Treatment Outcomes in Global Systematic Review and Patient Meta-Analysis of Children with Extensively Drug-Resistant Tuberculosis

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Learning Objectives

Upon completion of this activity, participants will be able to:

• Evaluate the presentation of children from 11 countries who were managed between 1999 and 2013 for bacteriologically confirmed extensively drug-resistant tuberculosis (XDR TB), based on a global systematic review and individual patient meta-analysis

• Assess the treatment of children from 11 countries who were managed between 1999 and 2013 for bacteriologically confirmed XDR TB, based on a global systematic review and individual patient meta-analysis

• Determine the outcome of children from 11 countries who were managed between 1999 and 2013 for bacteriologically confirmed XDR TB, based on a global systematic review and individual patient meta-analysis.

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Extensively drug-resistant tuberculosis (XDR TB) has extremely poor treatment outcomes in adults. Limited data are available for children. We report on clinical manifestations, treatment, and outcomes for 37 children (<15 years of age) with bacteriologically confirmed XDR TB in 11 countries. These patients were managed during 1999-2013. For the 37 children, median age was 11 years, 32 (87%) had pulmonary TB, and 29 had a recorded HIV status; 7 (24%) were infected with HIV. Median treatment duration was 7.0 months for the intensive phase and 12.2 months for the continuation phase. Thirty (81%) children had favorable treatment outcomes. Four (11%) died, 1 (3%) failed treatment, and 2 (5%) did not complete treatment. We found a high proportion of favorable treatment outcomes among children, with mortality rates markedly lower than for adults. Regimens and duration of treatment varied considerably. Evaluation of new regimens in children is required.

Extensively drug-resistant tuberculosis (XDR TB) is a public health emergency that threatens global TB control. Multidrug-resistant TB (MDR TB) is caused by Mycobacterium tuberculosis that shows resistance to isoniazid and rifampin, and XDR TB includes additional resistance to any fluoroquinolone and a second-line injectable drug (1). In 2016, a total of 8,014 cases of XDR TB were reported to the World Health Organization (WHO) by 72 countries (2). Treatment success rates for XDR TB remain poor; only 30% of patients show cure or treatment completion, and costs for care far exceed those for drug-susceptible TB (2). There is an increasing awareness that children are also affected by MDR TB and XDR TB. Modeling studies estimated that as many as 2 million children currently have MDR TB globally, and MDR TB develops in an estimated 30,000 children <15 years of age each year (3,4). Estimates suggest that of children with MDR TB, 4.7% have XDR TB (4).

We recently completed a systematic review and individual patient data (IPD) meta-analysis commissioned by WHO that described clinical presentation, treatment, and outcomes for children treated for MDR TB (5). We reported data for 975 children with MDR TB from 18 countries; 731 (75%) had bacteriologically confirmed MDR TB, and 244 (25%) had clinically diagnosed MDR TB (6). Overall, 764 (78%) of 975 children had a successful treatment outcome (6), as defined by WHO guidelines (7). This meta-analysis provided information on pediatric aspects for the revised 2016 WHO drug-resistant TB treatment guidelines, and specific recommendations were subsequently made for treatment of MDR TB in children (5). Children with confirmed XDR TB were excluded because they were a distinct subgroup and insufficient evidence was available to make treatment recommendations for children with XDR TB at that time (5).

The management of XDR TB in children is challenging because of the limited availability of new drugs and appropriate treatment regimens. XDR TB treatment regimens for children have historically been individualized on the basis of mycobacterial drug-susceptibility testing (DST) of the organism of the child or the putative source case (5). There are limited data on the optimal combination of medications and the duration of treatment for XDR TB and major research gaps remain (5). Therefore, we aimed to describe clinical manifestations, routine treatment, and outcomes for children with confirmed XDR TB in the era preceding access to novel anti-TB drugs for children.

Methods

Data Collection

As part of a systematic review and IPD meta-analysis, we collected data from global collaborators on children <15 years of age treated for MDR TB as part of a defined treatment cohort (6). We identified published and unpublished data from retrospective and prospective studies by using a broad search strategy. Eligible studies were identified, and individual level patient data were requested from each author by using a standardized data collection tool. We requested demographics, clinical details, and outcomes on the basis of specified definitions. Additional interpretation of data was conducted by the study team, and the primary authors were contacted to resolve queries. Data were obtained for 1,012 children treated during 1999-2013 (6). Detailed methods and outcomes of IPD meta-analysis for 975 children with MDR TB and pre-XDR TB (MDR TB with additional resistance to either a fluoroquinolone...
or an injectable drug, but not both) was reported (6). We report on children identified through systematic review of drug-resistant TB in children who had bacteriologically confirmed XDR TB. Therefore, this analysis only included children who were investigated for TB, had presence of *M. tuberculosis* confirmed bacteriologically, had isolates tested for resistance to second-line anti-TB drugs, and received a treatment outcome during the episode of TB.

**Definitions**

We defined TB as pulmonary TB when disease was localized to the lungs or intrathoracic lymph nodes and as extrapulmonary TB when disease was found at site distant from the lungs, including pleural effusions and miliary TB. We classified pulmonary TB as severe or nonsevere by using adapted criteria of Wiseman et al. (8) on the basis of a review of reported chest radiographs by 2 independent reviewers; a third reviewer arbitrated discordance. Previous TB treatment history and type of treatment previously received (for drug susceptible TB or drug-resistant TB), was documented when known. We defined malnutrition as being underweight for age (weight-for-age z-score < –2) or per the report of the treating clinician.

Because the review spanned many years and sites, DST methods varied by region and period. Data collected for treatment of XDR TB varied, and sites inconsistently submitted data on individual drugs used and drug dose or duration of treatment. We report intensive phase and continuation phase as submitted, for which investigators defined these 2 stages according to their clinical practice. The intensive phase typically refers to initial months of treatment, which include more drugs and the use of an injectable drug. The continuation phase refers to a second phase of treatment generally with a reduction in the number of drugs.

We defined TB treatment outcomes by using standard 2014 WHO MDR TB outcome definitions as classified by treating clinicians: cure (treatment completed as recommended by the national policy without evidence of failure and ≥3 consecutive cultures taken at least 30 days apart were negative after the intensive phase of treatment); treatment complete (as for cure but without records of negative cultures); treatment failed (treatment stopped or requiring change of 2 drugs because of persistent positive cultures at end of the intensive phase or reversion to positive cultures in the continuation phase, or evidence of additional acquired resistance or adverse drug reactions); death (for any reason while receiving treatment); or loss to follow-up (treatment interruption for 2 consecutive months) (9). We defined favorable (cure and completed treatment) and unfavorable (treatment failure, death, or loss to follow-up) outcomes. Adverse events were inconsistently included in primary data and are not reported here.

**Statistical Analysis**

We completed descriptive statistics for demographic and clinical variables. Missing data were noted, and each analysis reflects the sample size used. Categorical variables are presented as frequencies and percentages, and continuous variables (duration of intensive or continuation phase) are presented as median and interquartile ranges (IQRs). Age of children was categorized as <2 years, 2–4 years, 5–9 years, or 10–14 years. We used logistic regression with a preset 95% level of significance and calculated odds ratios (ORs) and 95% CIs to estimate and predict unfavorable outcomes. Children with unknown outcomes or loss to follow-up were excluded from regression analyses. We analyzed by using SAS software version 9.4 (https://www.sas.com).

**Ethical Approval**

The study protocol was approved by the Health Research Ethics Committee of the Faculty of Medicine and Health Sciences, Stellenbosch University (reference no. X14/09/020). The oversight committee at the location institution of each contributor approved collection of data and submission to the collaborative systematic review.

**Results**

Of 1,012 children included in the larger systematic review of children with MDR TB, 37 children from 11 countries had bacteriologically confirmed XDR TB (Table 1). We also compiled demographic and clinical characteristics at baseline (Table 2). Median age was 11 years (IQR 6.0–13.1 years). Thirty-two (87%) children had pulmonary TB only. Among children with pulmonary TB and chest radiographic findings, 20 (65%) of 31 had markers of severity, including disseminated/extensive disease, airway compression with lobar collapse, miliary opacification, or cavitation. HIV status was documented for 29 (78%) children; of these, 7 (24%) were infected with HIV.

We also obtained previous TB treatment history for children with XDR TB (Figure 1). Among 33 children who had documented knowledge of previous treatment, 17 (52%) had been previously treated for TB. Of children previously treated, only 10 had known TB treatment outcomes; 7 (70%) had a history of treatment failure.

Data for microbiological investigations were limited. All 37 children had culture-confirmed XDR TB, but only 30 had smear results before treatment, and 10 (33%) were smear positive. Follow-up culture results were infrequently available. The DST pattern for this cohort (Figure 2) showed that all children had isolates with proven resistance to rifampin and isoniazid. Isolates from some children were tested against multiple injectable drugs and fluoroquinolones (some of which were susceptible), but all children in the cohort had isolates that were resistant to ≥1 second-line injectable drug.
(kanamycin, amikacin, or capreomycin), and \( \geq 1 \) fluoroquinolone (moxifloxacin, levofloxacin, ofloxacin, or ciprofloxacin). DST for additional drugs was performed only for a limited number of children.

We also evaluated drugs used to treat these children (Figure 3). No children received bedaquiline or delamanid because both drugs became available only after the study period. The most commonly used drugs were an injectable (n = 27), a fluoroquinolone (n = 26), cycloserine/terizidone (n = 27), ethionamide/prothionamide (n = 24), and para-aminosalicylic acid (n = 25). Capreomycin was the most commonly used injectable drug (Figure 3); for some children, >1 injectable was used sequentially. Of 26 children who received a fluoroquinolone, 13 were given moxifloxacin alone, and 4 were switched between moxifloxacin and other fluoroquinolones. Limited information on timing and reason for changing drugs was available. Duration of use of individual drugs was recorded for a limited number of children, and estimates of individual treatment duration per drug could not be made. Median duration of the intensive phase for 26 children who completed treatment was 7 months (IQR 6.0–8.2 months), and median duration for 23 children who completed the continuation phase was 12.2 months (IQR 10.0–16.2 months).

Overall, 30 (81%) children had favorable treatment outcomes (Figure 4). Four (11%) children died during XDR TB treatment; 2 were HIV-infected and receiving antiretroviral therapy. One child (3%) had documentation of treatment failure (HIV infected when receiving antiretroviral therapy) and 2 (5%) were lost to follow-up during treatment (HIV uninfected). We obtained detailed demographic and clinical variables for the 7 children with unfavorable outcomes, including the DST pattern of the isolate and drugs used during treatment (Table 3).

We detected no associations between recorded variables and unfavorable treatment outcome by using univariate analyses. Although HIV-infected children had a 14.3 times (95% CI 1.2–174.8 times) greater odds of an unfavorable outcome than HIV-uninfected children, this logistic regression model used infected, uninfected, and unknown as 3 possible categories of HIV, and differences in the overall model were not significant (p = 0.098). Multivariable analysis was not completed because of lack of significance in all univariate analyses.

### Discussion

Data for treatment of XDR TB in children are limited. Our group of 37 confirmed cases over 15 years represents a small sample yet was larger than those in previous reports, highlighting the serious underdiagnosis and underreporting of XDR TB in children in the published literature. We included data for clinical manifestations, treatment, and outcomes for these children. Overall, we describe encouraging rates of successful treatment outcomes in this small cohort of children despite the limited drug options available to treat children in most instances and the notable lack of access to novel drugs, such as bedaquiline and delamanid, at the time of treatment. All children had confirmed XDR TB; more work is needed to include children given a diagnosis of probable XDR TB in the absence of bacteriological confirmation. Most children were >5 years of age and had pulmonary TB and severe TB. Half of the children had been previously given treatment for TB, and of those children, half had been previously treated with a drug-susceptible TB treatment regimen.

Of 975 children described in the MDR TB IPD, only 37 had confirmed XDR TB, which might reflect the limited access to second-line DST. The older median age (11 years) observed in this group than that for a pediatric MDR TB IPD review (median age 7.1 years) (6) might indicate that younger children might not have been as likely to have been investigated and given a diagnosis of XDR TB. Because it has been estimated that 80% of childhood TB deaths occur in children <5 years of age...
DST patterns in this cohort were emerging infectious diseases because of their typical paucibacillary disease and presentation (given a diagnosis based only on clinical findings) because older children are more likely to have adult-type TB with higher bacillary burden, bacteriological confirmation is more likely (22).

It is essential that empiric treatment for XDR TB in children be based on DST of the organism of the putative TB source case-patient, but the need for adequate sampling for TB bacteriology including gastric aspirates, or other samples, cannot be overstated (23–25). If one considers the risk for rapid disease progression (26), treatment initiation in children should not be delayed. Although DST for the XDR TB isolate might be used for development of a targeted treatment regimen, a good contact history with DST of the organism of the source case-patient might also be used where confirmation is not possible. In our study, 28 (76%) children had recorded information on source cases. Of these children, 20 (70%) had an identified source case during their episode of TB, but we did not have sufficient information to define the DST pattern of the source case. This finding highlights the need for contact investigation of source cases. Early screenings of households after diagnosis of TB is an opportunity to identify at-risk contacts. Symptom screening and sampling of child contacts could facilitate earlier management.

*M. tuberculosis* DST patterns in this cohort were limited because the review spanned 15 years starting in 1999. Determination of fluoroquinolone resistance varied and, for the definition we applied in this study, ciprofloxacin resistance was included. Data for additional testing of drugs was limited and did not enable us to make inferences about the DST pattern and the drug or regimen chosen or any changes in DST pattern during the course of treatment.

Although we were not able to evaluate the role of specific drugs because of the small number of children and the variability in treatment regimens, we found patterns of commonly used drugs. Most (n = 27) children received an injectable drug, and although more exact data were limited, the median duration used was 7 months. Isolates from all children had confirmed resistance to an injectable drug; thus, it is a concern that an injectable isolate might be used for development of a tar getsed therapy (27,28). It is possible that clinicians expected incomplete cross-resistance within the injectable drug group, a rationale supported by the predominance of capreomycin use (29). However, cross-resistance within this group is relatively common (30). It is also possible that children might have been given these agents before second-line DST results became available. *M. tuberculosis* strains resistant to earlier-generation fluoroquinolones might retain susceptibility to a later-generation fluoroquinolone (31), which might have supported fluoroquinolone use.
SYNOPSIS

Despite documented resistance, most (n = 26) children received a fluoroquinolone, and this drug was a later-generation treatment for >50% of case-patients. This finding might reflect the anticipation of clinicians for some activity of the drug, despite DST results and limited other treatment options.

Most children in our cohort were given treatment before availability of new drugs. Linezolid was only used for 3 children and clofazimine for 6 children, whereas bedaquiline and delamanid were not used. Paraaminosalicylic acid, ethionamide or prothionamide, and cycloserine or terizidone were used frequently, highlighting the limited drug options available and the necessity to use drugs with major toxicity and relatively poor efficacy. With new drugs, alternatives for children with drug-resistant TB are being explored. The WHO endorsement of delamanid for older children (>6 years of age) with MDR TB (32) and early reports on the use of bedaquiline in older children (≥10 years of age) (33) might be initial steps in finding the optimum treatment regimen for XDR TB in children.

Children with confirmed XDR TB had similar successful outcomes (81%) as children given treatment for clinically diagnosed and bacteriologically confirmed MDR TB (78%) over the same study period (6). Although loss to follow-up was lower (5.4% for XDR TB and 11.2% for MDR TB) (6), our cohort of children with XDR TB was smaller, and children with XDR TB might have had more intensive follow-up than children with MDR TB. WHO combined program data for 6,777 XDR TB case-patients (adults and children) reported from 52 countries during 2014 found treatment success in only 30%, with 28% deaths, 21% treatment failures, and 20% lost to follow-up (2). Overall mortality rates exceeded 40% in India and South Africa (2). High mortality rates for XDR TB in adults has been well documented.

Figure 1. Flow chart for 37 children with confirmed XDR TB and details of TB treatment history, type of TB treatment, and treatment outcome. DR, drug-resistant; DS, drug-susceptible, MDR, multidrug-resistant; TB, tuberculosis; XDR TB, extensively drug-resistant TB.

Figure 2. Mycobacterial drug susceptibility test pattern for children treated for extensively drug-resistant tuberculosis. All children had organisms that were resistant to rifampin and isoniazid. *Includes moxifloxacin, levofloxacin, ofloxacin, or ciprofloxacin. †Includes second-line injectable drugs kanamycin, amikacin, or capreomycin. AMK, amikacin; CAP, capreomycin; CIP, ciprofloxacin; CYS, cycloserine; EMB, ethambutol; ETH, ethionamide; KAN, kanamycin; LVX, levofloxacin; MXF, moxifloxacin; OFX, ofloxacin; PAS, para-aminosalicylic acid; PTH, prothionamide; PZA, pyrazinamide; STR, streptomycin; TRZ, terizidone.
In our cohort of children with XDR TB, 4 (10.8%) died during treatment. Considering the natural history of TB, we acknowledge that spontaneous cure in children occurs and might also contribute to the good outcomes seen. We note that none of the children included in our cohort had TB meningitis, which is well known to have poor outcomes (38), and this finding might contribute to the good outcomes we report. This lower mortality rate might be caused by this group representing a survival bias, only including children who were alive to make the diagnosis and initiate treatment. In addition, children tend to have lower organism load (paucibacillary), have less concomitant pathology, and be treated with more tailored individualized regimens compared with adults, which might improve survival. Diagnosis and treatment of XDR TB was historically limited to tertiary centers or specialized centers of excellence for TB care, providing more resources for care and improving diagnosis and treatment outcomes. Most (86%) children were admitted to a hospital and would have likely had good adherence support. We acknowledge that this cohort consisted of a small number of selected children and are therefore cautious about generalizability.

A further limitation of our study is poor reporting of adverse events, which could not be analyzed. Poor adverse event reporting has been identified by WHO as a problem (2). Systematic monitoring of TB drug safety for children is crucial, especially if one considers introduction of new drugs. Given the modest sample size and the limitations regarding the dose and exact duration of drugs, we could not analyze the effect of individual drugs, regimen combinations, or duration of treatment on final treatment outcomes. Newer drugs require evaluation because WHO has recommended use of delamanid for children and adolescents (32).
Access to newer TB drugs and effective shorter MDR TB regimens is improving. Systematic reviews of MDR TB or XDR TB have identified the benefits of linezolid, a larger number of effective drugs, the number of drugs used in each phase of treatment, and the duration of treatment (39–42). Although there might be some indication for use of more drugs during the intensive phase and continuation phase, with longer duration of treatment for XDR TB, this evidence is still limited (41). Studies to date have mainly included adults (39–42). Our findings highlight the need for more studies evaluating new drugs and treatment regimens in children with XDR TB.

In conclusion, we report treatment of XDR TB for children spanning 15 years. The limited number of children identified highlights a major gap in diagnosing and reporting XDR TB in children. The high proportion of favorable treatment outcomes and considerably lower mortality rates compared with those for adults is encouraging. We found considerable variability of regimens used and duration of treatment in children, but this review preceded availability and use of linezolid, clofazimine, bedaquiline, and delamanid. More collaborative, multicenter prospective cohorts are needed to collect better and more extensive data for children with drug-resistant TB. Evaluation of shorter effective and safe regimens for children with XDR TB is urgently needed.

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etymologia

Streptomycin [strep”to-mi’sin]

Ronnie Henry

In the late 1930s, Selman Waksman, a soil microbiologist working at the New Jersey Agricultural Station of Rutgers University, began a large-scale program to screen soil bacteria for antimicrobial activity. By 1943, Albert Schatz, a PhD student working in Waksman’s laboratory, had isolated streptomycin from Streptomyces griseus (from the Greek strepto- (“twisted”) + mykēs (“fungus”) and the Latin griseus, “gray”).

In 1944, William H. Feldman and H. Corwin Hinshaw at the Mayo Clinic showed its efficacy against Mycobacterium tuberculosis. Waksman was awarded a Nobel Prize in 1952 for his discovery of streptomycin, although much of the credit for the discovery has since been ascribed to Schatz. Schatz later successfully sued to be legally recognized as a co-discoverer of streptomycin.

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