Microbiological diagnosis of pulmonary tuberculosis in children by oral swab polymerase chain reaction

Mark P. Nicol1,2, Rachel C. Wood3, Lesley Workman4, Margaretha Prins4, Cynthia Whitman4, Yonas Ghebrekristos2, Slindile Mbhele1, Alaina Olson3, Lisa E. Jones-Engel5, Heather J. Zar4 & Gerard A. Cangelosi3

Microbiological diagnosis of pediatric pulmonary tuberculosis (TB) is challenging due to the difficulty of collecting and testing sputum from children. We investigated whether easily-obtained oral swab samples are useful alternatives or supplements to sputum. Oral swabs and induced sputum (IS) were collected from 201 South African children with suspected pulmonary TB. IS samples were tested by mycobacterial culture and Xpert MTB/RIF. Oral swabs were tested by PCR targeting IS6110. Children were categorized as Confirmed TB (microbiologic confirmation on IS), Unconfirmed TB (clinical diagnosis only), or Unlikely TB (recovery without TB treatment). Relative to Confirmed TB, PCR on two oral swabs per child was 43% sensitive and 93% specific. This sensitivity fell below that of sputum Xpert (64%). Among children with either Confirmed or Unconfirmed TB, PCR on two oral swabs per child was 31% sensitive and 93% specific, which was more sensitive than sputum testing among this group (21%). Although oral swab analysis had low sensitivity in sputum-positive children, it detected TB in a significant proportion of sputum-negative children who were clinically diagnosed with TB. Specificity at 93% was suboptimal but may improve with the use of automated methods. With further development, oral swabs may become useful supplements to sputum as samples for diagnosis of pulmonary TB in children.

Microbiological confirmation of pulmonary tuberculosis (TB) in children remains challenging despite recent advances1. Collection of specimens for testing, such as induced sputum (IS) or gastric aspirate, is perceived to be invasive and complex. Even when such specimens are obtained and tested, the diagnostic yield can be suboptimal, with microbiological confirmation achieved in only half of children diagnosed with pulmonary TB, even in well-resourced research studies2. Improved microbiological confirmation is important, not only for clinical care but also because assessment of novel diagnostic biomarkers for TB in children is severely constrained by the lack of a sensitive reference standard test.

One approach to improving TB diagnosis in children is to collect multiple different specimen types with the assumption that the yield would be cumulative. Potentially useful specimen types include gastric aspirates, induced sputum, nasopharyngeal aspirates, stool, urine, and the ‘string test’3–6. One of the simplest samples to obtain is a swab of the oral mucosa7. Originally demonstrated for diagnosis of TB in non-human primates8,9, we have shown that oral swab PCR could detect approximately 90% of microbiologically confirmed cases of pulmonary TB in adults10,11. As potential samples for pulmonary TB testing in young children, oral swabs (OS) have the advantage of being universally obtainable with minimal discomfort or need for operator training. We therefore...
investigated the diagnostic accuracy and incremental yield (over IS testing) of oral swab PCR for *Mycobacterium tuberculosis* DNA in children presenting to hospital with suspected pulmonary tuberculosis.

**Materials and Methods**

In a prospective study, we enrolled 201 children (<15 years of age) presenting to Red Cross War Memorial Children's Hospital, Cape Town, South Africa with suspected pulmonary TB between 13th November 2016 and 22nd September 2017. Enrolment criteria were cough of any duration plus one of: household contact with a TB source case within the preceding 3 months, loss of weight or failure to gain weight in the preceding 3 months, a positive tuberculin skin test (defined as ≥10 mm in HIV-uninfected children and ≥5 mm in HIV-infected children), or a chest radiograph suggestive of pulmonary TB. We excluded children if they had received more than 72 hours of TB treatment, were not resident in Cape Town, or if informed consent from a parent or guardian (and assent from older children) was not obtained.

Children provided two consecutive induced sputum specimens as previously described. Following decontamination with sodium hydroxide, centrifuged sputum deposits were re-suspended in 1.5 ml of phosphate buffer. 0.5 ml of the re-suspended pellet was used for automated liquid culture (BACTEC MGIT, Becton Dickinson, Cockeysville, MD, USA). Positive cultures were identified by MTBDRplus testing (Hain Lifescience, Nehren, Germany). For the 1st induced sputum, a further aliquot of 0.5 ml was used for Xpert MTB/RIF testing, as per manufacturer's recommendations.

TB diagnostic categorization was based on clinical and microbiological investigations and follow-up visits, in line with revised NIH consensus definitions. 'Confirmed TB' (any induced sputum culture or Xpert MTB/RIF positive for *M. tuberculosis*), 'Unlikely TB' (microbiologically negative, no TB treatment given and documented resolution of symptoms and signs at 3 month follow-up visit), or 'Unconfirmed TB' (all other children, including (i) children clinically diagnosed who were placed on TB treatment and (ii) children who were not placed on TB treatment and who had persistent symptoms and signs at follow-up).

Prior to sputum collection, two oral swabs (OS1 and OS2) were collected from each child, from the inside of the left cheek and the inside of the right cheek, respectively. In most cases the swabs were collected at the same time, one immediately after the other. In some cases they were collected up to 3 days apart. Swab collection always preceded sputum collection. GE Healthcare OmniSwabs or Puritan PurFlock swabs were used to scrape the buccal surface for 10 seconds. Swabs were immediately placed in a sterile, in-house lysis buffer for storage and frozen at −80°C.

DNA was extracted from the swab samples by using a modified Qiagen protocol described previously. Quantitative PCR targeting IS6110, a multi-copy insertion element unique to the *M. tuberculosis* complex, was done as described. DNA extraction and qPCR were conducted in a 3-room laboratory suite with one-way workflow to minimize sample contamination. Laboratory-generated negative control swabs were extracted alongside sets of 10 OS samples and additional negative template controls were included in qPCR reactions. PCR results were categorized as positive or negative using a predetermined Cq cut-off of less than 40. The persons doing the PCR were blinded to reference standard test results and clinical information. Likewise, index test results and clinical information were not available to microbiology staff performing the reference standard testing, and the index test results were not available to the data analyst responsible for diagnostic categorization of children.

The attending physician, who made the decision whether to start TB therapy, had access to all laboratory results except the oral swab PCR. All enrolled children attended follow-up visits at 1- and 3-months post-enrolment. Children on TB treatment attended an additional follow-up visit at 6-months post-enrolment. Research staff assessed response to treatment at follow-up by recording symptoms, signs and weight gain. The study was approved by the Research Ethics Committee of the Faculty of Health Sciences, University of Cape Town. All procedures were performed in accordance with the relevant guidelines and regulations. Informed consent was obtained through parents or guardians. No identifying information is presented in this report.

**Results**

Of the 201 children enrolled with suspected pulmonary TB, 36 (18%) were excluded from further analysis for the following reasons: no second oral swab collected (n = 4), confirmed extrapulmonary TB only (n = 15), no valid induced sputum culture result (n = 11), or not started on TB treatment and did not return for follow-up assessment (n = 6). Of the 165 children included in the analysis, the median age was 31 (IQR 13–81) months, 87 (53%, 95% CI 45–61) were female, 18 (11%, 95% CI 7–17) were HIV-infected, 84/137 (61%, 95% CI 53–70) had a positive TST, and 121 (73%, 95% CI 66–80) were started on TB treatment. Forty children (24%, 95% CI 18–32) were classified as confirmed TB, 81 (49%, 95% CI 41–57) as unconfirmed TB, and 44 (27%, 95% CI 20–34) as unlikely TB. The characteristics of the 165 children included in the analysis are detailed in Table 1. For 116 of the 165 children, the two swabs were collected at the same time, one immediately after the other. They were collected at different times for 49 of the children, ranging from 1 minute to 3 days apart. A valid Xpert MTB/RIF result from an IS sample was available in 154/165 (93%) children.

We first calculated test accuracy (sensitivity and specificity) using children with Confirmed TB as the positive reference standard and those with Unlikely TB as the negative reference standard. As a secondary analysis we used a composite reference standard, combining the groups of Confirmed TB and Unconfirmed TB as the positive reference standard. We determined the accuracy of PCR on one oral swab (OS1, the first collected) and two oral swabs (OS1 and OS2, the first two collected). Reporting is as recommended by STARD guidelines for diagnostic accuracy studies. With an estimated proportion of children with Confirmed TB of 15%, this sample size was predicted to give 95% confidence intervals of 63–90% around a point estimate of 80% for oral swab PCR sensitivity.

Of the 165 children included in the analysis, 23 (14%) had a positive PCR on OS1, 27 (16%) had a positive PCR on OS2, and 39 (23%) had a positive PCR on either OS1 or OS2 (Table 2). Among those children with
Confirmed TB, OS1 or OS2 were positive in 17/40 (43%). Among children with Unconfirmed TB, OS1 or OS2 were positive in 19/81 (24%). Among children with Unlikely TB, OS1 or OS2 were positive in 3/44 (7%). Oral swab positivity was significantly more frequent among children with Unconfirmed TB than among children with Unlikely TB (p = 0.020 by two-tailed z test).

Comparing OS results to Confirmed TB and Unlikely TB (Table 3), PCR on one oral swab (OS1) had a sensitivity of 30% (95% CI 17–47) and a specificity of 100% (95% CI 92–100). PCR on two oral swabs had a sensitivity of 43% (95% CI 27–59) and specificity of 93% (95% CI 81–99). For PCR on two OS, the positive likelihood ratio was 6.2 (95% CI 2.0–20) and the negative likelihood ratio was 0.6 (95% CI 0.5–0.8). By comparison, sensitivity of a single Xpert MTB/RIF test on induced sputum was 64% (95% CI 46–79). Although Confirmed TB was defined as any sputum culture-positive or Xpert MTB/RIF-positive for *M. tuberculosis* (per NIH consensus definitions), no Xpert MTB/RIF-positive, culture-negative cases were observed within the study population. Therefore, a secondary analysis was used to quantify the specificity of sputum Xpert MTB/RIF relative to sputum culture. This value was 100% (95% CI 92–100). The numbers of HIV-infected children were too small for assessment of HIV vs. non-HIV.

When OS positivity was compared to the combined group of Confirmed plus Unconfirmed TB (Table 3), in those 154 children with valid OS1, OS2, and induced sputum Xpert MTB/RIF results, the sensitivity of PCR on OS1 was 19% (21/111, 95% CI 12–27) and of PCR on OS1 or OS2 was 31% (34/111, 95% CI 22–40). By comparison, a single Xpert MTB/RIF test was positive in 21% (23/111, 95% CI 14–29) of children within this group.

Table 1. Characteristics of children. *Tuberculin skin test results available for a subset of the study population (N = 137).

| Age months | Median (IQR) |
|------------|-------------|
| <5 years   | 30.5 (13.3–80.8) |
| ≥5 years   | 23.0 (13.5–86.4) |

| Age category | Median (IQR) |
|--------------|-------------|
| <5 years     | 113 (68.5) |
| ≥5 years     | 52 (31.5) |

| Male sex | Median (IQR) |
|----------|-------------|
| 78 (47.3%) |

| HIV infected | Median (IQR) |
|--------------|-------------|
| 18 (10.9%) |

| Started on TB treatment | Median (IQR) |
|-------------------------|-------------|
| 121 (73.3%) |

| Tuberculin skin test positive | Median (IQR) |
|------------------------------|-------------|
| 84 (61.3%) |

Table 2. Number of positive tests, by TB diagnostic category and test type. OS1, first oral swab PCR; OS2, second oral swab PCR. IS, induced sputum. Of 36 IS culture-positive subjects, 23 (64%) were positive on culture and Xpert, and 13 (36%) were positive on culture only. 154 samples were tested by Xpert MTB/RIF.

| Sensitivity reference standard | Index method | Sensitivity n (%) (95% CI) | Specificity n (%)a (95% CI) | Positive LRb (95% CI) | Negative LR (95% CI) |
|-------------------------------|-------------|-----------------------------|-----------------------------|---------------------|---------------------|
| Confirmed TB                  | OS1         | 12/40 (30%) (17–47)         | 44/44 (100) (92–100)        | NA                  | 0.7 (0.6–0.9)       |
| Confirmed + Unconfirmed TB    | OS1 OR OS2  | 17/40 (43%) (27–59)         | 41/44 (93) (81–99)          | 6.2 (2–20)          | 0.6 (0.5–0.8)       |
| Confirmed + Unconfirmed TB    | IS Xpert MTB/RIF | 23/36 (64%) (46–79) | 43/43 (100) (92–100)        | NA                  | 0.4 (0.2–0.6)       |
| Confirmed TB                  | IS Xpert MTB/RIF | 21/111 (19%) (12–27) | 43/43 (100) (92–100)        | NA                  | 0.8 (0.7–0.9)       |
| Confirmed + Unconfirmed TB    | IS Xpert MTB/RIF | 34/111 (31%) (22–40) | 40/43 (93) (81–99)          | 4.4 (1–14)          | 0.8 (0.6–0.9)       |
| Confirmed + Unconfirmed TB    | IS Xpert MTB/RIF | 23/111 (21%) (14–29) | 43/43 (100) (92–100)        | NA                  | 0.8 (0.7–0.9)       |

Table 3. Accuracy of one or two oral swab PCR tests when using two different positive reference standards and Unlikely TB as the negative reference standard. OS1, first oral swab PCR; OS2, second oral swab PCR. IS, induced sputum. Positive LR could not be calculated when specificity was 100%.
Discussion

Although PCR of oral swabs had low sensitivity for identification of children with culture-confirmed pulmonary TB, the method was positive in a substantial proportion of children who were clinically diagnosed with TB but had negative mycobacterial culture of induced sputum (Unconfirmed TB). This proportion was significantly greater than that seen in the Unlikely TB group, indicating that at least some of these children were likely to have had the disease. Therefore, while oral swab PCR is unlikely to replace testing of more traditional samples for diagnosing TB in children, it may have a useful complementary role.

This study focused on PCR testing of swabs, rather than bacteriological culture of swabs, because the former method yields results far more quickly. A limitation of this study was the use of a manual real-time PCR assay. This strategy was used in this proof-of-principle study because commercial sputum processing methods such as Xpert MTB/RIF are extensively engineered for sputum processing and may be suboptimal for swab analysis. However, manual PCR assays are prone to false-positive results due to laboratory cross-contamination. Two swabs were collected per child, so a total of 88 swabs were tested from 44 children in the Unlikely TB group. Of these, 3 were positive (3.4%). This false positivity rate was similar to that seen with “air swabs” tested by the same manual method in a previous study (3.6%)11. Therefore, at least some of the background signal seen in oral swabs may have resulted from contamination during manual analysis. We did not attempt to use sequencing or other methods to exclude the possibility of false-positive results arising from environmental exposure or non-pathogenic carriage. Overall, automation of oral swab methodology is needed to reduce false-positivity.

In addition, we compared the accuracy of OS PCR with that of liquid culture of a single induced sputum specimen. We, and others, have previously shown that there is substantial incremental yield from testing additional respiratory specimens13,14, and we therefore have likely over-estimated the sensitivity of OS PCR in relation to true disease status. Additional limitations included a small sample size and a focus on a single geographical area.

Given the challenges associated with diagnosing pulmonary TB in children, there is increasing interest in testing of additional sample types that are easily and universally obtainable from children. If oral swab testing is successfully adapted to cartridge-based nucleic acid amplification tests such as Xpert MTB/RIF Ultra, it may hold substantial promise in this regard.

References

1. Nicol, M. P. et al. A Blueprint to Address Research Gaps in the Development of Biomarkers for Pediatric Tuberculosis. *Clinical Infectious Diseases* 61, S164–S172 (2015).
2. Connell, T. G., Zar, H. J. & Nicol, M. P. Advances in the Diagnosis of Pulmonary Tuberculosis in HIV-Infected and HIV-Uninfected Children. *The Journal of Infectious Diseases* 204, S151–S158 (2011).
3. Walters, E. A. et al. Molecular Detection of Mycobacterium tuberculosis from Stools in Young Children by Use of a Novel Centrifugation-Free Processing Method. *J. Clin. Microbiol.* 56 (2018).
4. Marcy, O. et al. Performance of Xpert MTB/RIF and Alternative Specimen Collection Methods for the Diagnosis of Tuberculosis in HIV-Infected Children. *Clinical Infectious Diseases* 62, 1161–1168 (2016).
5. Ios, V., Cordel, H. & Bonnet, M. Alternative sputum collection methods for diagnosis of childhood intrathoracic tuberculosis: a systematic literature review. *Arch Dis Child* (2018).
6. LaCourse, S. M. et al. Stool Xpert MTB/RIF and urine lipoarabinomannan for the diagnosis of tuberculosis in hospitalized HIV-infected children. *AIDS* 32 (2018).
7. McBride, C. M., Wade, C. H. & Kaphingst, K. A. Consumers’ Views of Direct-to-Consumer Genetic Information. *Annu. Rev. Genom. Hum. Genet.* 11, 427–446 (2010).
8. Wilbur, A. K. et al. From the Mouths of Monkeys: Detection of Mycobacterium tuberculosis Complex DNA From Buccal Swabs of Synanthropic Macaques. *Am. J. Primatol.* 74, 676–686 (2012).
9. Engel, G. A. et al. Naturally acquired Mycobacterium tuberculosis complex in laboratory pig-tailed macaques. *Emerg Microbes Infect* 1, e30 (2012).
10. Wood, R. C. et al. Detection of Mycobacterium tuberculosis DNA on the oral mucosa of tuberculosis patients. *Sci. Rep.* 5 (2015).
11. Luabeya, A. K. et al. Noninvasive Detection of Tuberculosis by Oral Swab Analysis. *J. Clin. Microbiol.* 57, e01847–18 (2019).
12. Nicol, M. P. et al. Accuracy of the Xpert MTB/RIF test for the diagnosis of pulmonary tuberculosis in children admitted to hospital in Cape Town, South Africa: a descriptive study. *The Lancet Infectious Diseases* 11, 819–824 (2011).
13. Graham, S. M. et al. Clinical Case Definitions for Classification of Infrathoracic Tuberculosis in Children: An Update. *Clinical Infectious Diseases* 61, S179–S187 (2015).
14. Zar, H. J. et al. Rapid Molecular Diagnosis of Pulmonary Tuberculosis in Children Using Nasopharyngeal Specimens. *Clinical Infectious Diseases* 55, 1088–1095 (2012).

Acknowledgements

We gratefully acknowledge the contributions of the NHLS diagnostic microbiology laboratory at Groote Schuur Hospital, the study, laboratory, and clinical staff at Red Cross Children’s Hospital and the Division of Medical Microbiology, and the children and their care-givers. We are also grateful to Kris Weigel, Divya Naidoo, and Rita Olson for help in analysing samples. This work was supported by grant OPP1140728 from the Bill & Melinda Gates Foundation and by the Regional Prospective Observational Research in Tuberculosis (RePORT TB) Consortium which is co-funded by the Medical Research Council of South Africa and the US Office of AIDS Research of the National Institutes of Health of the USA (DAAD-16-62066-1). Additional funding was received from the Medical Research Council of South Africa for the Tuberculosis Collaborating Centre for Child Health and for the MRC Unit on Child and Adolescent Health and from the NIH (RO1HD058971).

Author Contributions

M.P.N. designed the study, contributed to interpretation of findings and wrote the first draft of the manuscript. R.W. designed experiments, conducted laboratory work, interpreted results and contributed to the manuscript. L.W. supervised data collection, performed data analysis, and contributed to the manuscript. M.P. recruited participants and contributed to the manuscript. C.W. managed the clinical study site and contributed to the manuscript. Y.G. performed diagnostic laboratory testing and contributed to the manuscript. S.M. coordinated
laboratory aspects of the study and contributed to the manuscript. A.O. contributed to laboratory analysis and interpretation of results. L.J.E. helped design experiments and contributed to the manuscript. H.J.X.Z. designed the study, oversaw recruitment of participants, contributed to interpretation of findings, and contributed to the manuscript. G.C. led the study, designed experiments, interpreted results, and contributed to the manuscript.

Additional Information

Competing Interests: The authors declare no competing interests.

Publisher’s note: Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made. The images or other third party material in this article are included in the article’s Creative Commons license, unless indicated otherwise in a credit line to the material. If material is not included in the article’s Creative Commons license and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2019