Detection of *Mycobacterium tuberculosis* in pediatric stool samples using TruTip technology

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Abstract

Background: Rapid and accurate diagnosis of childhood tuberculosis (TB) is challenging

because children are often unable to produce the sputum sample required for conventional tests.

Stool is an alternative sample type that is easy to collect from children, and studies investigating

the use of stool for molecular detection of *Mycobacterium tuberculosis (Mtb)* have led to

promising results. Our objective was to evaluate stool as an alternative specimen to sputum for

Mtb detection in children. We did so using the TruTip workstation (Akonni Biosystems), a novel

automated lysis and extraction platform.

Methods: We tested stool samples of 259 children who presented with TB symptoms, aged 0-14

years old, in Lima, Peru. Following extraction with TruTip, we detected the presence of *Mtb*

DNA by IS6110 real-time PCR. We calculated assay sensitivity in two groups: (1) children with

culture confirmed TB (N=22); and (2) children with unconfirmed, clinically diagnosed TB

(N=84). We calculated specificity among children in whom TB was ruled out (N=153). Among

children who were diagnosed with TB, we examined factors associated with a positive stool test.

Results: Assay sensitivity was 59% (95% confidence interval [CI]: 39%-80%) and 1.2% (95%

CI: 0.0%-6.5%) in children with culture-confirmed and unconfirmed clinically-diagnosed TB,

respectively, and specificity was 97% (95% CI: 93%-99%). The assay detected *Mtb* in stool of

7/7 children with smear-positive TB (100% sensitivity; 95% CI: 59%-100%), and in 6/15 of

children with smear-negative, culture-confirmed TB (40% sensitivity; 95% CI: 16%-68%). Older

age, smear positivity, culture positivity, ability to produce sputum and cavitary disease were

associated with a positive stool result.

Conclusion: Testing of stool samples with the TruTip workstation and IS6110 amplification,

yielded sensitivity and specificity estimates comparable to other tests such as Xpert. Future work

should include detection of resistance using the TruTip closed amplification system and assay

optimization to improve sensitivity in children with low bacillary loads..

Keywords: Tuberculosis; Children; Molecular diagnostics, TruTip

<u>Background</u>

The World Health Organization estimates that one million new pediatric tuberculosis (TB) cases

and 194,000 childhood TB deaths occurred in 2017 (1). Rapid case detection and treatment

initiation is critical to minimizing TB morbidity and mortality in children but is hampered by the

absence of a rapid, accurate diagnostic tool for this group. Bacteriologic confirmation of

Mycobacterium tuberculosis (Mtb) in children is often difficult to achieve because they are

frequently unable to expectorate sputum for bacteriologic testing and often have paucibacillary

disease that cannot be detected using sputum smear microscopy, culture, and/or molecular testing

[e.g. Xpert (Cepheid, Sunnyvale CA, USA)]. Sputum induction and gastric aspiration can be

used to obtain respiratory specimens from children unable to expectorate sputum; however,

gastric aspiration is invasive and neither procedure is widely implemented in resource-

constrained settings. Due to these diagnostic challenges, bacteriologic confirmation of TB is

obtained in only a small minority of children diagnosed with TB (2–4).

Stool can be easily obtained from most children, and *Mtb* can be detected in stool using Xpert

(5–10) or other laboratory-developed PCR assays (11–13). In the case of Xpert, the process is

automated, but detection of drug resistance is currently limited to rifampin-associated resistance

mutations in rpoB. Non-integrated merthods for DNA, isolation and amplification of DNA using

extraction kits and in-house tests are often laborious, multistep procedures. An ideal test would

be an automated point-of-care workstation with integrated capacity for both *Mtb* and expanded

drug resistance testing-criteria listed in the target product profile for novel TB diagnostics in low

resource settings (14). The TruTip workstation is an automated platform including lysis and

homogenization with TruTip nucleic acid extraction and purification (Akonni Biosystems,

Frederick, MD, USA) (15–17). TruTip has been used for nucleic acid isolation from a variety of

pathogens and sample types and has demonstrated efficient *Mtb* DNA recovery from raw sputum

(15,18). The platform can be connected to a closed amplicon system for amplification and

microarray-based detection of *Mtb* as well as a number of drug resistance-associated mutations

(17,19–21). The aim of the present study was to estimate sensitivity and specificity of *Mtb*

detection in stool from children with symptoms compatible with intrathoracic TB in Lima, Peru,

using this novel technology with IS6110 real-time PCR.

Methods

Ethics

Study participants' guardians provided written informed consent to participate, and children

eight years of age and older provided written assent. Consent for publication was not applicable.

All study procedures were approved by the Ethics Committee of Peru's National Institute of

Health and the Office of Human Research Administration at the Harvard Medical School.

Study population

Between May 2015 and February 2018, we recruited children to participate in a pediatric TB

diagnostics study. Eligible children were less than 15 years of age, had a history of contact with

an adult with TB within the previous two years, and presented to a participating public sector

health center in Lima, Peru with symptoms compatible with TB (i.e., persistent cough for more

than two weeks; unexplained weight loss; unexplained fever for more than one week; and/or

unexplained fatigue or lethargy) (22). For this analysis, we included the subset of children with

culture-confirmed TB or clinically-diagnosed unconfirmed TB who had at least one stool sample

available. We also included a subset of the children in whom TB had been ruled out (i.e,

controls), matching on age and sample collection date when possible.

Study procedures and sample collection

Children were evaluated for TB per Peruvian National Tuberculosis Strategy guidelines (23). In

brief, children provided up to two gastric aspirate (GA) and/or sputum samples (expectorated or

induced) for smear and culture, and Ministry of Health pediatric pulmonologists considered these

results as well as medical history, physical examination, chest X-ray findings and tuberculin skin

testing (TST) results to diagnose or rule out TB. GA samples were neutralized to a pH of 6.8-7.2

upon collection. We requested two stool samples from all children for research purposes. From

children who were diagnosed with TB, we aimed to collect these samples prior to TB treatment

initiation. Stool collection took place at home or the health center. For children in diapers, plastic

wrap was inserted on the inside of the diaper and a urine collection bag was attached to the child.

The latter served to collect a urine sample as well as to prevent mixing of urine and stool.

Following collection, all samples were transported by cold-chain to the laboratory, aliquoted and

stored at -80°C until testing. Because testing of stool did not occur in real time and because the

assay was experimental, stool testing results were not shared with families or healthcare

providers. HIV testing was not performed because the HIV prevalence among children in the

study setting is <0.1% (24).

Laboratory procedures

Sample processing and testing was performed in the BSL-3 and BSL-2 facilities at the Socios En

Salud Sucursal Laboratory in Lima. We centrifuged and decontaminated sputum and GA

samples in 2% NaOH/0.25% n-acetyl-L-cysteine (NALC). GA and sputum samples were

analyzed by Ziehl-Neelsen staining/smear microscopy and liquid culture (BACTEC MGIT 960,

BD Franklin Lakes, USA).

For lysis and DNA extraction procedures using the TruTip workstation we followed methods

similar to previously described for sputum samples (18). In brief, 500 milligrams of thawed stool

was homogenized in TE buffer, in a magnetically-induced vortexing (MagVor) element, and

after a 20 minute heating step at 56°C, the samples were transferred to the TruTip workstation

for DNA extraction (16). *Mtb* DNA was amplified in duplicate by real-time PCR using a Roche

480 Lightcycler (Basel, Switzerland) with primers targeting the insertion element IS6110 (25).

We considered a sample positive for *Mtb* DNA if, for both PCR replicates, the cycle threshold

(Ct) value was <37 and the change in fluorescence was >6 units. We repeated the PCR step in

duplicate for discordant replicates and considered the sample to be indeterminate for *Mtb* DNA if

the two replicates remained discrepant after a second PCR run. Laboratory staff were blinded to

participants' clinical status.

Data analysis

Data were analyzed using SAS (Version 9.4. SAS Institute Inc., Cary, NC, USA). We calculated

sensitivity and specificity and 95% confidence intervals (CI) both at the child-level (including all

samples from each child) and at the sample-level. Sensitivity of *Mtb* detection from stool was

calculated separately for children with confirmed TB (i.e., based on a positive culture from

sputum or GA) and clinically-diagnosed unconfirmed TB. Specificity was estimated among

those in whom TB had been ruled out. In per-sample analyses we adjusted confidence intervals

for clustering from multiple samples per child. Among all children with a TB diagnosis, we used

chi-square tests to examine the associations between a positive stool test and the following

variables: age, sex, smear and culture results, ability to spontaneously expectorate sputum, and

cavitation on chest x-ray. Numerical precision for all percentages, p-values and 95% CIs is

presented based on the evidence-based recommendations proposed by Cole (26).

Results

Study population

From 628 children enrolled in the study, we selected 259 children based on stool sample

availability (22 children with culture confirmed TB, 84 children with unconfirmed, clinically

diagnosed TB, and 153 children in whom TB had been ruled out (Fig 1)). These children had a

median age of 5.1 years and 48% were female. Nearly all children (255/259; 98%) had at least

one respiratory sample collected. Of the four with no respiratory sample, three were diagnosed

based on clinical criteria and TB was ruled out in one. Children between the ages of 11 and 14

years old were more likely than younger children to have a culture-confirmed diagnosis, and all

seven children with smear-positive sputum belonged to this older age group (Table 1). Thirteen

of 22 children with confirmed TB were diagnosed based on expectorated sputum analysis and the

remaining nine on gastric aspirate analysis.

Table 1: Demographic characteristics of children, stratified by TB status.

Variable		All children (N=259) n (%)	SM+, CU+ TB[a] (N=7) n (%)	SM-, CU+ TB[a] (N=15) n (%)	All CU+ TB (N=22) n (%)	Unconfirmed clinically diagnosed TB (N=84) n (%)	No TB diagnosis (N=153) n (%)
Age group (years)	0-4	125 (48)	0 (0)	8 (53)	8 (36)	42 (50)	75 (49)
	5-10	37 (14)	0 (0)	5 (33)	5 (23)	34 (40)	58 (38)
	11-14	97 (37)	7 (100)	2 (13)	9 (41)	8 (10)	20 (13)
Female sex		124 (48)	5 (71)	6 (40)	11 (50)	47 (56)	66 (43)

[a]CU: culture, SM: smear

From the 259 children, we collected 484 stool samples. Most children had two stool samples;

however 34 children (13%) had only one sample available. Seventeen samples from children

diagnosed with TB were collected after TB treatment initiation (median time from treatment

initiation to collection: 3 days; range [1 – 11 days]). Eight samples were collected 72 hours or

more after treatment start. Four of these eight were in children with smear-positive sputum

because they were eligible to initiate TB treatment immediately, resulting in a narrow window

between enrollment and treatment initiation during which pre-treatment samples could be

collected.

Per sample analyses

The assay detected *Mtb* in 25/484 samples. Only one sample, from a child in the control group,

had an indeterminate result, which we excluded from further analyses. Among 13 stool samples

from seven children with smear positive sputum, 100% (95% confidence interval [CI]: 59%-

100%) were positive by the stool assay (Table 2). Sensitivity was lower in the smear-negative

culture-confirmed group (23%; 95% CI: 5.6%-41%, Table 2). We detected *Mtb* in 4/282 (1.4%)

samples from control children, resulting in a specificity of 99% (95% CI: 97-99.8%).

Table 2: Per sample analyses of *Mtb* stool assay in 259 children.

	Positive stool result (n/N)	Indeterminate stool result (n/N)	Sensitivity (95% CI[a])	Specificity (95% CI[a])
All samples	25/484	1/484		
Confirmed TB (SM+)[b]	13/13	0/13	100% (59%-100%)	
Confirmed TB (SM-)	7/30	0/30	23.3% (5.6%-41%)	
Confirmed TB (all smear grades)	20/43	0/43	47% (27%-66%)	
Unconfirmed, clinically diagnosed TB	1/159	0/159	0.6% (0.00%-1.9%)	
No TB diagnosis	4/282	1/282		99% (97%-100%)

[a]CI: confidence interval; 95% CI are adjusted for multiple samples per patient. [b]SM: smear

Per child analyses

In child-level analyses, sensitivity of the stool assay conducted in each child's first sample was

55% (95% CI: 33%-75%). When we included the results of both samples, we identified one

additional confirmed TB case, which increased sensitivity to 59% (95% CI: 39%-80%) (Table

3). While the majority of children with a positive stool sample also had a spontaneous sputum

sample that was culture positive (11/14, 79%), we detected *Mtb* in the stool of two children in

whom TB was confirmed based on gastric aspirate and one child with unconfirmed, clinically

diagnosed TB (this child had one expectorated and one induced sputum sample, both of which

tested negative for *Mtb*). All three children were under five years of age.

Table 3: Child-level analyses of *Mtb* stool assay in 259 children.

		Positive stool result (n/N)	Sensitivity (95% CI[a])	Specificity (95% CI)
First sample	All children	16/259		
	Confirmed TB (SM+)[b]	7/7	100% (59%-100%)	
	Confirmed TB (SM-)	5/15	33% (13%-61%)	
	Confirmed TB (all smear grades)	12/22	55% (33%-75%)	
	Unconfirmed, clinically diagnosed TB	1/84	1.2% (0.0%-6.5%)	
	No TB diagnosis	3/153		98% (94%-99.5%)
Both samples	All children	18/259		
	Confirmed TB (SM+)	7/7	100% (59%-100%)	
	Confirmed TB (SM-)	6/15	40% (16%-68%)	
	Confirmed TB (all smear grades)	13/22	59% (36%-79%)	
	Unconfirmed, clinically diagnosed TB	1/84	1.2% (0.0%-6.5%)	
	No TB diagnosis	4/153		97% (93-99.3%)

[a]CI: confidence interval, [b]SM: smear

Factors associated with stool test positivity among TB cases

Features of adult-like TB (i.e., a positive smear result, a positive culture result, the ability to

spontaneously expectorate sputum and cavitary TB), were associated with detection in stool and

more common in the 11-14 year age group (Table 4). There was no association between stool

positivity and sex.

Table 4: Factors associated with *Mtb* stool assay positivity among children with culture-

confirmed or unconfirmed, clinically diagnosed TB (N=106).

Variable		Positive stool result (N=14) n (%)	Negative stool result (N=92) n (%)	P-value
Age group (years)	0-4	3 (21)	47 (51)	
	5-10	4 (29)	35 (38)	0.0008[a]
	11-14	7 (50)	10 (11)	
Female sex		7 (50)	51 (55)	0.8[b]
Positive smear result		7 (50)	0 (0.0)	<0.0001[b]
Positive culture result		13 (93)	9 (9.8)	<0.0001[b]
Cavitary disease (x-ray)		5 (36)	0 (0.0)	<0.0001[b]
Ability to produce sputum		12 (86)	33 (36)	0.0006[a]

[a]Chi-square test [b]Fisher's Exact test

<u>Discussion</u>

We found that our stool test was 100% sensitive for *Mtb* detection in stool among children with

smear-positive confirmed TB and 40% sensitive in children with smear-negative confirmed TB.

Importantly, we detected *Mtb* in stool of three children below the age of five that were not able

to expectorate sputum (two of whom were diagnosed based on GA samples and one who was

culture negative by sputum, but clinically diagnosed). The sensitivity (59%) and specificity

(97%) of this assay relative to liquid culture of respiratory samples was comparable to previous

studies using Xpert for stool testing in children (7,27). Unfortunately, similar to other studies

(6,28–30), we rarely detected TB in children with culture negative, clinically diagnosed TB.

Overall there is room for optimization to improve the sensitivity of smear-negative TB cases.

Considerations for assay optimization include testing of multiple sample aliquots, incorporating

an initial homogenization step and altering the sample volume. Our stool test protocol required a

relatively small input volume (500 mg), which may minimize the impact of PCR inhibitors in

stool that can impede the detection of *Mtb*. In fact, several studies show reduced sensitivity with

larger sample volumes. For example, Banada et al reported higher sensitivity and fewer invalid

test results when stool input volume was decreased from 1.2 to 0.6 grams (5). Recently, DiNardo

et al obtained 67% sensitivity with an in-house developed using only 50 mg, demonstrating that

even smaller volumes can be sufficient to reach high sensitivity.

In line with previous reports (8,31) we found strong associations between a positive stool test

and features of more adult-like TB (i.e., cavitary disease; positive sputum culture; positive

sputum smear; the ability to produce sputum), which were most common in the oldest age group.

This emphasizes the impact of age and smear status on the outcomes of stool testing and has

critical implications for the reporting of pediatric TB diagnostic studies because stool assay

sensitivities may vary widely based on the percentage of children in the cohort with adult-like

TB and/or high bacterial burden. Because children with adult-like TB are often able to produce a

sputum sample for testing, a stool assay will only begin to fill the pediatric TB diagnostic gap if

it is sensitive in children who cannot already spontaneously expectorate sputum. To facilitate an

understanding of stool assay performance and its utility across settings, we recommend

stratifying stool sensitivity results by some measure of disease severity, ideally smear status, and

reporting the added yield among children unable to spontaneously expectorate sputum.

The TruTip workstation had a number of attractive attributes for *Mtb* DNA extraction from stool.

Extraction procedures required limited pipetting steps compared to commercially available kits.

Additional future advantages include that the extracted DNA eluate can be isolated for other

purposes, such as sequencing. Most importantly, the TruTip workstation can be integrated with a

microarray-based detection of *Mtb* as well as a number of drug resistance-associated mutations

(17,19,20), which is a focus of future studies.

A limitation of our study was that we lacked follow-up data on the clinical evolution of children,

a key criterion for classifying pediatric cases of clinically diagnosed, unconfirmed TB. To the

extent that unconfirmed clinical TB was overdiagnosed, sensitivity will be underestimated in this

group. Similarly, missed TB diagnoses among children in whom TB was ruled out could lead to

an underestimate of assay specificity. The influence of these potential biases appears limited

given the very low sensitivity among children with unconfirmed clinical TB and a high overall

specificity. A second limitation of our study is the absence of children living with HIV. Studies

have shown that sensitivity of stool assays may be higher in children living with HIV (6,7), in

particular with severe immunosuppression (27), therefore, our results may not be generalizable to

children living with HIV.

Conclusion

In conclusion, the results of our study show that we can detect *Mtb* in pediatric stool samples.

For detection we used the TruTip workstation in combination with realtime IS6110 PCR, and its

performance is comparable to other platforms. Future work should include detection of

resistance in stool samples using the TruTip closed amplification system and assay optimization

to improve sensitivity in children with low bacillary loads.

Figure legends

Figure 1. Flowchart of cohort enrollment and stool study inclusion.

Abbreviations

AFB – acid fast bacilli; Ct – cycle threshold; DST – drug sensitivity testing; Mtb –

Mycobacterium tuberculosis; PCR – Polymerase Chain Reaction; TB – tuberculosis; TST –

tuberculin skin test; WHO – World Health Organization

Declarations

Ethics Approval and Consent to participate

Study participant guardians' provided written informed consent, and children eight years of age

and older provided written assent. All study procedures were approved by the Ethics Committee

of Peru's National Institute of Health and the Office of Human Research Administration at the

Harvard Medical School.

Consent for publication

Our manuscript does not contain individual person's data. Consent for publication was not

applicable.

Availability of data and material

Reported data are available on request to the corresponding author.

Competing interests

RH is an employee Akonni Biosystems inc. The company was not involved in study design.

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Author's contributions

AM: led the data analysis and interpretation and wrote the manuscript. MS: implementation of

the study and interpretation of results. JC: analysis and interpretation of results. JA:

implementation and data analysis. RC: study design and data analysis. NP: study design and

interpretation of results. MM: implementation of the study. FM: implementation of the study.

CM: implementation of the study. MM: study design and interpretation of results, LL: study

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results. MF conceptualized and designed the study, interpreted the results, and contributed to

writing and editing. All authors critically reviewed the manuscript and approved the final

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References

1.   World Health Organization. Global Tuberculosis Report. 2018.

2.   Starke JR. Pediatric tuberculosis: Time for a new approach. Tuberculosis. 2003;83(1–

3):208–12.

3.   Newton S, Brent A, Anderson S, Whittaker E, Kampmann B. Paediatric Tuberculosis.

Lancet Infect Dis. 2008;8(8):498–510.

4.   DiNardo AR, Detjen A, Ustero P, Ngo K, Bacha J, Mandalakas AM. Culture is an

imperfect and heterogeneous reference standard in pediatric tuberculosis. Tuberculosis.

2016;101:S105–8.

5.   Banada PP, Naidoo U, Deshpande S, Karim F, Flynn JL, O'Malley M, et al. A novel

sample processing method for rapid detection of tuberculosis in the stool of pediatric

patients using the Xpert MTB/RIF assay. PLoS One. 2016;11(3):1–13.

6.   Nicol MP, Spiers K, Workman L, Isaacs W, Munro J, Black F, et al. Xpert MTB/RIF

testing of stool samples for the diagnosis of pulmonary tuberculosis in children. Clin

Infect Dis. 2013 Aug;57(3):e18-21.

7.   Chipinduro M, Mateveke K, Makamure B, Ferrand RA, Gomo E. Stool Xpert MTB / RIF

test for the diagnosis of childhood pulmonary tuberculosis at primary clinics in

Zimbabwe. Int J Tuberc Lung Dis. 2017;21(2):161–6.

8.   Walters E, Scott L, Nabeta P, Demers A-M, Reubenson G, Bosch C, et al. Molecular

detection of mycobacterium tuberculosis from stools in young children by use of a novel

centrifugation-free processing method. J Clin Microbiol. 2018 Sep;56(9).

9. Walters E, Gie RP, Hesseling AC, Friedrich SO, Diacon AH, Gie RP. Rapid diagnosis of

pediatric intrathoracic tuberculosis from stool samples using the xpert MTB/RIF assay: A

pilot study. Pediatr Infect Dis J. 2012 Dec;31(12):1316.

10. Hasan Z, Shakoor S, Arif F, Mehnaz A, Akber A, Haider M, et al. Evaluation of Xpert

MTB/RIF testing for rapid diagnosis of childhood pulmonary tuberculosis in children by

Xpert MTB/RIF testing of stool samples in a low resource setting. BMC Res Notes. 2017

Sep;10(1):473.

11. Wolf H, Mendez M, Gilman RH, Sheen P, Soto G, Velarde AK, et al. Diagnosis of

pediatric pulmonary tuberculosis by stool PCR. Am J Trop Med Hyg. 2008

Dec;79(6):893–8.

12. DiNardo AR, Kay AW, Maphalala G, Harris NM, Fung C, Mtetwa G, et al. Diagnostic

and treatment monitoring potential of a stool-based quantitative polymerase chain reaction

assay for pulmonary tuberculosis. Am J Trop Med Hyg. 2018 Aug;99(2):310–6.

13. Oberhelman R a, Soto-Castellares G, Caviedes L, Castillo ME, Kissinger P, Moore D a J,

et al. Improved recovery of Mycobacterium tuberculosis from children using the

microscopic observation drug susceptibility method. Pediatrics. 2006;118(1):e100-6.

14. Denkinger CM, Dolinger D, Schito M, Wells W, Cobelens F, Pai M, et al. Target product

profile of a molecular drug-susceptibility test for use in microscopy centers. J Infect Dis.

2015;211(Suppl 2):S39–49.

15. Chandler DP, Griesemer SB, Cooney CG, Holmberg R, Thakore N, Mokhiber B, et al.

Rapid , simple influenza RNA extraction from nasopharyngeal samples. J Virol Methods.

2012;183(1):8–13.

16. Thakore N, Garber S, Bueno A, Qu P, Norville R, Villanueva M, et al. A bench-top

automated workstation for nucleic acid isolation from clinical sample types. J Microbiol

Methods. 2018;148(January):174–80.

17. Linger Y, Kukhtin A, Golova J, Perov A, Lambarqui A, Bryant L, et al. Simplified

microarray system for simultaneously detecting rifampin,isoniazid, ethambutol, and

streptomycin resistance markers in mycobacterium tuberculosis. J Clin Microbiol.

2014;52(6):2100–7.

18. Thakore N, Norville R, Franke M, Calderon R, Lecca L, Villanueva M, et al. Automated

TruTip nucleic acid extraction and purification from raw sputum. PLoS One. 2018;1–12.

19. Chandler D, Bryant L, Griesemer S, Gu R, Knickerbocker C, Kukhtin A, et al. Integrated

Amplification Microarrays for Infectious Disease Diagnostics. Microarrays.

2012;1(3):107–24.

20. Linger Y, Knickerbocker C, Sipes D, Golova J, Franke M, Calderon R, et al. Genotyping

Multi-Drug Resistant M. tuberculosis from Primary Sputum and Decontaminated

Sediment with an Integrated Microfluidic Amplification Microarray. J Clin Microbiol.

2018;(January):JCM.01652-17.

21. Kukhtin AC, Sebastian T, Golova J, Perov A, Knickerbocker C, Linger Y, et al. Lab-on-a-

Film disposable for genotyping multidrug-resistant Mycobacterium tuberculosis from

sputum extracts. Lab Chip. 2019; doi: 10.1039/C8LC01404C

22. Cuevas LE, Browning R, Bossuyt P, Casenghi M, Cotton MF, Cruz AT, et al. Evaluation

of tuberculosis diagnostics in children: 2. Methodological issues for conducting and

reporting research evaluations of tuberculosis diagnostics for intrathoracic tuberculosis in

children. Consensus from an expert panel. J Infect Dis. 2012;205(SUPPL. 2):209–15.

23.   MINSA. Norma técnica de la salud para el control de la Tuberculosis. Peru: Ministerio de

Salud. 2013.

24.   UNAIDS. UNAIDS Data 2017. 2017.

25.   Savelkoul PHM, Catsburg A, Mulder S, Oostendorp L, Schirm J, Wilke H, et al. Detection

of Mycobacterium tuberculosis complex with Real Time PCR□: Comparison of different

primer-probe sets based on the IS6110 element. J Microbiol Methods. 2006;66:177–80.

26.   Cole TJ. Too many digits: The presentation of numerical data. Arch Dis Child.

2015;100(7):608–9.

27.   Lacourse SM, Pavlinac PB, Cranmer LM, Njuguna IN, Mugo C, Gatimu J, et al. Stool

Xpert MTB/RIF and urine lipoarabinomannan for the diagnosis of tuberculosis in

hospitalized HIV-infected children. Aids. 2018;32(1):69–78.

28.   Walters E, van der Zalm MM, Palmer M, Bosch C, Demers A-M, Draper H, et al. Xpert

MTB/RIF on Stool Is Useful for the Rapid Diagnosis of Tuberculosis in Young Children

with Severe Pulmonary Disease. Pediatr Infect Dis J. 2017 Sep;36(9):837–43.

29.   Memon SS, Sinha S, Sharma S, Kabra S, Lodha R, Soneja M. Diagnostic accuracy of

xpert MTB/RIF assay in stool samples in intrathoracic childhood tuberculosis. J Tuberc

Ther. 2018;3(2).

30.   Orikiriza P, Nansumba M, Nyehangane D, Bastard M, Mugisha IT, Nansera D, et al.

Xpert MTB/RIF diagnosis of childhood tuberculosis from sputum and stool samples in a

high TB-HIV-prevalent setting. Eur J Clin Microbiol Infect Dis. 2018 Aug;37(8):1465–

73.

31.   Walters E, Demers A-M, van der Zalm MM, Whitelaw A, Palmer M, Bosch C, et al. Stool

culture for the diagnosis of pulmonary tuberculosis in children. J Clin Microbiol.

2017;55(12):JCM.00801-17.

Figure 1

^aStool samples from a number of TB cases had been used in pilot experiments (data not shown)