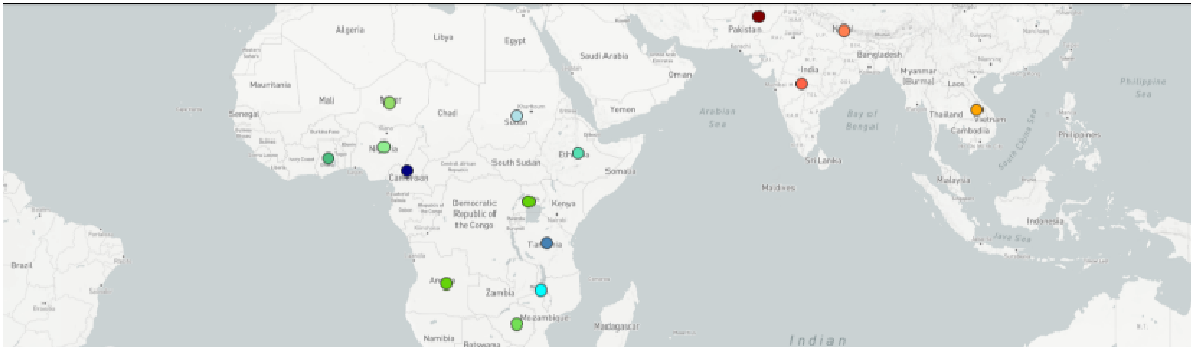
- Strongly associated with water & indirect transmission in endemic regions

## Culture remains important



Figure 1: Electron micrograph of a salmonella taken by Matthew Hannah, PHE

## Twenty isolates from the PHE culture collection were used



*Figure 2: Global distribution of strains used in optimisation & evaluation of environmental isolation (Satheesh Nair, PHE)*

Isolates from different regions selected for laboratory testing

The use of a novel chromogenic agar, mCASE was selected for the isolation of salmonella



*Figure 3: Pure S. Typhi on mCASE*

## Methods for the isolation of *Salmonella* Typhi were assessed

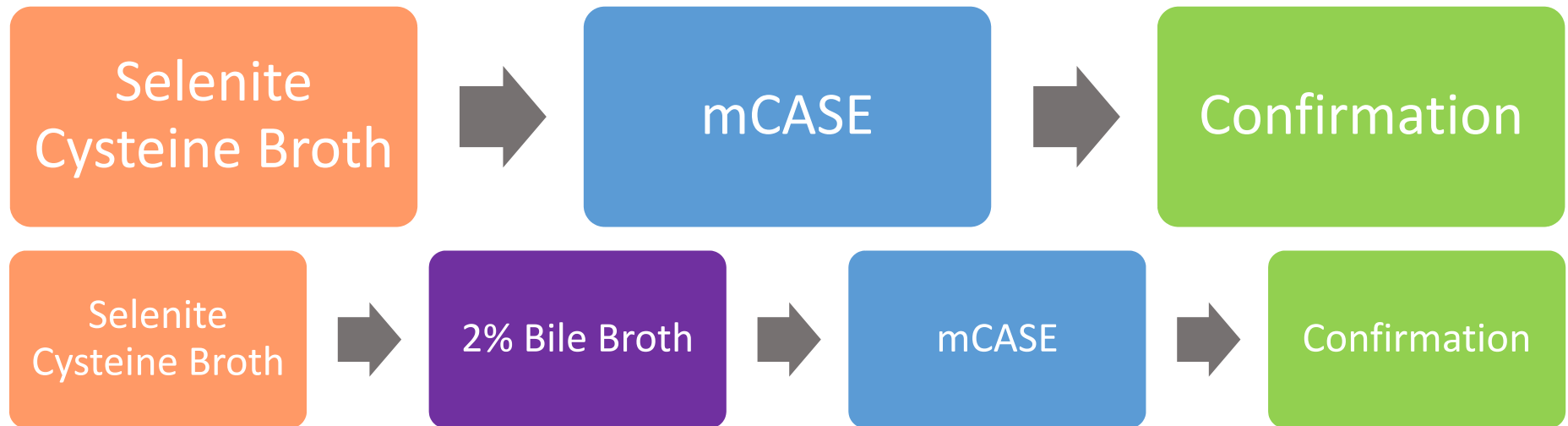


Figure 4: Two examples of the pathways assessed

- Pathways were narrowed to four options from original 15
  - Selenite cysteine broth was preferred due to its selectivity but does not remove *S. Typhi* like alternatives (e.g. Rappaport Vassiliadis broth)
  - Bile broth: infection starts post exposure to bile

# Too Many Plates

## Preliminary working at PHE Food, Water & Environmental Laboratory

- Hundreds of plates a week processed
  - ❖ Under laboratory conditions with control strains, *S. Typhi* was reliably retrieved
  - ❖ Environmental testing required



Photo Credit: <https://www.flickr.com/photos/erikaleef>

# Following narrowing of the pathways, challenges set up with blind, mixed cultures

## Challenge Organisms

- *Salmonella* Nottingham – same colour as *S. Typhi* on mCASE
- *Bacillus cereus* – similar colour & overgrows
- Fungi - blind cultures grew relatively well in the broths

## Immuno-magnetic bead separation ran in parallel

- Pan-*Salmonella* bead, developed by Ezzeddine Elmerhebi, Neogen LabM
- Further work required *in situ* as artificial mixtures showed no major difference in recovery rates



Figure 5: Mixed culture on mCASE, where *S. Typhi* was successfully selected

# Molecular approaches utilised to reinforce culture methods developed

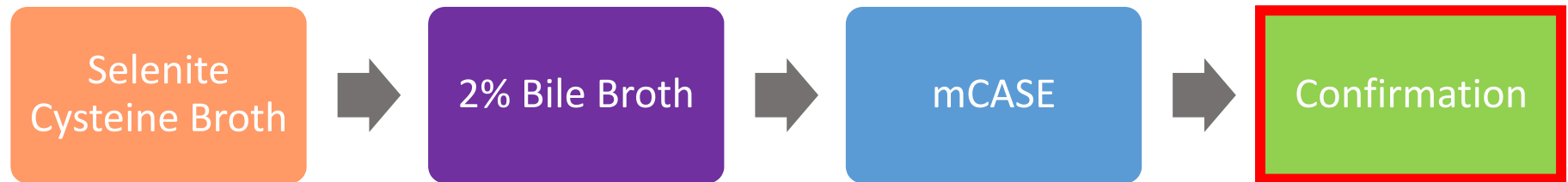


Figure 6: Pathway emphasising confirmation step

- The main aim of the project is to develop a method for culturing environmental *S. Typhi*
- Molecular methods are more cost effective & high-throughput
- Quantitative PCR also allows for screening of incoming samples
  - Inhibitors such as environmental, chemical & faecal contaminants

Note: Current DNA-based molecular methods cannot prove viability



# Novel assay from PHE used for rapid diagnostics



Satheesh Nair, PHE, designed an assay for diagnostics

- The original assay included primer's for Paratyphi A, B & C
- This project aims to multiplex the *S. Typhi* targets
- Multiplexed assay is currently being optimised
- Probes changed from original to minimise interference in multiplex

Table 1: Primer sequences of PHE assay and source publications

Gene	Name	Sequence 5'-3'	NCBI Accession Number	Reference
<i>ttr</i>	ttr_F	CTCACCAGGAGATTACAACATGG	AF282268	Hopkins, 2009
	ttr_R	AGCTCAGACCAAAAGTGACCATC		
	ttr_P	FAM-CACCGACGGCGAGACCGACTTT-BHQ1		
<i>sseJ</i>	sseJ_F	CGAGACTGCCGATGCATTTA	AF294582	Nair, 2019
	sseJ_R	GTACATAGCCGTGGTGAGTATAAG		
	sseJ_P	YY-TGGAGGCGGCCAGTAATATTGGTT-BHQ1		
<i>tviB</i>	tviB_F	TGTGGTAAAGGAACTCGGTAAA	NC_003198	Nair, 2004
	tviB_R	GACTTCCGATACCGGGATAATG		
	tviB_P	CY3-TGGATGCCGAAGAGGTAAGACGAGA-BHQ2		
<i>staG</i>	staG_F	CGCGAAGTCAGAGTCGACATAG	AL513382	Nga, 2010
	staG_R	AAGACCTCAACGCCGATCAC		
	staG_P	CY5-CATTTGTTCTGGAGCAGGCTGACGG-BHQ2		

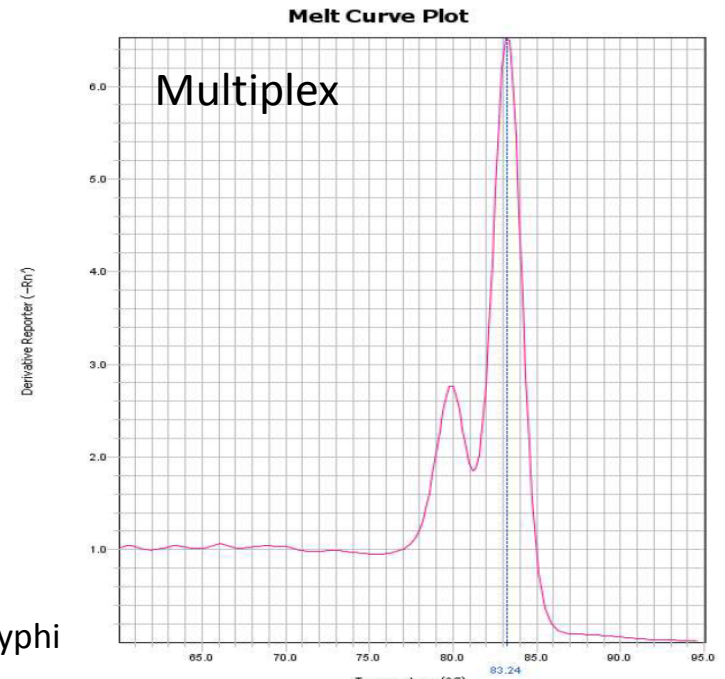
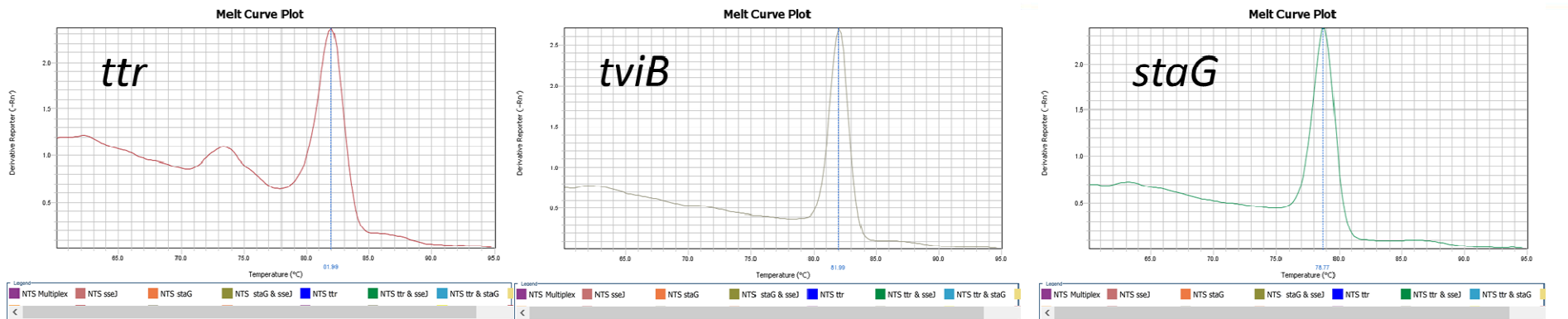
# High-Resolution Melt (HRM) PCR



The multiplex assay is being adapted to HRM

- Thomas Edwards, LSTM, consulted in assay conversion
  - ❖ Results consistent with *S. Typhi* & non-typhoidal salmonella control strains
  - ❖ Further optimisation & design changes

Figure 7: HRM melt curves for positive *Salmonella Typhi*



# Role of the assay

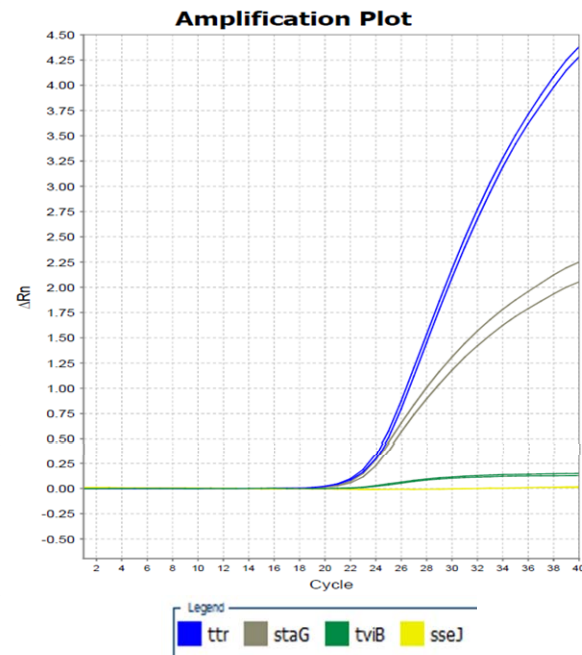


Figure 8: q-PCR assay for blind broth "A"

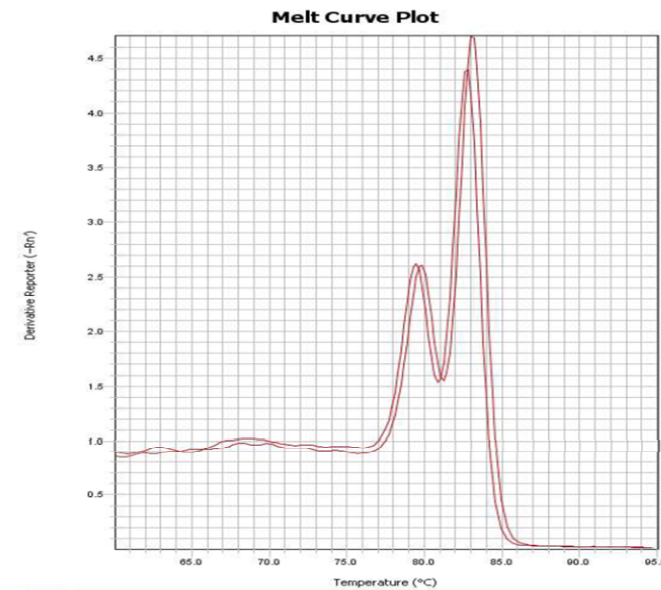


Figure 9: HRM-PCR assay for blind broth "A"

As a tool for screening & confirmation, the blind cultures were processed with both the probe based q-PCR & the HRM PCR

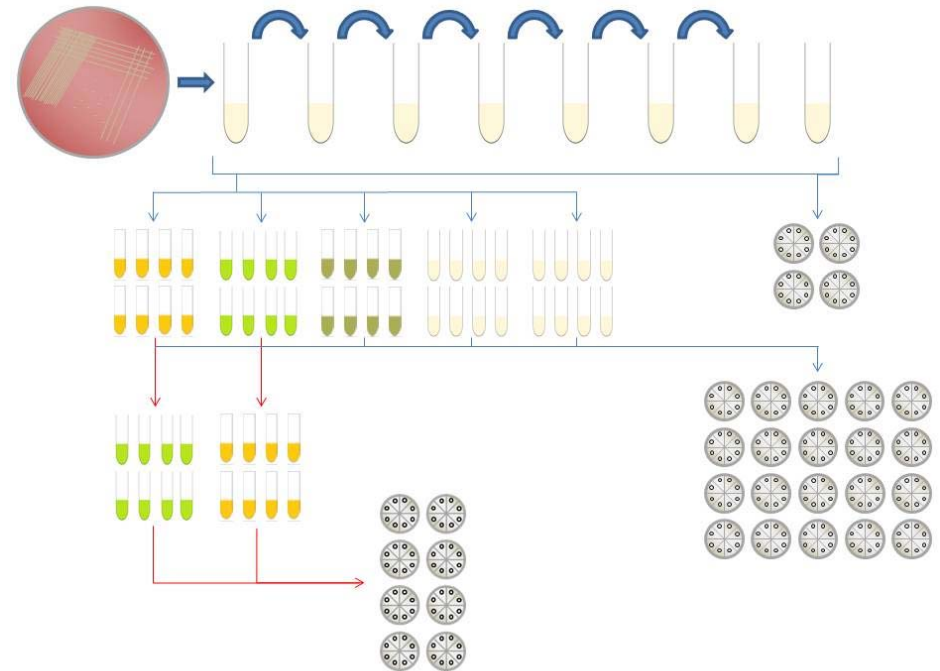
HRM: <£0.40 (<\$0.5)

Probe: <£0.50 (<\$0.6)

# Moving forwards, work to be done in Malawi

## Final Culture Pathways

- Limit of detection (LOD) & limit of quantification (LOQ) work still to be performed
- To finalise culture pathways based on further evaluation in Malawi



# Moving forwards, work to be done in Malawi

Sampling strategy – in collaboration with Jillian Gauld, University of Lancaster

- Pilot Study
- Challenges
  - ❖ Road conditions & water access
  - ❖ Logistics

## Methodology

- Continued optimisation
- Immuno-magnetic separation & microfluidics
- Adapt to new challenges from environmental samples

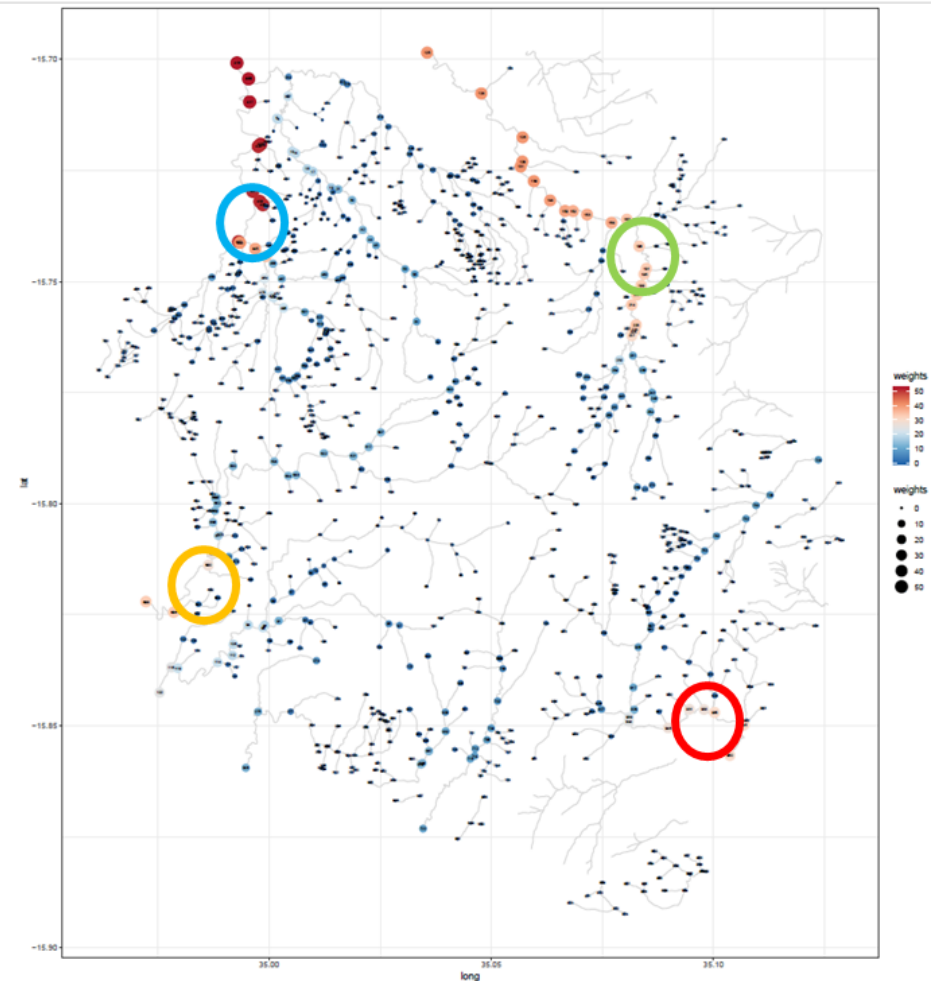


Figure 10: Cumulative map of Typhoid cases with highlighted sampling environmental areas

# Likhubula River

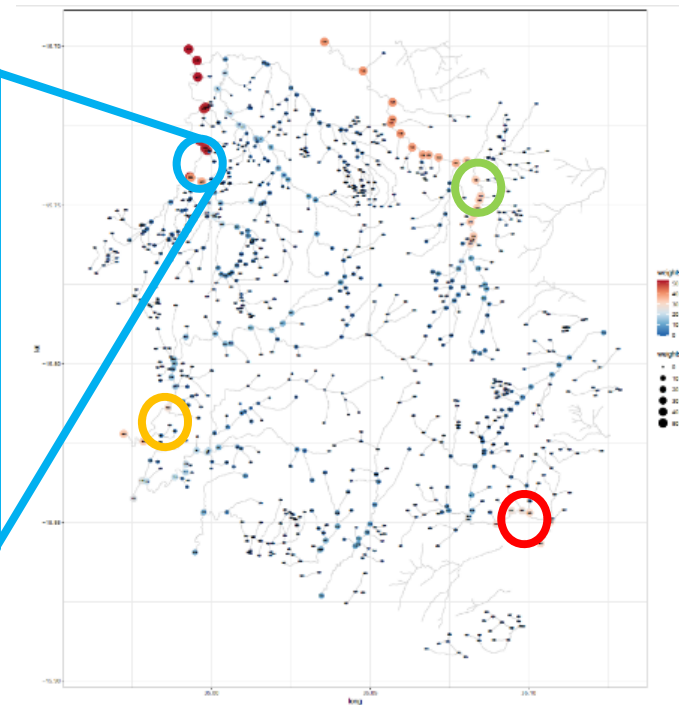


Figure 11: Examples of water access on the Likhubula River

# Mudi River

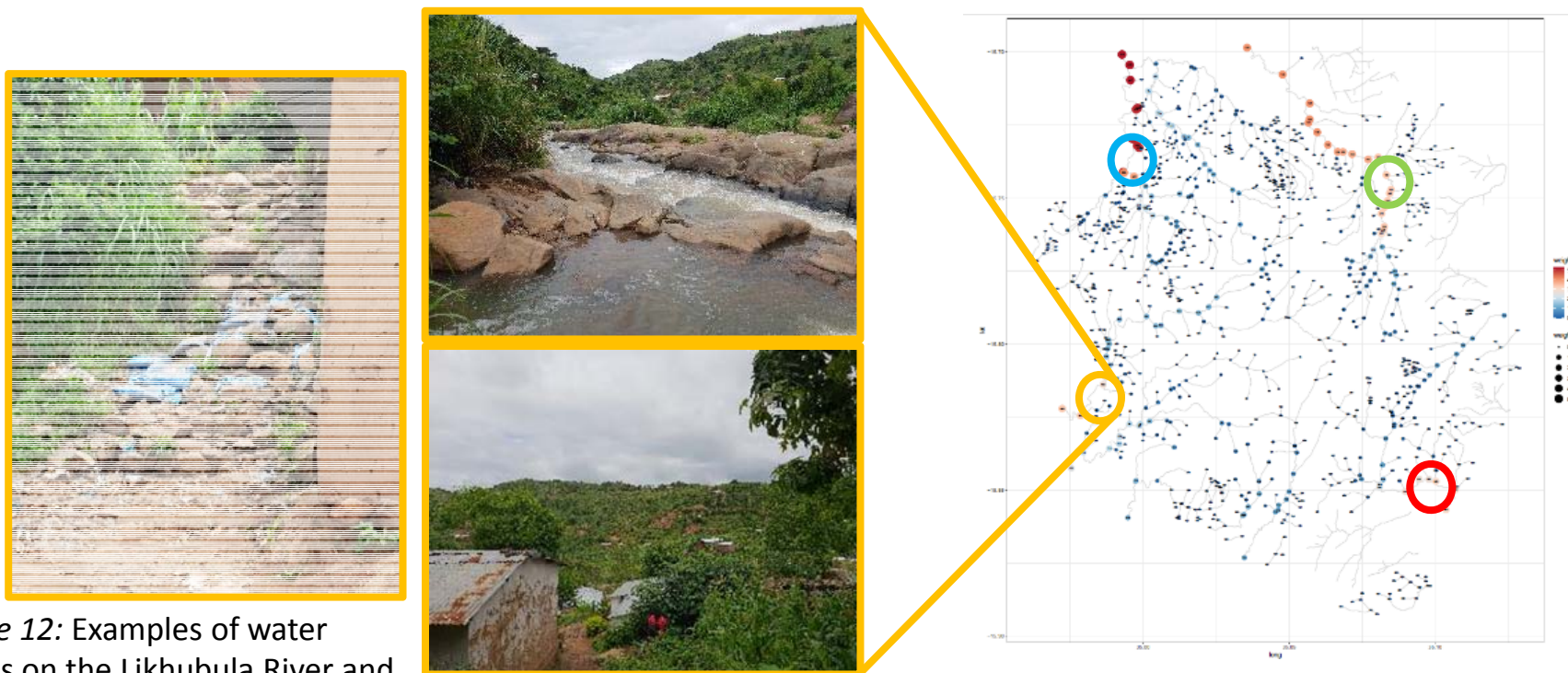


Figure 12: Examples of water access on the Likhubula River and walkway through village

# Lunzu River



Figure 13: Examples of water access on the Lunzu River



# Thuchila River

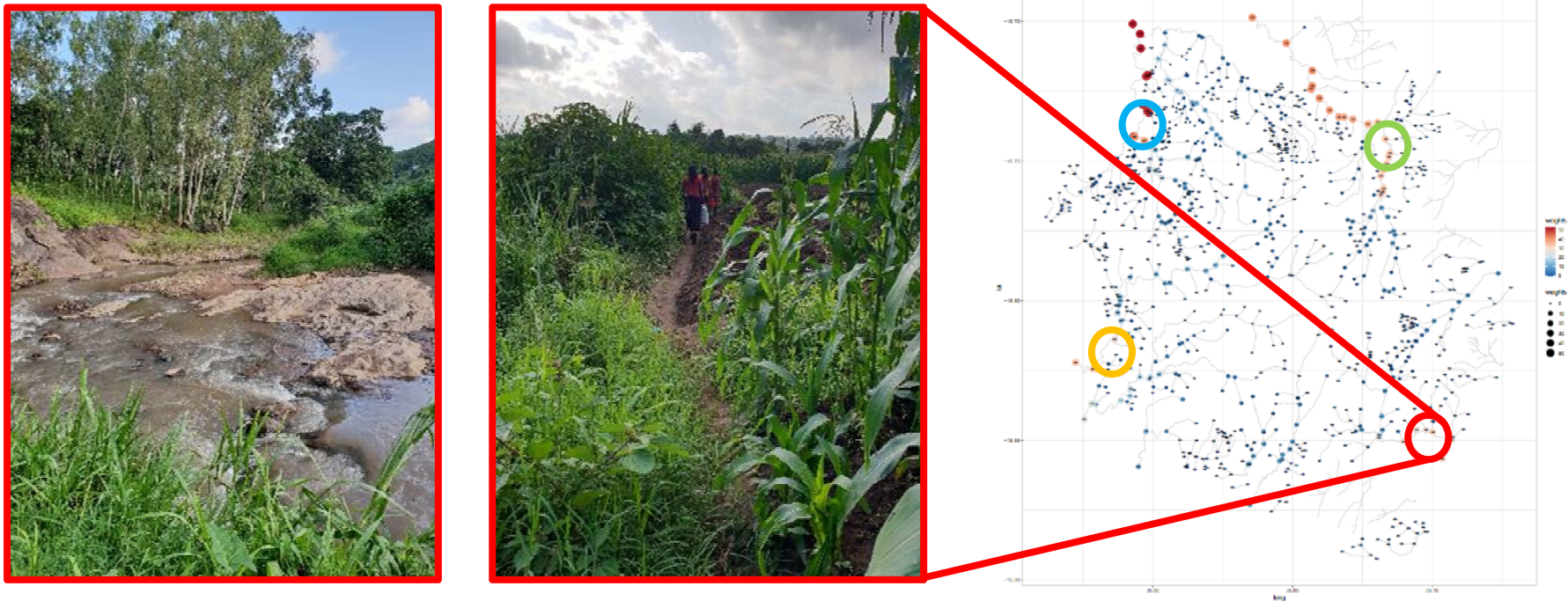


Figure 14: Examples of water access on the Thuchila River

# acknowledgements



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