



ELSEVIER



<http://intl.elsevierhealth.com/journals/ijid>

Mycobacterium tuberculosis and *Mycobacterium africanum* in stools from children attending an immunization clinic in Ibadan, Nigeria

S.I.B. Cadmus^a, A.O. Jenkins^b, J. Godfroid^c, K. Osinusi^d,
I.F. Adewole^e, R.L. Murphy^f, B.O. Taiwo^{f,*}

^a Department of Veterinary Public Health & Preventive Medicine, University of Ibadan, Nigeria

^b Department of Veterinary Tropical Disease, Faculty of Veterinary Sciences, University of Pretoria, Pretoria, South Africa

^c Department of Food Safety and Infection Biology, Norwegian School of Veterinary Science, Tromsø, Norway

^d Department of Paediatrics, University College Hospital, Ibadan, Nigeria

^e Department of Obstetrics and Gynaecology, University College Hospital, Ibadan, Nigeria

^f Division of Infectious Diseases, Northwestern University Feinberg School of Medicine, Chicago, IL, USA

Received 6 October 2008; accepted 30 November 2008

Corresponding Editor: William Cameron, Ottawa, Canada

KEYWORDS

Tuberculosis;
Stool;
Diagnosis;
Pediatric

Summary

Background: Tuberculosis is a major cause of childhood morbidity and mortality in Nigeria. Diagnosis of childhood tuberculosis is a global challenge making early treatment a mirage. In this study we investigated the stools of children for the presence of mycobacteria.

Methods: Stool samples from children aged 3 days to 3 years who presented for postnatal immunization at a large university-based clinic in Nigeria, were subjected to Ziehl–Neelsen staining. Samples with acid-fast bacilli were further processed using mycobacterial culture, spoligotyping, and deletion typing.

Results: One hundred and ninety-two stool samples from different children were collected and processed. Thirty (15.6%) had acid-fast bacilli. Of these, eight had *Mycobacterium tuberculosis* and one had *Mycobacterium africanum*.

Conclusions: Approximately 5% (9/192) of apparently well children had evidence of potentially serious tuberculosis infection. The usefulness of stool specimens for diagnosing pediatric tuberculosis warrants further investigation.

© 2009 International Society for Infectious Diseases. Published by Elsevier Ltd. All rights reserved.

Introduction

More than 2 billion people, approximately one-third of the global population, are infected with *Mycobacterium tuberculosis*, the major causative organism of tuberculosis (TB).¹

* Corresponding author. Tel.: +1 312 6950009; fax: +1 312 6955088.
E-mail address: b-taiwo@northwestern.edu (B.O. Taiwo).

Mycobacterium africanum and *Mycobacterium bovis*, also members of the *Mycobacterium tuberculosis* complex, are much less frequent causes of TB in humans. The incidence and prevalence of pediatric TB varies significantly across the globe, driven largely by the burden of the disease in different countries. About 1 million children under 15 years of age develop TB worldwide annually, representing 11% of all TB cases.² The majority of these cases occur in low-income countries where the prevalence of HIV/AIDS is high.³ Nigeria currently ranks fourth on the list of TB burdened nations globally,¹ and pediatric TB accounts for a substantial proportion of these cases.⁴ Almost 2 million people per year die as a result of TB, mostly in developing countries like Nigeria, but the mortality in children is often underreported. Despite this, TB is one of the ten leading causes of childhood mortality.⁵

Young children and especially newborns are at a high risk when exposed to a contagious source.⁶ A comprehensive review of the natural history of childhood TB showed that primary infection before 2 years of age frequently progressed to active disease within 12 months.⁷ As such, pediatric TB is a sentinel event reflecting recent TB transmission from an infectious contact in the community. The number of children with TB in a community is an indirect parameter for assessing the effectiveness of the local TB control program.³

The diagnosis of pulmonary TB (PTB) in children is challenging.⁸ Children rarely expectorate adequate amounts of sputum, and the limitations of using other specimens or techniques, such as first morning gastric aspirates (considered the best clinical specimens for young children with suspected PTB), nasopharyngeal swabs, sputum induction, and laryngeal swabs, are well known. Accordingly, there is a strong imperative to evaluate the diagnostic utility of clinical specimens that are more readily collectable. Some investigators have suggested that stool microscopy and culture for *M. tuberculosis* may be diagnostic in some children with tuberculosis,^{9–12} but other investigators have described stool evaluation as 'worthless' since non-pathogenic acid-fast bacilli (AFB) may be found in the normal intestinal contents of adults.^{13,14}

Following the identification of AFB in the stools of apparently well children who were being screened for cryptosporidiosis in our immunization clinic, we designed this study to characterize AFB in the stools of children attending the clinic.

Methods

Subjects

Subjects were consecutive children who presented for immunization at the University of Ibadan Health Services Clinic, Ibadan, Nigeria. Stool samples from the children were evaluated using AFB staining. All the AFB-positive stool specimens were evaluated further for the presence of mycobacteria.

Ethical approval

The Institutional Review Committee of the University of Ibadan and the University College Hospital, Ibadan, Nigeria approved the study. Oral informed consent was obtained from the parents of the children.

Clinical specimens

Stool samples were collected from each child into a sterile plastic container and kept in the refrigerator at 4 °C prior to processing using Ziehl–Neelsen (ZN) staining. The ZN stain was carried out as described by Shrestha et al.¹⁵

Processing of samples

From the stool samples positive by ZN staining, 2–3 g was suspended in 5 ml of sterile distilled water, mixed, and left for 15 min to separate, after which 3 ml of the supernatant was processed. Using a sterile centrifuge tube, equal amounts of specimen and activated *N*-acetyl-L-cysteine (NALC)–NaOH (3 ml each) was added. The contents of the tube were mixed until the specimen was liquefied, and then allowed to stand for 15 min. Phosphate buffer was added to the 10 ml mark on the centrifuge tube and mixed, followed by centrifugation for 15 min at 3000 × *g*. The supernatant was decanted; 2 ml of phosphate buffer pH 6.8 was added to resuspend the pellet. The suspension was inoculated onto Lowenstein–Jensen slopes with pyruvate and/or glycerol and incubated at 37 °C for between 8 and 12 weeks. Isolates were harvested for molecular typing analysis by scraping the growth from slopes into 200 µl of sterile distilled water and heating at 80 °C for 1 h.

Spoligotyping

This was carried out as previously described with minor modifications.¹⁶ The direct repeat (DR) region was amplified by PCR with oligonucleotide primers derived from the DR sequence. Twenty-five microliters of the following reaction mixture was used for the PCR: 12.5 µl of HotStarTaq Master Mix (Qiagen; this solution provides a final concentration of 1.5 mM MgCl₂ and 200 µM each deoxynucleoside triphosphate), 2 µl of each primer (20 pmol each), 5 µl of the suspension of heat-killed cells (approx. 10–50 ng), and 3.5 µl of distilled water. The mixture was heated for 15 min at 96 °C and subjected to 30 cycles of 1 min at 96 °C, 1 min at 55 °C, and 30 s at 72 °C. The amplified product was hybridized to a set of 43 immobilized oligonucleotides, each corresponding to one of the unique spacer DNA sequences within the DR locus. After hybridization, the membrane was washed twice for 10 min in 2× SSPE (1× SSPE is 0.18 M NaCl, 10 mM NaH₂ PO₄, and 1 mM EDTA (pH 7.7))–0.5% sodium dodecyl sulfate (SDS) at 60 °C and then incubated in 1:4000-diluted streptavidin-peroxidase conjugate (Boehringer) for 45 to 60 min at 42 °C. The membrane was washed twice for 10 min in 2× SSPE–0.5% SDS at 42 °C and rinsed with 2× SSPE for 5 min at room temperature. Hybridizing DNA was detected by the enhanced chemiluminescence method (Amersham) and by exposure to X-ray film (Hyperfilm ECL; Amersham) as specified by the manufacturer.

Deletion typing

The use of deletion analysis for the typing of *M. tuberculosis* complex strains has been previously described.^{17,18} For this work, the deletion typing method described by Warren et al.¹⁹ was used. In our analysis, we used primers directed against the RD4 and RD9 loci to generate a deletion profile

Table 1 Profile of pediatric patients with a positive culture result for *Mycobacterium* species

ID	Sex	Age	SpolDB4 type	Family in SpolDB4 type	Species
JC1	F	2 months	52	T2	<i>M. tuberculosis</i>
JC2	F	1 week	53	T1	<i>M. tuberculosis</i>
JC3	M	6 weeks			<i>M. tuberculosis</i> (deletion analysis)
JC4	F	5 weeks	61	LAM10_CAM	<i>M. tuberculosis</i>
JC5	M	3 months	774	T1	<i>M. tuberculosis</i>
JC6	F	1 year, 3 months	61	LAM10_CAM	<i>M. tuberculosis</i>
JC7	M	4 months			<i>M. africanum</i> (deletion analysis)
JC8	F	3 months	Not in SpolDB4	New strain (probably evolved from LAM10_CAM)	<i>M. tuberculosis</i>
JC9	F	6 months	358	T1	<i>M. tuberculosis</i>

that would allow speciation of the isolate. The multiplex master mix system from Qiagen was used for the PCRs, with primers previously described by Warren and colleagues.¹⁹ The PCR mixture was a multiplex reaction, with each PCR reaction containing 1 µl of DNA template, 5 µl Q-buffer, 12.5 µl multiplex master mix (Qiagen), and 0.5 µl of each primer (50 pmol/µl). The total volume of the reaction was made up to 25 µl with water. The reaction was allowed to run for 15 min at 95 °C, followed by 45 cycles at 94 °C for 1 min, 62 °C for 1 min, and 72 °C for 1 min. After the last cycle, the samples were incubated at 72 °C for 10 min. Products were visualized by electrophoresis on 3% agarose gels.

The positive controls included a known *M. bovis* isolate (AN5) and a known *M. tuberculosis* isolate provided by the Medical Research Council, Center for Molecular and Cellular Biology, Stellenbosch University, Cape Town, South Africa, whilst the negative control was water. The resulting gel images were analyzed on the basis of their alignment on the gel (i.e., same band size with either of the controls). The RD9 deletion analysis was done to discriminate *M. tuberculosis* from other *Mycobacterium tuberculosis* complex (MTC). Those with a deletion at this region were further investigated with primers targeting the RD4 region and this discriminated *M. bovis* from the other members of the MTC. Primers targeting the RD1^{mic} and RD2^{seal} regions were later used to confirm the presence of *M. africanum* relative to *Mycobacterium pinnipedii* and *Mycobacterium microti* as described by Warren and colleagues.¹⁹

Results

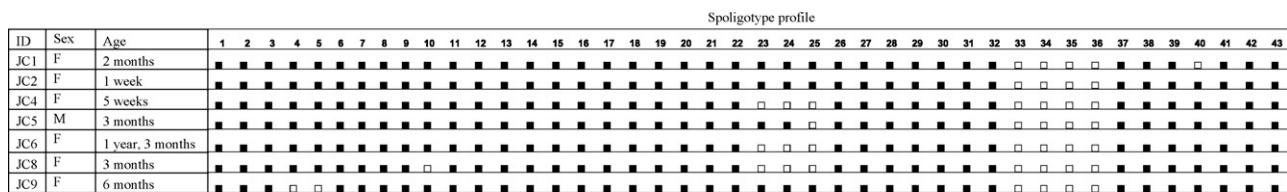
One hundred and ninety-two children were recruited into the study, comprising 95 males and 97 females, aged 3 days to 3 years. Thirty children had AFB present in their stool specimens. Mycobacterial culture of stool samples from the 30

children yielded a growth in nine (30%). Spoligotyping and deletion analysis confirmed these isolates as *M. tuberculosis* complex (eight *M. tuberculosis* and one *M. africanum*; Table 1 and Figure 1). The patients with the positive stool mycobacterial cultures included three males and six females, with ages ranging from 1 week to 15 months (Table 1).

Discussion

The diagnosis of TB in pediatric patients is often based on case definitions that incorporate signs and symptoms of TB, suggestive chest radiograph, positive tuberculin skin test, and contact with an active TB patient. It is less frequently based on laboratory isolation of *M. tuberculosis*. Existing algorithms, however, have serious shortcomings, and the development of reliable and widely applicable algorithms is a high research priority.

In this study, we diagnosed *M. tuberculosis* in 27% (8/30) and *M. africanum* in 3% (1/30) of children who had AFB-positive stool specimens. Other investigators have diagnosed tuberculosis based on isolation of *M. tuberculosis* from stool specimens, but in different patient populations. Mwachari et al.²⁰ cultured *M. tuberculosis* from the stools of 10 (13%) HIV-infected adults with chronic diarrhea in Kenya. Manatsathit et al.²¹ also found *M. tuberculosis* in the stools of eight (18%) adult AIDS patients in Thailand. In South Africa, 8% and 5% of 66 children with suspected PTB had stool specimens that were AFB-positive and *M. tuberculosis* culture-positive, respectively.⁹ In that study, AFB were identified only in the stools of children who had PTB that was confirmed with positive gastric aspirates, but stool testing was less sensitive than gastric aspirates overall. Our study is unique because the study population included apparently well children who were brought to the clinic for routine immunization. None of



Key:
 ■: Spacer present
 □: Spacer absent

Figure 1 Spoligotype profile of pediatric patients with a positive culture result for *Mycobacterium tuberculosis* complex.

the children had diarrhea, but four had cough, although clinical data were not collected prospectively in all children. The yield might be higher in children clinically suspected of having TB.

Some findings of this study provide potentially important epidemiological information. Firstly, *M. tuberculosis* has been implicated in most cases of TB in children; however, we found a case of *M. africanum*. This is, to our knowledge, the first published isolation of *M. africanum* in the stools of a Nigerian child. Cases of TB caused by *M. africanum* have been previously reported in adults from Nigeria^{22,23} and other African countries.^{24,25} In Cameroon, there has been a decline in the prevalence of *M. africanum*,²⁴ however more cases have recently been observed in HIV/AIDS patients in The Gambia²⁵ and Nigeria.²³ Secondly, one of the *M. tuberculosis* strains cultured in this study (isolate from patient JC8) has not been previously reported in the SpolDB4 database, which contains a comprehensive listing of the *M. tuberculosis* strains around the world (Figure 1). This implies the possible circulation of poorly characterized or emerging *M. tuberculosis* strains in Nigeria.

This study should be interpreted in the context of its limitations. Since we did not have records of HIV testing in the mothers or children, we were unable to correlate our findings with the HIV infection status of the subjects. Also, full contact tracing was possible only in some of the children with positive stool tests, seriously constricting our ability to comment on the public health significance of the results. Our data cannot be extrapolated to older children because the oldest subject in this study was 3 years old. Finally, we do not have complete data on the subsequent clinical course of the patients. Despite these limitations, our findings add to the evidence that directed stool studies may be useful in pediatric TB diagnosis.

In conclusion, we have described the isolation of *M. tuberculosis* from the stools of a significant proportion of apparently well children attending the University of Ibadan Health Services Clinic. We have also described the first isolation of *M. africanum* from the stools of a Nigerian child. The problematic nature of diagnosing TB disease in this age group justifies further investigation of the diagnostic potential of stool specimens and other readily obtainable specimens, perhaps using more sensitive techniques. The limitations of such testing and the population for which it would be applicable are also fertile areas for clinical and laboratory studies.

Acknowledgments

The project described was supported by grant No. D43TW007995 from the Fogarty International Center. The content is solely the responsibility of the authors and does not necessarily represent the official views of the Fogarty International Center or the National Institutes of Health.

Conflict of interest: No conflict of interest to declare.

References

1. World Health Organization. Global tuberculosis control: surveillance, planning, financing. WHO Report 2007. WHO/HTM/TB/2007.376. Geneva: World Health Organization; 2007. Available at: http://www.who.int/tb/publications/global_report/2007/en/index.html (accessed January 2009).
2. Corbett EL, Watt CJ, Walker N, Maher D, Williams BG, Raviglione MC, et al. The growing burden of tuberculosis: global trends and interactions with the HIV epidemic. *Arch Intern Med* 2003;163:1009–21.
3. Morcillo N. Tuberculosis in children. In: Palomino JC, Leao SC, Ritacco V, editors. Tuberculosis 2007—from basic science to patient care. Bernard Sebastian Kamps and Patricia Bourcillier Fund; 2007, p. 525–58. Available at: <http://www.tuberculosis-textbook.com> (accessed January 2009).
4. Osinusi K. Clinical and epidemiological features of childhood tuberculosis in Ibadan, Nigerian. *J Paediatr* 1998;25:15–9.
5. Rheka B, Swaminathan S. Childhood tuberculosis—global epidemiology and the impact of HIV. *Pediatric Resp Rev* 2007;8:99–106.
6. Dye C, Scheele S, Dolin P, Pathania V, Raviglione MC, for the WHO Global Surveillance, Monitoring Project. Global burden of tuberculosis: estimated incidence, prevalence and mortality by country. *JAMA* 1999;282:677–86.
7. Marais BJ, Gie RP, Schaaf HS, Hesselting AC, Obihara CC, Starke JJ, et al. The natural history of childhood intra-thoracic tuberculosis: a critical review of literature from the pre-chemotherapy era. *Int J Tuberc Lung Dis* 2004;8:392–402.
8. Starke JR, Taylor-Watts KT. Tuberculosis in the pediatric populations of Houston, Texas. *Pediatrics* 1989;84:28–35.
9. Donald PR, Schaaf HS, Gie RP, Beyers N. Stool microscopy and culture to assist the diagnosis of pulmonary tuberculosis in childhood. *J Trop Pediatr* 1996;42:311–2.
10. Allen BW. Isolation of *Mycobacterium tuberculosis* from faeces. *Med Lab Sci* 1989;46:101–6.
11. Allen BW. Comparison of three methods for decontamination of faeces for isolation of *Mycobacterium tuberculosis*. *Tubercle* 1991;72:214–7.
12. Oberhelman RA, Soto-Castellares G, Caviedes L, Castillo ME, Kissinger P, Moore DA, et al. Improved recovery of *Mycobacterium tuberculosis* from children using the microscopic observation drug susceptibility method. *Pediatrics* 2006;118:e100–6.
13. Collins CH, Grange JM, Yates MD. *Organization and practice in tuberculosis bacteriology*. London: Butterworth-Heinemann; 1985. p. 36–43.
14. Portaels F. The importance and evaluation of mycobacterial diseases as assessed by a mycobacteriological laboratory. *Bull Int Union Tuberc Lung Dis* 1988;63:13–6.
15. Shrestha D, Bhattachaya SK, Rajendra K. Evaluation of different staining techniques (Ziehl–Neelsen stain, Kinyoun stain, modified cold stain, fluorochrome stain) for the diagnosis of pulmonary tuberculosis. *J Nepal Health Res Council* 2005;3:8–16.
16. Kamerbeek J, Schouls L, Kolk A, van Agterveld M, van Soolingen D, Kuijper S, et al. Simultaneous detection and strain differentiation of *Mycobacterium tuberculosis* for diagnosis and epidemiology. *J Clin Microbiol* 1997;35:907–14.
17. Brosch R, Gordon SV, Marmiesse M, Brodin P, Buchrieser C, Eiglmeier K, et al. A new evolutionary scenario for the *Mycobacterium tuberculosis* complex. *Proc Natl Acad Sci U S A* 2004;99:3684–9.
18. Parsons LM, Brosch R, Cole ST, Somoskovi A, Loder A, Bretzel G, et al. Rapid and simple approach for identification of *Mycobacterium tuberculosis* complex isolates by PCR-based genomic deletion analysis. *J Clin Microbiol* 2002;40:2339–45.
19. Warren RM, Gey van Pittius NC, Barnard M, Hesselting A, Engelke E, de Kock M, et al. Differentiation of *Mycobacterium tuberculosis* complex by PCR amplification of genomic regions of difference. *Int J Tuberc Lung Dis* 2006;10:818–22.
20. Mwachari C, Batchelor BI, Paul J, Waiyaki PG, Gilks CF. Chronic diarrhoea among HIV-infected adult patients in Nairobi, Kenya. *J Infect* 1998;37:48–53.
21. Manatsathit S, Tansupasawasdikul S, Wanachiwanawin D, Setawarin S, Suwanagool P, Prakasvejakit S, et al. Causes of chronic

- diarrhea in patients with AIDS in Thailand: a prospective clinical and microbiological study. *J Gastroenterol* 1996;**31**:533–7.
22. Cadmus SI, Palmer S, Okker M, Dale J, Gover K, Smith N, et al. Molecular analysis of human and bovine tubercle bacilli from a local setting in Nigeria. *J Clin Microbiol* 2006;**44**:29–34.
 23. Cadmus SI. Molecular characterisation of human and bovine tubercle bacilli in Ibadan. PhD Thesis. University of Ibadan, Ibadan, Nigeria, 2007.
 24. Niobe-Eyangoh SN, Kauban C, Sorlin P, Thonnon J, Vincent V, Gutierrez MC, et al. Molecular characteristics of strains of the Cameroon family, the major group of *Mycobacterium tuberculosis* in a country with a high prevalence of tuberculosis. *J Clin Microbiol* 2004;**42**:5029–35.
 25. de Jong BC, Hill PC, Brookes RH, Otu JK, Peterson KL, Small PM, et al. *Mycobacterium africanum*: a new opportunistic pathogen in HIV. *AIDS* 2005;**19**:1714–5.