



Analytical and Clinical Evaluation of the Epistem Genedrive Assay for Detection of *Mycobacterium tuberculosis*

Shubhada Shenai,^a Derek T. Armstrong,^b Eloise Valli,^c David L. Dolinger,^c Lydia Nakiyingi,^d Reynaldo Dietze,^e Margareth Pretti Dalcolmo,^j Mark P. Nicol,^f Widaad Zemanay,^f Yuka Manabe,^{b,d} David Jamil Hadad,^e Patricia Marques-Rodrigues,^e Moises Palaci,^e Renata L. Peres,^e Mary Gaeddert,^g Sandra Armakovitch,^g Bareng A. S. Nonyane,^h Claudia M. Denkinger,^c Padmapriya Banada,^a Moses L. Joloba,ⁱ Jerrold Ellner,^g Catharina Boehme,^c David Alland,^a Susan E. Dorman,^b the TB Clinical Diagnostics Research Consortium

Division of Infectious Diseases, Center for Emerging Pathogens, Rutgers—New Jersey Medical School, Rutgers Biomedical & Health Sciences, Newark, New Jersey, USA^a; Johns Hopkins University School of Medicine, Baltimore, Maryland, USA^b; Foundation for Innovative New Diagnostics, Geneva, Switzerland^c; Infectious Disease Institute and Makerere University, Kampala, Uganda^d; Universidade Federal do Espirito Santo, Vitoria, Brazil^e; University of Cape Town and National Health Laboratory Service, Cape Town, South Africa^f; Boston Medical Center, Boston, Massachusetts, USA^a; Johns Hopkins Bloomberg School of Public Health, Baltimore, Maryland, USA^h; Makerere University Mycobacteriology Laboratory, Kampala, Ugandaⁱ; Centro de Referência Helio Fraga, FIOCRUZ, Rio de Janeiro, Brazilⁱ

The Epistem Genedrive assay rapidly detects the Mycobacterium tuberculosis complex from sputum and is currently available for clinical use. However, the analytical and clinical performance of this test has not been fully evaluated. The analytical limit of detection (LOD) of the Genedrive PCR amplification was tested with genomic DNA; the performance of the complete (sample processing plus amplification) system was tested by spiking M. tuberculosis mc²6030 cells into distilled water and M. tuberculosis-negative sputum. Specificity was tested using common respiratory pathogens and nontuberculosis mycobacteria. A clinical evaluation enrolled adults with suspected pulmonary tuberculosis, obtained three sputum samples from each participant, and compared the accuracy of the Genedrive to that of the Xpert MTB/RIF assay using M. tuberculosis cultures as the reference standard. The Genedrive assay had an LOD of 1 pg/ μ l (100 genomic DNA copies/reaction). The LODs of the system were 2.5 \times 10⁴ CFU/ml and 2.5×10^5 CFU/ml for cells spiked into water and sputum, respectively. False-positive *rpoB* probe signals were observed in 3/32 (9.4%) of the negative controls and also in few samples containing Mycobacterium abscessus, Mycobacterium gordonae, or Mycobacterium thermoresistibile. In the clinical study, among 336 analyzed participants, the overall sensitivities for the tuberculosis case detection of Genedrive, Xpert, and smear microscopy were 45.4% (95% confidence interval [CI], 35.2% to 55.8%), 91.8% (95% CI, 84.4% to 96.4%), and 77.3% (95% CI, 67.7% to 85.2%), respectively. The sensitivities of Genedrive and Xpert for the detection of smear-microscopy-negative tuberculosis were 0% (95% CI, 0% to 15.4%) and 68.2% (95% CI, 45.1% to 86.1%), respectively. The Genedrive assay did not meet performance standards recommended by the World Health Organization for a smear microscopy replacement tuberculosis test. Epistem is working on modifications to improve the assay.

Pulmonary tuberculosis (TB) is commonly diagnosed on the basis of patient history, clinical presentation, radiological findings, and sputum smear microscopy in high-TB-burden countries, but these approaches have limited sensitivity and specificity (1, 2). Culture-based diagnosis is more reliable but is labor-intensive, costly, and slow, which results in diagnostic delay even when available (3).

Epistem (Manchester, United Kingdom) developed and recently self-certified for CE-IVD a rapid molecular TB detection test that has the potential to speed up and simplify the diagnosis of pulmonary TB. This test uses a simple paper-based DNA extraction method coupled with PCR amplification and detection on Epistem's Genedrive instrument, a lightweight, portable, benchtop PCR platform with real-time PCR and melting temperature analysis capabilities. Epistem's Genedrive MTB/RIF (Genedrive) assay detects the Mycobacterium tuberculosis complex by targeting two different regions of the M. tuberculosis complex genome, a short repetitive region, rep13E12, and a segment of the rpoB gene (4-6). This test is CE marked for sale in European Economic Area (EEA) member states and is also licensed for distribution in India (7–9). However, this test's performance has not been extensively evaluated or reported in peer-reviewed journals apart from a single study published by the manufacturer and collaborators (6).

We assessed the analytical performance, biosafety, and diag-

nostic accuracy of the Genedrive assay for detection of *M. tuber-culosis*. In the analytical study, we used DNA and *M. tuberculosis* cells spiked into buffer or *M. tuberculosis*-negative human sputum samples. In a multicenter prospective diagnostic accuracy clinical study, adults with suspected pulmonary TB were recruited, and the accuracy of Genedrive was compared to that of Xpert MTB/

Received 23 October 2015 Returned for modification 18 November 2015 Accepted 16 December 2015

Accepted manuscript posted online 10 February 2016

Citation Shenai S, Armstrong DT, Valli E, Dolinger DL, Nakiyingi L, Dietze R, Dalcolmo MP, Nicol MP, Zemanay W, Manabe Y, Hadad DJ, Marques-Rodrigues P, Palaci M, Peres RL, Gaeddert M, Armakovitch S, Nonyane BAS, Denkinger CM, Banada P, Joloba ML, Ellner J, Boehme C, Alland D, Dorman SE, the TB Clinical Diagnostics Research Consortium. 2016. Analytical and clinical evaluation of the Epistem Genedrive assay for detection of *Mycobacterium tuberculosis*. J Clin Microbiol 54:1051–1057. doi:10.1128/JCM.02847-15.

Editor: G. A. Land

Address correspondence to Susan E. Dorman, dsusan1@jhmi.edu.

S.S. and D.T.A. contributed equally to this article.

Supplemental material for this article may be found at http://dx.doi.org/10.1128 /JCM.02847-15.

Copyright © 2016, American Society for Microbiology. All Rights Reserved.

RIF (Cepheid, Sunnyvale, CA) using a reference standard of mycobacterial culture.

MATERIALS AND METHODS

Ethics statement. The clinical study was approved by institutional review boards at Johns Hopkins Medicine, Boston University Medical Center, and each local study site. Participants provided written informed consent.

Analytic studies. The complete protocol of the analytic studies is described in the supplemental material.

Analytic sensitivity. Initial limit of detection (LOD) studies were performed with genomic M. tuberculosis DNA (ATCC 25618) to test the Genedrive PCR assay and the instrument without using the assay's sample processing cassette. For these studies, 19 µl of nuclease-free water and 1 µl diluted DNA were added to each PCR vial of the Genedrive PCR cartridge as described in the supplemental Materials and Methods. Studies designed to test the sample processing and PCR detection capacity of the Genedrive system used quantified M. tuberculosis mc²6030 cells. For studies using cells without spiking into sputum, 20 µl of each dilution was applied at the center of the sample loading cassette. Twenty microliters of cell dilution was also spiked into 480 µl of distilled water and was tested for the presence of *M. tuberculosis* using the Xpert assay. For spiking studies into sputum, deidentified discarded sputum samples were collected from patients who were not suspected of having TB (10). Thirty-two microliters of homogenized sputum was spiked with 8 μ l of the mc²6030 strain, where the last dilution was adjusted as needed to obtain the final cell concentration desired. Twenty microliters of this mixture was then applied to the center of the Genedrive sample loading cassette. Eight microliters of each cell dilution was also spiked into 492 µl of sputum and was tested for the presence of M. tuberculosis using the Xpert assay as described in the supplemental Materials and Methods. At least one positive and one negative control were included in each set of experiments.

For DNA and sputum spiking studies, at least 10 samples were tested at each dilution. The highest dilution with 100% positive test results was then identified, after which the test was repeated with the same dilution for a total of at least 10 additional times. If any negative test results were obtained, then the next lowest dilution was tested in the same manner to find the percentage of positivity at that concentration. The lowest dilution with at least 20 test replicates that had 100% positive tests was used to define the limit of detection (LOD).

Analytic specificity. Sixteen clinically occurring nontuberculosis mycobacterium (NTM) species (Table 1) and 11 Gram-positive and Gramnegative bacterial and fungal species (see Table S1 in the supplemental material) that are among those commonly reported as causes of respiratory infections or that have been identified as commonly isolated respiratory tract colonizers were analyzed at a concentration of 10⁷ CFU/ml. All strains were obtained from either the ATCC or the microbiology laboratory at the New Jersey Medical School, Rutgers Biomedical and Health Sciences, Rutgers University.

Biosafety studies. We also studied the extent to which M. tuberculosis present in sputum samples was killed using the Genedrive DNA extraction cassette. Different concentrations (107 to 105) of M. tuberculosis mc²6030 cells were applied onto the sample processing cassette directly or were first spiked into TB-negative sputum specimens and then applied to the cassette. The cassettes were allowed to stand at room temperature for 20 min. A 1 cm by 1 cm area of the membrane where the sample had been spotted in the center was cut out using a sterile scalpel and was inoculated into Bactec MGIT vials. Three different punches (1 mm by 1 mm in size) were also cut from the center of the membrane using a sterile disposable biopsy punch and were inoculated into MGIT vials. Similarly, 500 µl of 7H9 medium was added to the center of the sample loading cassette, the membrane was washed twice, and the membrane washings were inoculated in MGIT vials. All inoculated vials were incubated in a Bactec 320 MGIT instrument until they were flagged as positive by the instrument or remained negative for 42 days. Positive controls (*M. tuberculosis* mc²6030 cultures of various concentrations with and without plain membrane)

TABLE 1 Analytical specificity for nontuberculous mycobacteria panel

		Genedrive assay			
NTM species	ATCC no.	<i>rep13E12</i> target	<i>rpoB</i> target	Final result	
M. abscessus	19977	Negative	1/7 Positive	MTB detected Low	
M. asiaticum	25274	Negative	Negative	Undetected	
M. avium	15769	Negative	Negative	Undetected	
M. celatum	51131	Negative	Negative	Undetected	
M. chelonae	35749	Negative	Negative	Undetected	
M. fortuitum	35754	Negative	Negative	Undetected	
M. gordonae	14470	Negative	2/6 Positive	MTB detected Low	
M. intracellulare	35771	Negative	Negative	Undetected	
M. kansasii	12478	Negative	Negative	Undetected	
M. malmoense	29571	Negative	Negative	Undetected	
M. scrofulaceum	19073	Negative	Negative	Undetected	
M. smegmatis	Mc ² 155	Negative	Negative	Undetected	
M. simiae	25275	Negative	Negative	Undetected	
M. szulgai	23069	Negative	Negative	Undetected	
M. thermoresistibile	19527	Negative	1/3 Positive	MTB detected Low	
M. xenopi	19250	Negative	Negative	Undetected	

and negative controls (MGIT medium plus supplements plus *panD*) were incorporated in the study. Confirmation of *M. tuberculosis* growth in selected positive MGIT vials was performed by testing 1 ml of the culture using the Xpert assay.

Diagnostic accuracy clinical study. (i) Design, setting, and participants. A multicenter, cross-sectional, blinded diagnostic accuracy study was conducted. Participants were enrolled at outpatient clinics in Brazil, South Africa, and Uganda; hospitalized individuals were also enrolled in Uganda. Key inclusion criteria included an age of ≥ 18 years, a cough for ≥ 2 weeks, and one or more of the following: fever, night sweats, or weight loss. Individuals were excluded if they had received ≥ 2 days of anti-TB treatment within the prior 60 days. Each study participant was asked to provide a total of two sputum samples over days 1 and 2; a third sputum sample was collected within 7 days after enrollment (Fig. 1).

(ii) Mycobacteriology testing. Tests were performed at designated on-site labs, each of which had a quality assurance program in place. Lab staff was trained in Genedrive procedures by the manufacturer, and all laboratory tests were performed according to the manufacturer's protocol that is described in the supplemental material. Briefly, 20 µl of direct or processed sputum sample was placed on the Genedrive sample processing cassette and was allowed to dry at room temperature to extract nucleic acids from each sample. A 1-mm sterile disposable biopsy punch (Miltex, NY, USA) was used to cut 3 discs from the center of the cassette, and each disc was added individually to each of the 3 PCR vials of the Genedrive PCR cartridge. The sealed PCR cartridge was inserted in the Genedrive instrument, and the assay was run as per the manufacturer's instruction. Laboratory technologists performing Genedrive tests were blinded to the results of other tests for that specimen or participant. For each participant, the first sputum sample (sputum 1) was tested as a direct sample by fluorescence smear microscopy and Genedrive; the remaining portion of the sputum sample was then homogenized with 3-mm glass beads (Fisher Scientific, USA) and was tested using Xpert and Genedrive (Fig. 1). The second sputum sample (sputum 2) was digested and decontaminated using N-acetyl-L-cysteine (NALC)-NaOH (11). The resulting pellet was resuspended in 1.5 ml phosphate-buffered saline, mixed, and used for Genedrive (~50 µl), Lowenstein-Jensen culture (LJ; 200 µl inoculum), and MGIT culture (500 µl inoculum). Sputum sample 3 was digested and decontaminated using NALC-NaOH; the pellet was resuspended as above and was used to inoculate an LJ culture and an MGIT culture. Genedrive and Xpert testing were performed per the manufacturer's instructions. A participant was considered to be culture positive if any study culture was positive for growth of M. tuberculosis and culture negative if no culture

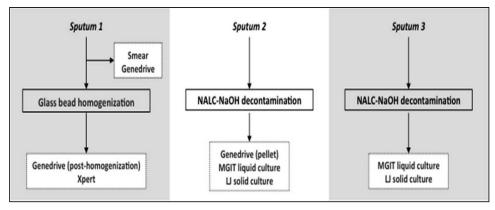


FIG 1 Specimen processing and testing schema for the prospective multicenter diagnostic accuracy clinical study. Each participant provided 3 sputum samples. Sputum 1 was used to perform direct smear microscopy and the Genedrive assay before being homogenized with glass beads. After homogenization, it was split into 2 portions for testing with the Genedrive assay and the Xpert MTB/RIF assay. Sputum 2 was decontaminated by the standard NALC-NaOH method, and then the resuspended pellet was used to perform liquid MGIT and solid Lowenstein-Jensen (LJ) cultures and the Genedrive assay. Sputum 3 was decontaminated by the standard NALC-NaOH method, and then the resuspended pellet was used to perform MGIT and LJ cultures.

grew *M. tuberculosis*. All cultures growing NTM or other bacterial contaminants were excluded from the data analysis. Similarly, specimens with missing results for any of the smear microscopy, culture, Genedrive, or Xpert assays were excluded from the analysis.

(iii) Statistical analysis. The LOD for the assay was analyzed by determining the percentage of positive tests for each concentration of *M. tuberculosis* isolates tested. The sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV) of the Genedrive assay with 95% confidence intervals (CIs) were estimated for prehomogenized and posthomogenized sputum specimens using a reference standard of 4 valid culture results using online statistical software MedCalc, SigmaPlot (7.0), and Stata (Stata Corp., TX). Similarly, the sensitivities, specificities, PPVs, and NPVs of the Xpert assay and the acid-fast bacilli (AFB) smear were also calculated against culture standard. Additional secondary outcomes were sensitivity and specificity of Genedrive performed on raw sputum using Xpert as the reference standard.

RESULTS

Performance of Genedrive PCR for detection of *M. tuberculosis* **genomic DNA.** The Genedrive PCR assay reproducibly detected as little as 1 pg of DNA per reaction, which is equivalent to approximately 100 genomic copies per reaction (Fig. 2A). One hundred femtograms of DNA (approximately 10 genomic copies) was detected with 45% positivity, and 10 fg of DNA (approximately 1 genomic copy) was detected with 8% positivity.

Dynamic range and LOD of M. tuberculosis cell dilutions spiked into water. M. tuberculosis cells were detected 100% of the time tested down to dilutions of 2.5×10^4 CFU/ml. This defined the Genedrive assay LOD for M. tuberculosis cells in a water matrix. The assay also detected 1×10^4 CFU/ml in 85% of the tests and 1×10^3 CFU/ml in 20% of the tests (Fig. 2B). As a comparator, tests performed with the Xpert assay on the same 20-µl sample volume of each concentration (brought up to a larger test volume with buffer as described in the Materials and Methods) detected 1×10^4 CFU/ml in 100% of tests and detected 1×10^3 CFU/ml in 60% of tests. We also tested the LOD of the Xpert assay using a 1-ml sample of each dilution (instead of the 20-µl sample of each dilution tested above) since this is the normal test volume for the Xpert assay. Using this normal test approach, the Xpert assay detected 250 CFU/ml in 100% of tests and detected 100 CFU/ml in 83.3% of tests, which is consistent with the previously reported Xpert LOD (10).

LOD of *M. tuberculosis* cell dilutions spiked into sputum. We next tested the LOD of the Genedrive using known *M. tuberculosis* sis-negative clinical sputum specimens spiked with known numbers of *M. tuberculosis* cells. The assay correctly identified *M. tuberculosis* in sputum samples containing 10^7 and 10^6 CFU/ml and 5×10^5 CFU/ml in 100% of all tests, defining the LOD in sputum as 5×10^5 CFU/ml (Fig. 2C). The assay also detected 2.5×10^5 CFU/ml in 90% of tests, 1×10^5 CFU/ml in 70% of tests, and 1×10^4 CFU/ml in 16.7% of tests. The Xpert assay tested with the same volume (8 µl) of 5×10^4 CFU/ml cell dilution spiked into 492 µl of sputum (i.e., approximately 400 CFU per Xpert cartridge) detected TB in 100% of tests and the 5×10^3 CFU/ml dilution (i.e., approximately 40 CFU per Xpert cartridge) in 91.7% of tests. Using 1-ml samples of each dilution, the Xpert assay showed an LOD of 250 CFU/ml in 100% of tests.

Cross-reactivity with NTM and other bacterial species. The *rep13E12*-targeting portion of the Genedrive assay did not show any cross-reactivity with nonmycobacterial respiratory pathogens or bacteria (Table 1; see also Table S1 in the supplemental material). However, false-positive *M. tuberculosis* results were observed due to a melting temperature (T_m) shift in the *rpoB* probe of three NTM species, *Mycobacterium abscessus*, *Mycobacterium thermoresistibile*, and *Mycobacterium gordonae* (Table 1; see also Fig. S1a to d in the supplemental material). False positive reports due to T_m shifts of the *rpoB* probe were also observed in 3 of 32 (9.4%) no-target or negative-control tests, of which one was in sputum matrix and two were with nonsputum negative controls (see Fig. S2 in the supplemental material).

Biosafety. *M. tuberculosis* growth occurred in all of the MGIT vials inoculated with an entire membrane, membrane washings, or three punches with cell concentrations ranging from 10^7 to 10^5 cells/ml (Table 2). Only the negative controls were culture negative. MGIT vials inoculated with the highest bacterial load, i.e., 10^7 cells/ml, took less time (6 to 8 days) to turn positive than the MGIT vials with the lowest load, i.e., 10^5 cells/ml of membrane (10 to 25 days).

Characteristics of participants in the clinical study. Between August 2014 and December 2014, 504 participants were enrolled. A total of 42 participants did not submit three sputum samples and were therefore excluded from the analysis (Fig. 3). An additional 80 participants did not have a Genedrive result for sputum

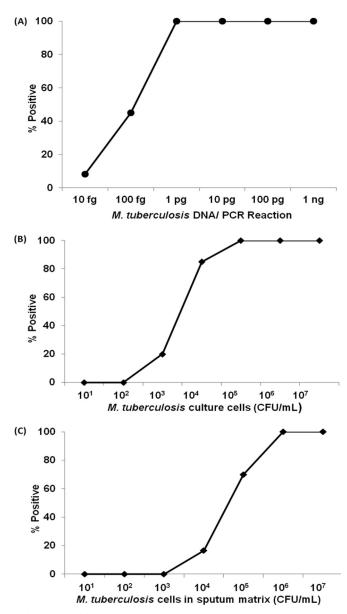


FIG 2 Limit of detection (LOD) of the Genedrive assay for detection of *M. tuberculosis*. LOD of Genedrive assay using (A) *M. tuberculosis* DNA. *M. tuberculosis* DNA at final concentration of 10 fg, 1pg, 10 pg, 100 pg, and 1 ng per PCR vial was loaded into Genedrive PCR cartridges and was processed according to the Genedrive protocol. (B) *M. tuberculosis* H37Rv mc²6030 quantified culture cells in water. *M. tuberculosis* mc²6030 quantified cells diluted in sterile water were tested by the Genedrive assay at final concentrations ranging from 10¹ to 10⁷ CFU/ml (n = 15); the percentage of *M. tuberculosis* H37Rv mc²6030 quantified culture cells in Sputum. *M. tuberculosis* mc²6030 cells were added to TB negative sputum to final concentrations of 10¹ to 10⁷ CFU/ml (n = 20) and were then processed according to the Genedrive assay protocol. The percentage of assays where *M. tuberculosis* was detected was then plotted for each concentrations of cells.

1, and 6 participants did not have an Xpert result for sputum 1. Forty participants were excluded from the analysis for other reasons per the protocol (Fig. 3). The resulting 336 participants comprised the main analysis population; 120 (36%) were enrolled in Brazil, 121 (36%) in Uganda, and 95 (28%) in South Africa. Of these patients, 202/336 (60%) were male, 60/336 (18%) were documented to be HIV-positive, and 120/336 (36%) had cavitation on chest radiograph (see Table S2 in the supplemental material). Among 336 analyzed participants, 97 (28.9%) had at least one culture that was positive for *M. tuberculosis*.

Clinical performance of single rapid tests versus culture. Using culture as a reference standard, the sensitivity for TB case detection of a single Genedrive assay prehomogenization (44/97 [45.4%]) was lower than that of Xpert (89/97 [91.8%]) (Table 3). The sensitivity of Genedrive performed prehomogenization was also lower than that of smear microscopy (75/97 [77.3%]) for a difference favoring smear microscopy of 32.0 percentage points (95% CI, 22.5 to 41.4). Genedrive failed to detect *M. tuberculosis* in any of 22 smear-negative/culture-positive specimens and detected *M. tuberculosis* in only 44/75 (58.7%) smear-positive/culture-positive specimens (Table 3). There was no difference in specificity between Genedrive performed prehomogenization (232/239 [97.5%]) and Xpert (234/239 [97.9%]).

Compared with the sensitivity of Genedrive performed prehomogenization on sputum 1 (44/97 [45.4%]), the sensitivities were higher for Genedrive performed posthomogenization on sputum 1 (57/97 [58.8%]; P = 0.004) and for Genedrive performed on the sputum 2 pellet (64/95 [67.4%]; P < 0.001). However, the sensitivity for TB case detection of a single Genedrive assay performed on the sputum 2 pellet remained lower than that of Xpert (a difference favoring Xpert of 24.4 percentage points [95% CI, 15.8 to 34.6]) and also significantly different from that of smear microscopy (difference favoring smear microscopy of 9.9 percentage points [95% CI, 3.5 to 24.4]).

DISCUSSION

We report the first comprehensive evaluation of the Epistem Genedrive assay for detection of M. tuberculosis. The sensitivity of Genedrive was low compared to other methods, and this finding was consistent across the analytic studies using sputum spiked with *M. tuberculosis* and a clinical study using fresh sputum from adults with signs/symptoms of pulmonary TB. Analytic study results indicated that the observed suboptimal sensitivity can be traced to the sample processing component of the assay. The amplification portion of the assay appeared to perform well, with an LOD of 100 genomes when tested with purified M. tuberculosis DNA that did not require any additional sample processing. Therefore, it is likely that the assay's limited performance is due to the relatively small amount of sample that is processed on the paper cassette and then placed into the assay amplification vial. Insufficient cell lysis or insufficient removal of PCR inhibitors from sputum by the cassette may also adversely affect sensitivity, although we did not specifically study these parameters. Our results differ markedly from the only other paper published on this assay to date (6).

The Genedrive assay did not show any positive results when tested with common Gram-positive and Gram-negative respiratory pathogens. However, there were false-positive *rpoB* peaks when tested with *M. abscessus*, *M. gordonae*, and *M. thermoresistibile*. False-positive results for *M. tuberculosis* detection were also observed in 9% of negative controls, although the reasons are unclear. A test design concern is that the assay internal control is run in a different assay tube than the two *M. tuberculosis* detection tubes. This control format may make it difficult to detect tube-

Inoculum (CFU/ml)	Amount of material inoculated into MGIT vial ^a	MGIT culture	Time to positivity	Bacterial load/ MGIT vial	Confirmation $(C_T)^b$
107	3 punches ^c (replicate a)	Positive	7 days, 1 h	3×10^4	Xpert MTB (17.8)
10 ⁷	3 punches ^{<i>c</i>} (replicate b)	Positive	8 days, 10 h	3×10^4	
107	Entire membrane (1 by 1 cm)	Positive	6 days, 1 h	1×10^{5}	
10 ⁷	Washings from SLC^d	Positive	8 days, 10 h		
10 ⁷	20- μ l culture + plain membrane	Positive	4 days, 10 h	2×10^5	
107	20-µl culture (positive control)	Positive	5 days	2×10^{5}	
10 ⁶	3 punches ^c (replicate a)	Positive	9 days, 13 h	3×10^3	Xpert MTB (22.3)
10 ⁶	3 punches ^c (replicate b)	Positive	12 days, 7 h	3×10^3	· · ·
10 ⁶	Entire membrane (1 by 1 cm)	Positive	7 days, 17 h	1×10^4	Xpert MTB (18.2)
10 ⁶	Washings from SLC	Positive	9 days, 9 h		• · · ·
10 ⁶	20- μ l culture + plain membrane	Positive	7 days	$2 imes 10^4$	
10 ⁵	3 punches ^c (replicate a)	Positive	25 days, 1 h	3×10^{2}	
10 ⁵	3 punches ^c (replicate b)	Positive	25 days, 13 h	3×10^2	
10 ⁵	Entire membrane (1 cm by 1 cm)	Positive	10 days, 11 h	1×10^{3}	Xpert MTB (20.1)
10 ⁵	20 µl Culture (positive control)	Positive	7 days, 17 h	2×10^{3}	÷ , ,
	Negative control (replicate a)	Negative	No growth at the end	of 42 days	
	Negative control (replicate b)	Negative			

TABLE 2 Biosafety experiment: ability to detect viability of M. tuberculosis after paper-based DNA extraction

^a Experiments conducted in duplicate are labeled as replicate a and b.

^b C_T, Xpert cycle threshold defined as the number of cycles required for the fluorescent signal to cross the threshold (i.e., exceed background level).

^c Three punches were cut from the center of the Genedrive sample loading cassette and inoculated into the MGIT vial.

^d SLC, sample loading cassette.

specific reagent issues or uneven distribution of PCR inhibitors from the test sample.

in viable *M. tuberculosis* CFU sufficient to make the sample biosafe. At least a six-log reduction in viability is generally required to consider that a procedure has sterilizing ability (12, 13). We recommend that biosafety level 3 (BSL3) lab facilities or at least a

The analytical component of our study demonstrated that the Genedrive sample-processing cassette did not result in a reduction

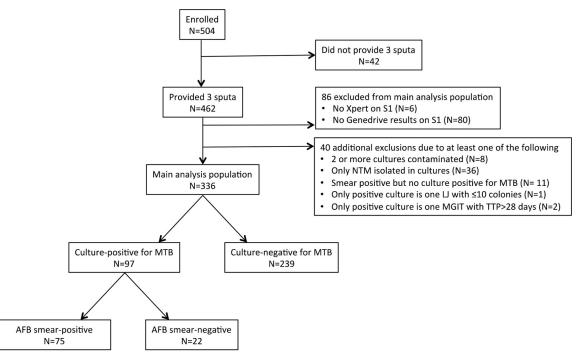


FIG 3 Overview of patient enrollment and study outcomes. A total 504 suspected pulmonary TB patients were recruited in this study, of which 42 patients were unable to produce sputum specimens. Of the remaining 462, 80 participants did not have a determinate Genedrive result on sputum 1 (12 with test result of "error" and 68 with Genedrive testing not performed due to instrument inoperability or no intact cartridges at the testing site), and 6 participants did not have a determinate Xpert result for sputum 1 (2 with test result of 'error' and 4 with Xpert testing not performed due to human error) and hence were excluded from the study. Of the 336 participants who completed this study, 97 were culture positive and 239 were culture negative. Among culture positives, 75 were AFB positive and 22 were AFB negative by smear microscopy. S1, sputum 1; NTM, nontuberculous mycobacteria; MTB, *M. tuberculosis*; LJ, Lowenstein-Jensen culture; MGIT, mycobacterial growth indicator tube culture; TTP, time to positivity; AFB, acid-fast bacilli.

	Compared to 4 cultur specimens					
Prospective diagnostic accuracy	Sensitivity (n/n [%] [95% CI])	Specificity (n/n [%] [95% CI])	PPV (n/n [%] [95% CI])	NPV (n/n [%] [95% CI])	AFB Sm ⁺ /Cx ⁺ , Sensitivity (n/n [%] [95% CI]) ^{<i>a</i>}	AFB Sm ⁻ /Cx ⁺ , Sensitivity (n/n [%] [95% CI])
Sputum 1 result vs culture gold standard ($n = 336$ participants)						
Genedrive, prehomogenization, sputum 1	44/97 (45.4) (35.2–55.8)	233/239 (97.5) (94.6–99.1)	44/50 (88.0) (75.7–95.5)	233/286 (81.5) (75.7–85.8)	44/75 (58.7) (46.7–70.0)	0/22 (0) (0-15.4)
Genedrive, posthomogenization, sputum 1	57/97 (58.8) (48.3–68.7)	233/239 (97.5) (94.6–99.1)	57/63 (90.5) (80.4–96.4)	233/273 (85.3) (80.6–89.3)	56/75 (74.7) (63.3–84.0)	1/22 (4.5) (0.0–22.8)
Xpert MTB/RIF, sputum 1	89/97 (91.8) (84.4–96.4)	234/239 (97.9) (95.2–99.3)	89/94 (94.7) (88.0–98.3)	234/242 (96.7) (93.6–98.6)	74/75 (98.7) (92.8–99.9)	15/22 (68.2) (45.1–86.1)
AFB smear microscopy, sputum 1	75/97 (77.3) (67.7–85.2)	239/239 (100) (98.5–100)	75/75 (100) (95.1–100)	239/261 (91.6) (87.5–94.6)	N/A	N/A
Sputum 2 result vs culture gold standard $(n = 326 \text{ participants})$						
Genedrive, pellet, sputum 2	64/95 (67.4) (57.0–76.6)	225/231 (97.4) (94.4–99.0)	64/70 (97.0) (81.5–96.6)	225/256 (87.9) (83.3–91.6)	63/74 (85.14) (74.9–92.3)	1/21 (4.7) (0–23.8)

TABLE 3 Diagnostic accuracy clinical study: accuracy of single rapid tests using as a gold standard four cultures performed from two sputum samples

^a Cx, culture; Sm, smear; N/A, not available.

BSL2 facility with a biosafety-2 cabinet be used to process samples for this assay until additional work can be done to confirm that assay procedures are biosafe. Sample processing cassettes should be treated with 20% bleach or another acceptable decontaminant for 20 to 30 min before being discarded.

Our analytic LOD study was critically dependent on proper mixing and dispersion of the spiked *M. tuberculosis* cells into the test matrix. Only a small (20 μ l) volume of this matrix was applied to the processing cassette, and even smaller portions (two small punches) were placed into the assay amplification tubes. Therefore, poor mixing of the sample may have resulted in some 20- μ l aliquots having unexpectedly low numbers of spiked *M. tuberculosis* cells. However, our procedures addressed this issue as much as possible by vortexing and sonicating samples for homogenization. Furthermore, the Xpert assay was able to detect *M. tuberculosis* from dilution aliquots containing as little as 10³ CFU/ml when we placed the same volume of cell suspension (20 μ l) in the Xpert and on the Genedrive sample processing cassette. This parallel study with the Xpert assay demonstrated that *M. tuberculosis* was detectable even from the 20- μ l aliquots.

Findings from the diagnostic accuracy clinical study confirmed the low sensitivity observed in the analytic study. Genedrive sensitivity was inferior to that of Xpert overall and was also substantially lower than that of smear microscopy. Genedrive's sensitivity was well below minimum targets (60% for smear-negative/culture-positive TB and 99% for smear-positive/culture-positive TB) recommended by the World Health Organization for high-priority new TB diagnostics (14). It is worth noting, however, that the observed Genedrive performance generally met or exceeded the WHO targets with respect to specificity, number of assay steps, time to result, and instrument and power requirements.

In the clinical study, the sensitivity of Genedrive was lower in direct sputum specimens tested before homogenization versus those tested after homogenization and was highest when NALC-NaOH-processed sediment was used. This is likely a consequence of heterogeneous distribution of bacilli in nonmanipulated sputum, a challenge also encountered in the analytic study. While it is encouraging that sensitivity can be increased through sputum manipulation, a requirement for sputum manipulation would nevertheless be a disadvantage in practice and limit the test's use at point-of-care. A limitation of the current study is that we did not assess the ability of the Genedrive assay to detect genotypic changes associated with rifampin resistance.

In summary, results from analytical and clinical studies showed low Genedrive sensitivity for the detection of *M. tuberculosis*. There was cross-reactivity with three nontuberculosis mycobacterial species, and the sample-processing cassette did not result in a reduction in viable *M. tuberculosis* CFU sufficient to make the sample biosafe. The Genedrive assay is undergoing further development, and the test developers are aiming to address the different issues highlighted in this work.

ACKNOWLEDGMENTS

We thank Epistem Holdings Plc for donation of instruments and supplies for the Genedrive assay. We thank the clinical study participants, as well as the TB-CDRC Data Coordinating Center.

In addition to the authors, the contributions of the following individuals are gratefully acknowledged: Carolyn Namaganda, Eric Bugumirwa, and Carol Kayiza (Infectious Disease Institute and Makerere University, Kampala, Uganda); Luiz Guilherme Schmidt, Paolo Poloni, and Pedro Sousa (Nucleo de Doencas Infecciosas, Universidade Federal do Espirito Santo, Vitoria, Brazil); Slindile Mbele, Layla Hendricks, Michelle Mento, and Judi van Heerden (University of Cape Town, Cape Town, South Africa); and David Hom and Rachel Kubiak (Boston University Medical Center, Boston, Massachusetts, USA).

This work was supported by the Foundation for Innovative New Diagnostics and also by the National Institute of Allergy and Infectious Diseases, National Institutes of Health, Department of Health and Human Services under contract HHSN2722000900050C, "TB Clinical Diagnostics Research Consortium." Additional support was provided by NIH to S.E.D. under contract K24AI104830.

D.A. is one of a group of coinvestigators who invented molecular beacon technology and who receive income from licensees, including a license to Cepheid for *M. tuberculosis* detection. To manage this conflict of interest, the income attributable to the Xpert MTB/RIF assay, which he may receive, has been irrevocably capped at \$4,999 per year. D.A. also reports receiving two research contracts from Cepheid. E.V., C.M.D., D.L.D., and C.B. are employed by FIND, a nonprofit organization that

collaborates with industry partners, including BD, Cepheid, and Hain Lifesciences among others for the development, evaluation, and demonstration of new diagnostic tests for poverty-related diseases. M.P.N. and the University of Cape Town have received funding from Epistem (subsequent to the conduct of the study described here) to test a modified version of the assay. The remaining authors declare no conflict of interest.

FUNDING INFORMATION

The National Institutes of Health provided funding to the TB Clinical Diagnostics Research Consortium under grant number HHSN2722000900050C. The National Institutes of Health provided funding to Susan E. Dorman under grant number K24AI104830. The Foundation for Innovative New Diagnostics provided funding to Shubhada Shenai, Mark P. Nicol, Widaad Zemanay, Padmapriya Banada, and David Alland.

This work was supported by the Foundation for Innovative New Diagnostics and also by the National Institute of Allergy and Infectious Diseases, National Institutes of Health, Department of Health and Human Services, under contract no. HHSN2722000900050C, "TB Clinical Diagnostics Research Consortium." Additional support was provided by NIH K24AI104830 to S.E.D.

REFERENCES

- 1. Hobby GL, Holman AP, Iseman MD, Jones JM. 1973. Enumeration of tubercle bacilli in sputum of patients with pulmonary tuberculosis. Antimicrob Agents Chemother 4:94–104. http://dx.doi.org/10.1128 /AAC.4.2.94.
- Harries AD, Dye C. 2006. Tuberculosis. Ann Trop Med Parasitol 100: 415–431.
- 3. Urbanczik R, Rieder HL. 2009. Scaling up tuberculosis culture services: a precautionary note. Int J Tuberc Lung Dis 13:799–800.
- Lee TY, Lee TJ, Belisle JT, Brennan PJ, Kim SK. 1997. A novel repeat sequence specific to Mycobacterium tuberculosis complex and its implica-

tions. Tuber Lung Dis 78:13–19. http://dx.doi.org/10.1016/S0962-8479 (97)90011-3.

- 5. Gordon SV, Heym B, Parkhill J, Barrell B, Cole ST. 1999. New insertion sequences and a novel repeated sequence in the genome of *Mycobacterium tuberculosis* H37Rv. Microbiology 145:881–892.
- Castan P, de Pablo A, Fernandez-Romero N, Rubio JM, Cobb BD, Mingorance J, Toro C. 2014. Point-of-care system for detection of *My-cobacterium tuberculosis* and rifampin resistance in sputum samples. J Clin Microbiol 52:502–507. http://dx.doi.org/10.1128/JCM.02209-13.
- 7. Epistem. 2012. Epistem press release. Epistem signs tuberculosis channel partner agreement. Epistem Plc., Manchester, UK. http://www.epistem.co.uk /press-releases/EpistemPressReleaseTBCollaborationAgreement05Mar12 .pdf.
- DDNews. August 2011. Epistem and Xcelris Labs partner on TB diagnostics. DDNews, Rocky River, OH. http://www.ddn-news.com/index.php ?newsarticle=5251.
- Labmate. 15 May 2015. India approves licence for TB diagnosis. Labmate, Hertfordshire, UK. http://www.labmate-online.com/news/news-and-views/ 5/epistem_ltd/india_approves_licence_for_tb_diagnosis/34643/.
- Helb D, Jones M, Story E, Boehme C, Wallace E, Ho K, Kop J, Owens MR, Rodgers R, Banada P, Safi H, Blakemore R, Lan NT, Jones-Lopez EC, Levi M, Burday M, Ayakaka I, Mugerwa RD, McMillan B, Winn-Deen E, Christel L, Dailey P, Perkins MD, Persing DH, Alland D. 2010. Rapid detection of *Mycobacterium tuberculosis* and rifampin resistance by use of on-demand, near-patient technology. J Clin Microbiol 48:229–237. http://dx.doi.org/10.1128/JCM.01463-09.
- 11. Kent PT, Kubica GP. 1985. Public health mycobacteriology, a guide for level III laboratory. Centers for Disease Control, Atlanta, GA.
- Rutala WA, Weber DJ. 2001. New disinfection and sterilization methods. Emerg Infect Dis 7:348–353. http://dx.doi.org/10.3201/eid0702.010241.
- 13. Association for the Advancement of Medical Instrumentation. 2003. Sterilization of health care products: requirements for products labled "sterile." Standard ANSI/AAMI ST67. Association for the Advancement of Medical Instrumentation, Arlington, VA.
- 14. World Health Organization. 2014. High-priority target product profiles for new tuberculosis diagnostics: report of a consensus meeting. *In* Proceedings of the WHO/HTM/TB/2014.18. WHO, Geneva, Switzerland.