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RESEARCH ARTICLE

Identifying patients with multidrug-resistant tuberculosis who may benefit from shorter durations of treatment

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Abstract

Objective

Studying treatment duration for rifampicin-resistant and multidrug-resistant tuberculosis (MDR/RR-TB) using observational data is methodologically challenging. We aim to present a hypothesis generating approach to identify factors associated with shorter duration of treatment.

Study design and setting

We conducted an individual patient data meta-analysis among MDR/RR-TB patients restricted to only those with successful treatment outcomes. Using multivariable linear regression, we estimated associations and their 95% confidence intervals (CI) between the outcome of individual deviation in treatment duration (in months) from the mean duration of their treatment site and patient characteristics, drug resistance, and treatments used.

repository. Access to this is governed by an oversight committee and applications must be directed to that committee. There are legal restrictions in terms of signed data sharing agreements with all data contributors. Inquiries can be made here: https://www.ucl.ac.uk/global-health/ research/tb-ipd-platform.

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Competing interests: The authors have declared that no competing interests exist.

Results

Overall, 6702 patients with successful treatment outcomes from 84 treatment sites were included. We found that factors commonly associated with poor treatment outcomes were also associated with longer treatment durations, relative to the site mean duration. Use of bedaquiline was associated with a 0.51 (95% CI: 0.15, 0.87) month decrease in duration of treatment, which was consistent across subgroups, while MDR/RR-TB with fluoroquinolone resistance was associated with 0.78 (95% CI: 0.36, 1.21) months increase.

Conclusion

We describe a method to assess associations between clinical factors and treatment duration in observational studies of MDR/RR-TB patients, that may help identify patients who can benefit from shorter treatment.

Introduction

Multidrug-resistant tuberculosis (MDR-TB), defined as tuberculosis with resistance to both rifampicin and isoniazid, is a major global health burden [1]. Although treatment success has increased over time to 60–70% [1, 2], the estimated number of MDR-TB cases has increased from previous years to 450,000 in 2021 [1]. Current recommended treatment from the World Health Organization (WHO) for extensive or severe MDR-TB is as long as 18–20 months [3] and entails a high patient burden. There is no doubt that shorter regimens are attractive for patients, health systems, and providers, as they reduce the burden of treatment [4–7]. In the past 10 years, several studies [8–12] have investigated shorter regimens for treatment of MDR-TB in randomized controlled trials (RCTs), but these may not reflect treatment in programmatic settings.

Assessing the effect of MDR-TB treatment duration in non-randomized studies has several potential limitations. Individuals' treatment durations are determined by the outcomes of loss to follow-up, failure, and death. For those remaining on treatment, the regimens and duration are highly individualized and vary by provider and patient presentation, which entail method-ological challenges. Despite these challenges, investigators have used individual duration as an outcome [13], but inferences were limited and the evidence is considered by the WHO to be of very low quality [14].

Based on previous analyses using individual patient data (IPD) [13, 15] treatment duration varies widely between treatment sites and each site typically has a 'usual' duration of treatment targeted for patients, which may be based on local guidelines, experience, patient population, and availability of anti-tuberculosis drugs. However, there is substantial individual variation around that usual duration at each site. We hypothesize that analyzing individual differences from the site-specific mean treatment duration, among patients with successful treatment outcomes, may help address these methodologic challenges.

Our aim was to describe associations with site-specific average treatment durations and to use deviations from these site-specific average treatment durations to identify clinical and treatment factors associated with shorter duration of treatment among individual rifampicin-resistant (RR-TB) and MDR-TB patients with successful treatment outcomes.

Methods

We conducted this study using a dataset of the 2019 IPD in MDR/RR-TB described in detail previously [15]. This study began in 2016 when the dataset was initially assembled. Briefly, the dataset included data from studies conducted between January 1, 2009, and April 15, 2016 that were identified in a systematic review [16]. In addition, the IPD were updated with data contributed by authors of a 2010 IPD meta-analysis [17] and data from two public calls by the WHO in 2018 [18] and 2019 [19] (Supplemental Figure S1 in S1 File presents a timeline for important changes in WHO treatment guidelines for MDR-TB). For comprehensive details on search strategy, study eligibility, and quality assessment see Supplement 1 in S1 File. Studies exclusively in children were excluded.

The 2019 IPD in MDR/RR-TB contains records from 55 studies and 13,272 patients who initiated treatment between 1993 and 2019 in 38 countries and regions. The characteristics of studies included in the IPD have been described previously [20, 21] and the quality and completeness of all studies in the IPD are described in Supplement 2 (S1 File).

Study population

We included studies reporting individual treatment duration and excluded studies which did not provide information on duration, or only provided planned durations. From the included studies, we included only patients that had successful (cured or completed) treatment outcomes, as defined elsewhere [22, 23] and who had their individual treatment duration recorded. We verified outcomes provided by study investigators in their original study, and harmonized these to WHO 2013 definitions [23], as detailed elsewhere (see supplement to Ahmad et al. [15]). In those with death, failure, or loss to follow-up, their treatment duration is determined by their outcome, which may bias associations between characteristics and treatment duration, and were thus excluded. Any patients for which their individual treatment duration was missing were excluded from our primary analyses.

Outcomes

We assessed two outcomes among patients with successful treatment outcomes: i) the mean treatment duration at each treatment site, which was used in an ecological level analysis to explore potential associations with site-level factors; and ii) the difference between each individual's treatment duration and the mean treatment duration of all patients with treatment success at their site. The latter is our primary outcome in this analysis, which is the individual deviation from the site-specific mean treatment duration; this is referred to as *deviation in treatment duration* throughout the text and interpreted in terms of shorter (negative value) or longer (positive value) duration of treatment in months.

Statistical analysis

Ecological analysis of mean treatment duration of site. We first conducted an ecological analysis of the site-specific mean treatment duration in patients with successful outcomes where the unit of analysis was the treatment site, rather than the individual patient. Using available (non-imputed) data, we computed site-level proportions of categorical variables and means of continuous variables and described all using mean and standard deviation (SD), median and interquartile range [IQR], and range (minimum to maximum). We then performed univariable and multivariable linear regression in imputed data (described below) to examine associations between site-level characteristics and the mean treatment duration of the site (see Supplement 3 in S1 File for details).

Analysis of individual deviation from mean treatment duration of site. In our primary analysis, our approach was to construct an exploratory, hypothesis generating, multivariable model to identify factors conditionally associated with a change in deviation in treatment duration, while controlling for all variables selected into the model.

For clinical characteristics, drug susceptibility testing results, and treatments used we described categorical variables as n (%) while continuous variables were described using mean and standard deviation (SD) or median and interquartile range (IQR) using the available data (for detail on all variable specifications see Supplement 4 in <u>S1 File</u>). We also presented the regression coefficients (in months) and their 95% CI for age- and sex-adjusted univariable associations between deviation in treatment duration and each variable listed previously.

All regression analyses were conducted using data imputed with multivariate imputation by chained equations (MICE) with the assumption that data were missing at random (see Supplement 5 in <u>S1 File</u> for detail). The deviation in treatment duration was imputed for those with either only planned or missing deviation in treatment duration for our sensitivity analyses, along with the other variables, however we only included subjects with non-missing duration in our primary analysis. Twenty data sets were generated with 25 Gibb's sampling iterations [24].

To construct our exploratory model, we included variables known to be associated with treatment success in the published literature [13, 15, 20, 21]. Additionally, we ran adaptive Lasso regression [25], using each imputed data set, on the previously listed characteristics to identify other potentially important predictors of treatment duration that were not a priori identified. Pearson coefficients were used to assess correlation between variables to be included. When highly correlated variables were present, we chose the more clinically relevant variable. We then used multivariable linear mixed-effects models with a random intercept for study to estimate regression coefficients and 95% confidence intervals (CI) for each selected covariate, controlling for the others.

In subgroup analyses, we assessed the final model stratified by subpopulations of patients: i) with MDR/RR-TB plus resistance to both fluoroquinolones (FQ) and second-line injectables (SLI) and all others with MDR/RR-TB (including resistance to FQ or SLI but not both); ii) with or without extensive disease (defined as yes if acid-fast bacilli (AFB) smear positive at baseline, and if AFB smear status was missing then the presence of radiographic findings of cavitation or bilateral disease); and iii) with or without previous tuberculosis treatment. We also did additional exploratory analyses in subgroups of those with: i) extensive disease with only MDR/ RR-TB and those without extensive disease with MDR/RR-TB plus any additional resistance; and ii) those with past tuberculosis treatment with MDR/RR-TB only and those without any past treatment with MDR/RR-TB plus any additional resistance. Additionally, we explored the possible effect of selection bias on our population by analyzing our final model adjusted with inverse probability of selection weights for inclusion into the study population (see Supplement 6 in S1 File for detail). We also performed an analysis that included subjects with missing treatment durations whose durations were imputed in the MICE procedure. Finally, we explored the impact that unmeasured confounding may have on the largest associations estimated from our primary analysis by calculating E-values as described by VanderWeele et al. [26] (see Supplement 7 in <u>S1 File</u>). All analyses were conducted using R version 4.1.2. [27]

This study used individual patient data provided by the investigators of the original studies, who obtained informed consent from all participants as appropriate for their original study designs. All data received were anonymized. This analysis received ethical approval from the McGill University Health Centre Research Ethics Board. Ethics approval was also obtained at participating sites, if considered necessary.

Results

Of the 13,272 patients from 55 studies in the entire IPD, we included 6,702 from 49 studies that included 84 treatment sites in 34 countries (Fig 1). We excluded 6,570 patients in total. Six entire studies were excluded (2,235 patients) as they provided only planned duration or did not provide duration data (excluded and included studies were similar, see Supplement 2 in S1 File). Of the included studies, 4,335 patients were excluded: 44 had success but no duration data and 4,291 did not have treatment success. The characteristics of patients excluded from our analysis are presented in Supplementary Table S1 in S1 File.

Ecological analysis of mean treatment duration of site

Descriptions of the site-level characteristics are presented in <u>Table 1</u>. The mean treatment duration of all sites was 22.8 and ranged from 12 to 36 months (see Supplementary Table S2 in <u>S1 File</u> for mean treatment duration of each site).

In univariable analysis, the proportion of patients at the site with past first-line drug use, MDR/RR-TB plus resistance to both FQ and SLI (MDR-FQ+SLI), or resistance to pyrazinamide were associated with longer mean treatment duration at the site. However, in multivariable analysis, only the proportion of patients with MDR-FQ+SLI was associated with longer mean treatment duration of site (Table 1).

Analysis of individual deviation from mean treatment duration of site

The patients included in this analysis are described in Tables 2 and 3. The average total treatment duration was 22.0 months with SD of 4.6 (median 22 [IQR: 19, 24]). In univariable analyses, lower body mass index, past first- and second-line drug use, cavitation or bilateral disease on X-ray, and AFB smear positivity were all associated with longer treatment duration. Resistance to each drug, if tested (except linezolid, which was rarely tested) was associated with longer treatment duration. Longer treatment duration was associated with MDR/RR-TB plus resistance to SLI but FQ sensitive (MDR-SLI), or MDR/RR-TB plus resistance to FQ but SLI sensitive (MDR-FQ), or MDR-FQ+SLI. Within the treatment regimen of a patient, the use of capreomycin, kanamycin, moxifloxacin, levofloxacin, PAS, linezolid, clofazimine, Amx-Clv, clarithromycin, or bedaquiline, as well as greater number of drugs, were all associated with longer treatment duration in univariable analyses.

In the final multivariable model (see Supplement 8 in S1 File for detail on variable selection due to correlation of variables), longer treatment duration was associated with presence of cavitation, AFB smear positivity, HIV infection, past first-line drug use, and MDR/RR-TB with all types of additional resistance (Fig 2). Individual deviation from mean duration of site was also associated with several treatment factors. In contrast to univariable regression results, use of bedaquiline was associated with **shorter** treatment duration by -0.51 (95% CI -0.87 to -0.15) months in adjusted analyses. Longer treatment duration was associated with use of clarithromycin (1.12 months; 95% CI 0.71, 1.53), and with greater number of drugs used, or use of moxifloxacin, kanamycin, capreomycin, or Amx-Clv.

Results were similar when using inverse probability weights for selection into our study population from the entire IPD (Supplementary Table S3 in <u>S1 File</u>). However, in our sensitivity analysis including patients whose treatment durations were imputed, results were substantially different (Supplemental Figure S2 in <u>S1 File</u>).

E-values for the largest regression coefficients from our primary analysis are presented in Supplementary Table S4 (S1 File). For bedaquiline, an unmeasured confounder would need to have a risk ratio associated with both use of bedaquiline and treatment duration of 1.50 to completely explain away the association we observed with bedaquiline. The largest E-value



Fig 1. PRISMA diagram for studies and patients included and excluded from the study population.

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required of an unmeasured confounder to explain away our estimated associations was for use of clarithromycin, while the smallest was for cavitation.

Subgroup analyses. In subgroup analyses (<u>Table 4</u>) the direction of associations between shorter treatment duration and use of bedaquiline remained consistent across all subgroups (except in those with MDR-FQ+SLI), and regardless of disease extent.

Associations between bedaquiline and duration were similar between those with or without past treatment. Additionally, use of Amx-Clv and clarithromycin were consistently associated with longer treatment duration in all subgroups. Body mass index was not associated with treatment duration in any subgroup while HIV was associated with longer duration in those with extensive disease and past tuberculosis treatment.

In other exploratory analyses (Supplemental Table S5 in <u>S1 File</u>) results were similar for bedaquiline, Amx-Clv, clarithromycin, and body mass index. However, HIV was not associated with treatment duration in any exploratory subgroup.

Discussion

With this IPD meta-analysis of 6,702 MDR/RR-TB patients with treatment success, we have applied a novel approach to identify patients who may benefit from shorter MDR/RR-TB treatment. In ecological analysis of site-level factors, the only clinical or treatment characteristic associated with average treatment duration of a site was the proportion of MDR patients with added resistance to FQ and SLI. The lack of associations between mean treatment duration of site with many clinical factors (such as age, HIV infection, past treatment, or other patterns of drug resistance) may indicate that unmeasured factors like physician beliefs, site

				Mean site-specific treatment duration		
Variable: proportion at site unless stated otherwise (n = 84)	Mean (SD)	Median [IQR]	Range	Univariable Months (95% CI)	Multivariable Months (95% CI)	
Clinical characteristics						
Age (mean years)	37.5 (6.1)	37.6 [33.4, 41.5]	21 to 55	0.01 (-0.1, 0.2)	0.1 (-0.1, 0.3)	
Sex (Female)	0.37 (0.2)	0.38 [0.27, 0.48]	0 to 1	-3.5 (-7.8, 0.9)	-2.7 (-7.9, 2.5)	
HIV infection	0.09 (0.2)	0 [0, 0.09]	0 to 0.73	3.0 (-1.9, 7.9)	3.0 (-2.9,.8.8)	
2018 World Bank income category						
Low/lower-middle income—n (%)	13 (15.5)	NE	NE	Ref	Ref	
Upper-middle income—n (%)	31 (36.9)	NE	NE	-1.4 (-4.1, 1.2)	-1.7 (-4.7, 1.2)	
High income—n (%)	40 (47.6)	NE	NE	-2.1 (-4.7, 0.4)	-1.5 (-4.9, 1.9)	
Extensive disease*	0.72 (0.24)	0.74 [0.52, 0.95]	0 to 1	1.4 (-2.3, 5.2)	1.0 (-3.4, 5.4)	
Treatment history and drug resistance						
Past first-line TB drugs	0.70 (0.29)	0.79 [0.47, 0.98]	0 to 1	3.3 (0.2, 6.5)	1.3 (-4.1, 6.7)	
Past second-line TB drugs	0.25 (0.32)	0.08 [0, 0.5]	0 to 1	2.3 (-0.6, 5.1)	-0.2 (-5.4, 4.9)	
Number of effective drugs used	4.41 (0.78)	4.27 [4, 4.9]	2.4 to 7.08	-0.8 (-2.0, 0.3)	-0.2 (-1.6, 1.2)	
MDR/RR-TB + FQ & SLI sensitive	0.51 (0.36)	0.41 [0.18, 0.9]	0 to 1	3.0 (0.5, 5.5)		
MDR/RR-TB + FQ resistant & SLI sensitive	0.14 (0.20)	0.05 [0, 0.19]	0 to 1	1.2 (-3.2, 5.6)	0.1 (-5.2, 5.5)	
MDR/RR-TB + SLI resistant & FQ sensitive	0.13 (0.17)	0.08 [0, 0.18]	0 to 1	-2.4 (-7.4, 2.7)	-0.2 (-6.3, 6.0)	
MDR/RR-TB + SLI & FQ resistance	0.24 (0.31)	0.06 [0, 0.42]	0 to 1	4.1 (1.4, 6.9)	4.6 (0.2, 9.0)	
MDR/RR-TB + Pyrazinamide resistance	0.45 (0.30)	0.44 [0.21, 0.65]	0 to 1	4.3 (0.7, 7.9)		
Drugs used in treatment						
Patients received Bedaquiline	0.32 (0.44)	0 [0, 1]	0 to 1	1.5 (-0.6, 3.5)		
Patients received Linezolid	0.37 (0.40)	0.23 [0, 0.73]	0 to 1	0.03 (-2.2, 2.2)		
Bedaquiline used at site (%)	41 (48.8)	NE	NE	0.17 (-1.6, 2.0)	-0.3 (-2.6, 2.0)	
Linezolid used at site (%)	58 (69.0)	NE	NE	-0.59 (-2.5, 1.3)	-0.3 (-2.8, 2.2)	
Patients with success	79.8 (236.7)	34.5 [11, 76.5]	1 to 2128	-0.03 (-0.12, 0.07)		
Patients treated	131.3 (401.5)	52.5 [15.8, 123.3]	1 to 3626	-0.01 (-0.06, 0.05)	0.0 (-0.1, 0.1)	
Treatment duration (Months)	22.8 (4.1)	22.6 [20.3.24.6]	12 to 36	NE	NE	

 Table 1. Site-level characteristics and their univariable and multivariable associations with the site-specific mean treatment duration in patients with successful treatment outcomes. Estimates and 95% confidence interval (CI) from linear regression models.

Note: Extensive disease is defined as: AFB smear positive at baseline. If AFB smear information missing, then if radiographic findings of cavitation or bilateral disease. If value blank in multivariable coefficient column, then the variable was not included in the multivariable model. NE: not estimated; TB: tuberculosis; FQ: fluoroquinolones; SLI: second-line injectable. Note: proportion of patients receiving bedaquiline/linezolid, MDR/RR-TB FQ & SLI sensitive, MDR/RR-TB plus pyrazinamide resistance, and number of patients with success were not included in the multivariable model as they were highly correlated with other relevant variables that were included. For the multivariable model, R²: 0.24; adjusted R²: 0.05

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conventions, or access to medications are more important determinants of treatment duration. In contrast, several clinical and treatment factors were associated with individual treatment duration in our analysis, which have shown to be associated with treatment outcomes in several prior studies [15, 20, 21, 28]. Hence, our novel approach of using individual deviation from the site-specific mean treatment duration may provide a better method to assess clinical and treatment characteristics association with treatment duration.

By accounting for the mean treatment duration of a site in the duration outcome and by restriction to patients with successful treatment outcomes we aimed to create an outcome variable that accounts for the site-level variation and outcome-dependent complexities inherent in studying duration for treatment of MDR/RR-TB. The finding that factors predicting poor

		Total treatment duration	Individual Deviation in treatment duration from centre mean		
	n (%) unless specified otherwise	mean (SD) Months	mean (SD) Months	Univariable regression estimate months (95% CI)*	
All patients n = 6702		22.0 (4.6)	0.0 (4)		
Clinical characteristics					
$Sex = Male^{\Delta}$	3982 (59.4)	22.1 (4.6)	0.1 (4)	Ref	
Female	2719 (40.6)	21.9 (4.6)	-0.1 (4)	-0.18 (-0.37, 0.02)	
Age (mean (SD))	37.02 (13)	NE	NE	0.01 (-0.003, 0.01)§	
Body mass index (mean (SD))	20.47 (3.84)	NE	NE	-0.04 (-0.07, -0.01)§	
Body mass index category					
Normal	2024 (30.2)	22.4 (4.8)	0 (4.2)	Ref	
Underweight	1028 (15.3)	22.7 (4.3)	0.4 (3.7)	-0.21 (-0.62, 0.19)	
Overweight/Obese	377 (5.6)	22.3 (4.6)	-0.2 (3.9)	0.26 (0.00, 0.52)	
Missing	3273 (48.8)	21.5 (4.5)	-0.1 (3.9)	Not estimated	
2018 World Bank income category					
Low/Low-middle	1226 (18.3)	22.5 (4.2)	0 (3.5)	Ref	
Upper-Middle	3555 (53.0)	22.3 (4)	0 (3.6)	0.01 (-0.26, 0.26)	
High	1921 (28.7)	21.3 (5.6)	0 (4.9)	-0.02 (-0.30, 0.27)	
Smoking					
Ex-smoker or never smoker	1834 (27.4)	22.5 (5.3)	-0.1 (4.8)	Ref	
Current smoker	939 (14.0)	22.5 (5)	0.4 (4.1)	0.17 (-0.09, 0.42)	
Unknown	3929 (58.6)	21.7 (4.1)	-0.1 (3.5)	Not estimated	
HIV					
Negative	4771 (71.2)	22 (4.8)	0 (4.1)	Ref	
Positive	1859 (27.7)	22.1 (3.9)	0.1 (3.5)	0.13 (-0.08, 0.35)	
Unknown	72 (1.1)	22.9 (5.4)	-0.1 (5.1)	Not estimated	
If HIV positive, on ART	1686 (90.7)	22 (3.8)	0 (3.5)	-0.16 (-0.83, 0.50)	
Not on ART	173 (9.3)	23.2 (4.4)	0.5 (4.1)		
Diabetes					
No	3311 (49.4)	22.4 (5)	0 (4.3)	Ref	
Yes	466 (7.0)	21.9 (4.4)	0.3 (3.7)	0.21 (-0.18, 0.59)	
Unknown	2925 (43.6)	21.7 (4)	0 (3.6)	Not estimated	
Cavitation on X-ray					
No	1606 (24.0)	21.7 (4.9)	-0.4 (4.2)	Ref	
Yes	2308 (34.4)	22.5 (5.1)	0.3 (4.3)	0.60 (0.35, 0.86)	
Unknown	2788 (41.6)	21.8 (3.8)	0 (3.6)	0.37 (0.12, 0.61)	
Bilateral disease					
No	1122 (16.7)	21.4 (4.9)	-0.3 (4)	Ref	
Yes	1999 (29.8)	22.2 (4.9)	0.2 (4.1)	0.52 (0.22, 0.81)	
Unknown	3581 (53.4)	22.1 (4.3)	0 (3.9)	0.35 (0.09, 0.62)	
AFB smear result					
Neg	1974 (29.5)	21.4 (4.7)	-0.6 (4)	Ref	
Pos	4280 (63.9)	22.4 (4.5)	0.3 (4)	0.91 (0.70, 1.12)	
Unknown	448 (6.7)	21.2 (4.2)	0 (3.8)	0.65 (0.24, 1.05)	
Extensive disease					
No	2147 (32.0)	21.3 (4.6)	-0.6 (3.9)		
Yes	4512 (67.8)	22.4 (4.6)	0.3 (4)	0.90 (0.70, 1.11)	
Unknown	43 (0.0)	20.9 (4.7)	-0.1 (4.5)	Not estimated	

Table 2. Description of patient characteristics and their association (adjusted for age and sex) with deviation in treatment duration from site mean.

(Continued)

Table 2. (Continued)

		Total treatment duration	Individual Deviation in treatment duration from centre mean		
	n (%) unless specified otherwise	mean (SD) Months	mean (SD) Months	Univariable regression estimate months (95% CI)*	
Treatment history and markers of disease s	everity				
Past TB treatment					
No	2336 (34.9)	21.2 (4.4)	-0.4 (3.8)	Ref	
Yes	4271 (63.7)	22.5 (4.6)	0.2 (4.1)	0.56 (0.36, 0.77)	
Unknown	95 (1.4)	21.7 (4.5)	0.1 (3.8)	0.48 (-0.34, 1.29)	
Past first-line TB drug use					
No	2336 (34.9)	21.2 (4.4)	-0.4 (3.8)	Ref	
Yes	4271 (63.7)	22.5 (4.6)	0.2 (4.1)	0.56 (0.36, 0.76)	
Unknown	95 (1.4)	21.7 (4.5)	0.1 (3.8)	Not estimated	
Past second-line TB drug used					
No	5048 (75.3)	21.7 (4.1)	-0.2 (3.6)	Ref	
Yes	1226 (18.3)	23.3 (5.4)	0.6 (4.6)	0.71 (0.44, 0.98)	
Unknown	428 (6.4)	22.4 (6.2)	0.5 (5.6)	Not estimated	
Pre-treatment Drug susceptibility results					
DST Performed for FQ	6449 (96.2)			Not estimated	
If DST Performed, FQ Resistant = Yes	1172 (18.2)	23.6 (5.8)	0.8 (5)	1.04 (0.77, 1.31)	
If DST Performed, FQ Resistant = No	5277 (81.8)	21.6 (4.1)	-0.2 (3.6)	Ref	
DST Performed for SLIs	6455 (96.3)			Not estimated	
If DST Performed, SLI Resistant = Yes	1629 (25.2)	23 (5.3)	0.5 (4.5)	0.58 (0.36, 0.81)	
If DST Performed, SLI Resistant = No	4826 (74.8)	21.7 (4.2)	-0.1 (3.8)	Ref	
DST Performed for Linezolid	665 (9.9)			Not estimated	
If DST Performed, Linezolid Resistant = Yes	16 (2.4)	21.5 (3.6)	-0.8 (2.8)	-0.76 (-2.74, 1.22)	
If DST Performed, Linezolid Resistant = No	649 (97.6)	21.1 (4.4)	0 (3.7)		
DST Performed for Pyrazinamide	3490 (52.1)			Not estimated	
If DST Performed, Pyrazinamide Resistant = Yes	1859 (53.3)	22.2 (5.4)	0.3 (4.7)	0.51 (0.30, 0.72)	
If DST Performed, Pyrazinamide Resistant = No	1631 (46.7)	21.1 (4.6)	-0.4 (4)	Ref	
DST Performed for Clofazimine	252 (3.8)			Not estimated	
If DST Performed, Clofazimine Resistant = Yes	9 (3.6)	24.4 (5)	2 (5)	Not estimated †	
If DST Performed, Clofazimine Resistant = No	243 (96.4)	21.8 (5.8)	0.1 (4.4)		
DST Performed for Cycloserine‡	2034 (30.3)			Not estimated	
If DST Performed, Cycloserine Resistant = Yes	260 (12.8)	23.4 (6.2)	1 (5.2)	1.16 (0.65, 1.68)	
If DST Performed, Cycloserine Resistant = No	1774 (87.2)	21.9 (5.3)	-0.1 (4.5)		
MDR category					
MDR/RR-TB FQ &SLI sensitive	4337 (64.7)	21.5 (4)	-0.3 (3.5)	Ref	
MDR/RR-TB + FQ resistant & SLI sensitive	929 (13.9)	22.2 (4.5)	0.2 (3.9)	0.48 (0.20, 0.77)	
MDR/RR-TB + SLI resistant & FQ sensitive	475 (7.1)	23.2 (5.8)	0.9 (5.1)	1.24 (0.85, 1.63)	
MDR/RR-TB + SLI & FQ resistance	688 (10.3)	23.9 (5.9)	0.8 (5)	1.08 (0.76, 1.41)	

(Continued)

		Total treatment duration	Individual Deviation in treatment duration from centre	
	n (%) unless specified otherwise	mean (SD) Months	mean (SD) Months	Univariable regression estimate months (95% CI)*
No DST	273 (4.1)	22.5 (5.7)	0 (5)	Not estimated
MDR/RR-TB + SLI & FQ resistance vs. all others				
No	5741 (85.7)	21.8 (4.3)	-0.1 (3.8)	Ref
Yes	688 (10.3)	23.9 (5.9)	0.8 (5)	0.83 (0.52, 1.14)
Unknown	273 (4.1)	22.5 (5.7)	0 (5)	Not estimated

Table 2. (Continued)

https://doi.org/10.1371/journal.pone.0292106.t002

treatment outcomes such as MDR with additional resistance to FQ and/or SLI [1, 3], HIV infection [20], or cavitation [11, 29] were associated with longer treatment duration provides support for the use of this method. Our finding that treatment duration is shorter when bedaquiline was used, is supported by several studies that have established the efficacy of bedaquiline [3, 15, 28], which further supports the use of bedaquiline containing regimens for all MDR/RR-TB patients. Additionally, as this was observational data, we included patient populations in our analysis that were excluded from trials of shorter treatment, such as those with additional resistance to SLIs or FQs [9, 10], low body mass index [11, 29], low HIV CD4 cell counts [9-11, 29], "any comorbidity likely to compromise protocol assessments" [11, 29], or extensive disease and past treatment (the last two groups are not eligible for the 9-month alloral regimen in WHO guidelines [3]). Use of bedaquiline was associated with shorter treatment duration across the majority of subgroups, suggesting that inclusion of patients previously excluded from RCTs [9–11, 29] or considered ineligible for short MDR treatment in guidelines [3] could be included in future trials of shorter bedaquiline and/or pretomanid containing regimens (such as bedaquiline, pretomanid, & linezolid (BPaL), or BPaL plus moxifloxacin, BPaLM) most recently recommended by the WHO [3]. Additionally, our results indicate that certain patients with more complicated clinical profiles, such as MDR/RR-TB patients without additional resistance who also have extensive disease but either no past treatment or no HIV, may benefit from shorter treatment.

Although our analysis indicated that use of bedaquiline was associated with shorter treatment duration, these results require cautious interpretation as our models were constructed for the purposes of hypothesis generation. The association of shorter duration with use of bedaquiline may reflect the preferred use of the drug in regimens with planned shorter durations. However, this was not observed with linezolid or FQs, which are also used in regimens with shorter planned durations. Additionally, we assessed drugs that were received at any time (ever) during treatment, which does not adequately account for regimen changes. Some characteristics and drugs that were associated with longer duration may reflect clinical conventions. For instance, use of low efficacy drugs (e.g. clarithromycin, Amx-Clv, and injectables [15]) may reflect use of drugs in desperation for patients with more complicated disease with longer planned duration. Similar conventions apply to associations with cavitation.

Our study has limitations. Primarily, we conducted this analysis in a population treated between 1993 and 2019, and treatment practises, including use of fluoroquinolones, bedaquiline, and second-line injectables, as well as advancement in antiretroviral therapy (ART) and their uptake, have changed substantially in the last five years [3]. We also did not have data on the number of cavities, only the presence or absence, nor did we have data on level of AFB smear positivity (only positive or negative), and were unable to assess what effect this had on

		Total treatment duration	Individual Deviation in treatment duration from a mean	
All patients n = 6702	n (%) unless specified otherwise	mean (SD) Months	mean (SD) Months	Univariable regression estimate months (95% CI)*
Drugs used in treatment				
Used Bedaquiline Ever During Treatment = Yes	1605 (23.9)	22.4 (3.9)	0.2 (3.5)	0.27 (0.04, 0.49)
No	5097 (76.1)	21.9 (4.8)	-0.1 (4.1)	Ref
Used Ofloxacin Ever During Treatment = Yes	1373 (20.5)	22 (4.2)	-0.1 (3.6)	-0.13 (-0.36, 0.11)
No	5329 (79.5)	22 (4.7)	0 (4.1)	Ref
Used Ciprofloxacin Ever During Treatment = Yes	266 (4.0)	23 (5.8)	0 (5.4)	0.03 (-0.46, 0.52)
No	6436 (96.0)	22 (4.5)	0 (3.9)	Ref
Used Moxifloxacin Ever During Treatment = Yes	3459 (51.6)	22.1 (4.6)	0.2 (4.1)	0.41 (0.22, 0.60)
No	3243 (48.4)	21.9 (4.6)	-0.2 (3.9)	Ref
Used Levofloxacin Ever During Treatment = Yes	1889 (28.2)	21.8 (4.9)	0.1 (4.2)	0.19 (-0.03, 0.40)
No	4813 (71.8)	22.1 (4.4)	-0.1 (3.9)	Ref
Used Linezolid Ever During Treatment = Yes	1594 (23.8)	22.5 (5.1)	0.5 (4.4)	0.63 (0.41, 0.86)
No	5108 (76.2)	21.9 (4.4)	-0.1 (3.8)	Ref
Used Clofazimine Ever During Treatment = Yes	1101 (16.4)	22.5 (4.8)	0.3 (3.8)	0.35 (0.10, 0.61)
No	5601 (83.6)	21.9 (4.5)	-0.1 (4)	Ref
Used Cycloserine/Terizidone Ever During Treatment = Yes	5702 (85.1)	22.1 (4.4)	0 (4.0)	0.19 (-0.08, 0.46)
No	1000 (14.9)	21.7 (5.3)	-0.2 (4.1)	Ref
Used Ethambutol Ever During Treatment = Yes	2895 (43.2)	22 (4.5)	0.1 (3.9)	0.10 (-0.09, 0.29)
No	3807 (56.8)	22.1 (4.7)	0 (4.1)	Ref
Used Pyrazinamide Ever During Treatment = Yes	5175 (77.2)	22 (4.3)	0 (3.8)	-0.13 (-0.36, 0.09)
No	1527 (22.8)	22 (5.5)	0.1 (4.6)	Ref
Used Streptomycin Ever During Treatment = Yes	692 (10.3)	22.5 (5.1)	0.1 (4.5)	0.10 (-0.21, 0.42)
No	6010 (89.7)	22 (4.5)	0 (3.9)	Ref
Used Rifabutin Ever During Treatment = Yes	154 (2.3)	22.8 (6.7)	0.1 (5.8)	0.05 (-0.59, 0.69)
No	6548 (97.7)	22 (4.5)	0 (3.9)	Ref
Used Amikacin Ever During Treatment = Yes	1048 (15.6)	21.8 (4.8)	0 (4.2)	0.06 (-0.21, 0.32)
No	5654 (84.4)	22.1 (4.5)	0 (3.9)	Ref
Used Capreomycin Ever During Treatment = Yes	1446 (21.6)	23.1 (5.6)	0.5 (4.7)	0.66 (0.42, 0.89)
No	5256 (78.4)	21.7 (4.2)	-0.1 (3.8)	Ref
Used Kanamycin Ever During Treatment = Yes	3151 (47.0)	21.9 (3.9)	0.1 (3.5)	0.20 (0.01, 0.39)
No	3551 (53.0)	22.1 (5.1)	-0.1 (4.3)	Ref
Used Ethionamide/Prothionamide Ever During Treatment = Yes	5096 (76.0)	22.1 (4.5)	0 (4)	020 (-0.03, 0.42)
No	1606 (24.0)	21.8 (4.9)	-0.2 (4.1)	Ref
Used PAS Ever During Treatment = Yes	2759 (41.2)	22.7 (5.2)	0.3 (4.4)	0.48 (0.29, 0.68)
No	3943 (58.8)	21.6 (4.1)	-0.2 (3.6)	Ref
Used Amx-Clv Ever During Treatment = Yes	994 (14.8)	24 (6.2)	1 (5.5)	1.27 (0.99, 1.55)
No	5708 (85.2)	21.7 (4.1)	-0.2 (3.6)	Ref
Used Thioacetazone Ever During Treatment = Yes	68 (1.0)	21 (5.4)	0.2 (4)	0.20 (-0.75, 1.16)
No	6634 (99.0)	22 (4.6)	0 (4)	Ref
Used Clarithromycin Ever During Treatment = Yes	485 (7.2)	24.6 (7)	1.4 (6.2)	1.50 (1.14, 1.87)
No	6217 (92.8)	21.8 (4.3)	-0.1 (3.7)	Ref
Used Imipenem Ever During Treatment = Yes	237 (3.5)	23.4 (4.6)	0.4 (4.1)	0.37 (-0.14, 0.89)
No	6465 (96.5)	22 (4.6)	0 (4)	Ref
	. ,			

Table 3. Description of drugs used in treatment and their association (adjusted for age and sex) with deviation in treatment duration from site mean.

(Continued)

Table 3. (Continued)

		Total treatment duration	Individual Deviation in treatment duration from centre mean		
All patients n = 6702	n (%) unless specified otherwise	mean (SD) Months	mean (SD) Months	Univariable regression estimate months (95% CI)*	
Used Meropenem Ever During Treatment = Yes	61 (0.9)	21.1 (5.4)	-0.3 (4.4)	-0.32 (-1.33, 0.68)	
No	6641 (99.1)	22 (4.6)	0 (4)	Ref	
Used Delamanid Ever During Treatment = Yes	114 (1.7)	21 (4.4)	0 (3.6)	-0.02 (-0.76, 0.72)	
No	6588 (98.3)	22 (4.6)	0 (4)	Ref	
Number of drugs (median [IQR])	5 [4, 6]	NE	NE	0.37 (0.29, 0.46)§	
Number of effective drugs (median [IQR])	4 [4, 5]	NE	NE	0.03 (-0.06, 0.12)§	
Number of limited access drugs** (median [IQR])	0 [0, 1]	NE	NE	0.18 (0.09, 0.26)§	
Total treatment duration (median [IQR])	22 [19, 24]	NE	NE	Not estimated	
Deviation in treatment duration (median [IQR])	-0.15 [-2, 2]	NE	NE	Not estimated	

SD: standard deviation; XDR: extensively drug resistant tuberculosis; MDR: multidrug resistant tuberculosis; TB: tuberculosis; AFB: acid-fast bacillus; Amx-Clv: Amoxicillin-Clavulanic Acid

* Regression coefficients were estimated using imputed data and adjusted for age and sex.

† Too few observations to estimate.

‡ Drug susceptibility testing for cycloserine and terizidone combined.

§ per unit increase.

** includes bedaquiline, clofazimine, linezolid, imipenem, and meropenem.

 Δ One subject missing sex.

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duration. Further, studies conducted exclusively in children were excluded from the IPD, and we were not able to assess associations in this population. Additionally, there are site-level differences in treatment protocols that may affect treatment outcomes (availability of drugs), that may not be captured in variables we included in our models. However, as we used the average duration of treatment at a site in our duration outcome, we believe this may account for sitelevel heterogeneity in clinical practice. There is still potential for indication bias affecting duration of treatment in patients with complex profiles which may not be accounted for with our outcome. Although we conducted a 'new user' subgroup analysis [30] in those without previous treatment, this only addresses one aspect that may create indication bias for treatment duration. Because our population included only those with treatment success, our findings may not be generalizable to all patients with MDR/RR-TB, although we conducted inverse probability of selection weighted analyses which indicated no substantial differences in the excluded vs included populations. However, adverse events may be more common with longer treatment, and contribute to treatment failure (and as such were excluded in this study) yet it is possible that patients with adverse events would benefit even more from shorter treatment. Finally, our results do not reflect causal relationships and should be interpreted with caution, with additional consideration that variance of regression estimates may be underestimated due to the statistical selection of variables [31].

Despite that, our study has several strengths. We included a large population of patients who had detailed information on important clinical characteristics and treatment. Additionally, we conducted several subgroup analyses of important patient groups that were previously excluded from trials on shorter treatment [9–11, 29]. We also used E-values to assess unmeasured confounding (where a larger E-values implies a more robust observed estimate). Although plausible that an unmeasured confounder could account for the observed association with bedaquiline, we feel it is unlikely such an important predictor would not have been

Clinical characteristics

Sex (female)	-0.11 (-0.3, 0.08)
Body mass index (per unit increase)	-0.02 (-0.05, 0.01)
Age (per year increase)	0 (0, 0.01)
Bilateral disease on X-ray	0.24 (-0.07, 0.54)
Cavitation on X-Ray	0.3 (0.05, 0.56)
HIV infection	0.33 (0.06, 0.6)
AFB smear positive	0.82 (0.6, 1.04)



Treatment history and drug resistance

Past second line drug use	0.21 (-0.16, 0.58)
MDR with resistance to SLI	0.34 (0.03, 0.65)
Number of drugs used	0.34 (0.21, 0.47)
Past first line drug use	0.38 (0.14, 0.62)
MDR with resistance to both SLI & FQ	0.61 (0.21, 1.02)
MDR with resistance to FQ	0.78 (0.36, 1.21)



Drugs used in treatment

-			
Used Bedaquiline Ever During Treatment	-0.51 (-0.87, -0.15)		
Used Clofazimine Ever During Treatment	-0.15 (-0.51, 0.21)		
Used PAS Ever During Treatment	-0.03 (-0.27, 0.21)		
Used Linezolid Ever During Treatment	0.29 (-0.06, 0.63)	-	
Used Moxifloxacin Ever During Treatment	0.32 (0.09, 0.54)	-	_
Used Kanamycin Ever During Treatment	0.44 (0.2, 0.68)	-	_
Used Capreomycin Ever During Treatment	0.69 (0.41, 0.97)	-	_
Used Amx-Clv Ever During Treatment	0.73 (0.4, 1.06)	-	_
Used Clarithromycin Ever During Treatment	1.12 (0.71, 1.53)	-	
		-1 0) 1 2

Deviation from mean duration of site; months (95% CI)



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included in our data. Although RCTs can provide clearer evidence on optimal duration, these are expensive, time consuming, often lack generalizability, and can test only a limited number of durations and/or regimens at once. As well, evidence from cohorts are useful as potential indicators of success of these regimens in programmatic settings [32, 33]). Our use of observational data from a large population of MDR/RR-TB patients from 84 treatment sites and 34 countries provided evidence that should be more generalizable. We also describe an analysis of characteristics associated with duration at the level of the treatment site. We interpret that the lack of associations between patient or treatment characteristics and our outcome may reflect

				-		_
	Patients with additional SLI & FQ resistance	Patients without FQ and SLI resistance <u>*</u>	Patients with extensive disease	Patients without extensive disease	Patients with past TB treatment	Patients without past TB treatment
Characteristic	months (95% CI)	months (95% CI)	months (95% CI)	months (95% CI)	months (95% CI)	months (95% CI)
Clinical characteristics						
Age (per year increase)	0.02 (-0.01, 0.05)	0 (0, 0.01)	0 (-0.01, 0.01)	0.01 (0, 0.03)	0.01 (0, 0.02)	0 (-0.02, 0.01)
Sex (Female)	0.16 (-0.6, 0.92)	-0.14 (-0.34, 0.06)	-0.26-0.5, -0.02)	0.12 (-0,21, 0.46)	-0.08 (-0.32, 0.17)	-0.17 (-0.48, 0.14)
Body mass index (per unit increase)	0.01 (-0.1, 0.11)	-0.03 (-0.06, 0.01)	-0.02 (-0.06, 0.02)	-0.03 (-0.09, 0.02)	-0.03 (-0.07, 0.01)	-0.01 (-0.07, 0.05)
HIV infection	0.15 (-1.01, 1.31)	0.34 (0.06, 0.62)	0.33 (0.01, 0.66)	0.33 (-0.1, 0.76)	0.38 (0.02, 0.74)	0.22 (-0.18, 0.62)
AFB smear positive	0.77 (-0.08, 1.62)	0.79 (0.56, 1.01)	Not estimated	Not estimated	0.82 (0.52, 1.12)	0.77 (0.44, 1.1)
Cavitation on X-Ray	1.08 (0.1, 2.05)	0.22 (-0.05, 0.49)	Not estimated	Not estimated	0.23 (-0.09, 0.54)	0.59 (0.12, 1.06)
Bilateral disease on X-ray	0.64 (-0.56, 1.85)	0.16 (-0.16, 0.48)	Not estimated	Not estimated	0.36 (-0.01, 0.74)	-0.04 (-0.58, 0.5)
Treatment history and dru	ig resistance					
Past first-line drug use	-0.22 (-1.56, 1.11)	0.4 (0.16, 0.65)	0.43 (0.13, 0.74)	0.34 (-0.05, 0.73)	Not estimated	Not estimated
Past second-line drug use	0.72 (-0.41, 1.86)	0.15 (-0.24, 0.55)	0.13 (-0.32, 0.57)	0.4 (-0.16, 0.97)	Not estimated	Not estimated
Number of drugs used (per unit increase)	0.35 (-0.09, 0.79)	0.33 (0.19, 0.47)	0.4 (0.24, 0.56)	0.16 (-0.06, 0.38)	0.31 (0.14, 0.47)	0.29 (0.07, 0.51)
MDR/RR-TB + FQ resistant & SLI sensitive	Not estimated	Not estimated	0.59 (0.21, 0.98)	-0.2 (-0.75, 0.34)	0.12 (-0.29, 0.54)	0.62 (0.15, 1.08)
MDR/RR-TB + SLI resistant & FQ sensitive	Not estimated	Not estimated	0.92 (0.38, 1.45)	0.46 (-0.24, 1.15)	0.75 (0.26, 1.24)	0.97 (0.1, 1.83)
MDR/RR-TB + SLI & FQ resistance	Not estimated	Not estimated	0.84 (0.32, 1.36)	0.15 (-0.52, 0.81)	0.55 (0.08, 1.02)	0.86 (0.04, 1.68)
Drugs used in treatment						
Used Bedaquiline Ever During Treatment	-0.89 (-2.19, 0.41)	-0.47 (-0.86, -0.09)	-0.51 (-0.94, -0.09)	-0.61 (-1.25, 0.03)	-0.43 (-0.89, 0.03)	-0.52 (-1.09, 0.05)
Used Moxifloxacin Ever During Treatment	0.63 (-0.2, 1.47)	0.26 (0.02, 0.5)	0.42 (0.14, 0.7)	0.07 (-0.31, 0.45)	0.18 (-0.1, 0.46)	0.7 (0.31, 1.09)
Used Linezolid Ever During Treatment	-0.82 (-2.03, 0.38)	0.54 (0.15, 0.93)	0.22 (-0.19, 0.64)	0.44 (-0.18, 1.06)	0.12 (-0.32, 0.56)	0.69 (0.12, 1.27)
Used Clofazimine Ever During Treatment	0.54 (-0.51, 1.58)	-0.43 (-0.84, -0.02)	-0.37 (-0.8, 0.07)	0.3 (-0.31, 0.91)	0.01 (-0.45, 0.47)	-0.25 (-0.85, 0.34)
Used Capreomycin During Treatment	0.46 (-0.44, 1.36)	0.8 (0.49, 1.11)	0.74 (0.4, 1.09)	0.56 (0.08, 1.04)	0.75 (0.39, 1.1)	0.73 (0.25, 1.22)
Used Kanamycin Ever During Treatment	0.64 (-0.58, 1.86)	0.47 (0.22, 0.71)	0.48 (0.18, 0.77)	0.45 (0.02, 0.88)	0.21 (-0.09, 0.52)	0.9 (0.48, 1.31)
Used Amx-Clv Ever During Treatment	0.7 (-0.18, 1.59)	0.77 (0.39, 1.15)	0.84 (0.43, 1.25)	0.51 (-0.03, 1.05)	0.93 (0.53, 1.33)	0.43 (-0.16, 1.03)
Used Clarithromycin During Treatment	0.64 (-0.46, 1.73)	1.19 (0.73, 1.66)	0.88 (0.38, 1.39)	1.6 (0.9, 2.3)	1.25 (0.77, 1.73)	0.72 (-0.1, 1.55)

 Table 4. Associations of individual deviation in treatment duration from site mean with patient characteristics, resistance categories, and drugs used, within specified subgroups. Estimates and 95% confidence interval (CI) from multivariable linear mixed models including all variables shown (unless otherwise specified).

*Includes MDR/RR-TB FQ &SLI sensitive, MDR/RR-TB + FQ resistant & SLI sensitive, MDR/RR-TB + SLI resistant & FQ sensitive. Note: for MDR models were also adjusted for resistance to fluoroquinolone (FQ), second line injectables (SLI), pyrazinamide and cycloserine. For MDR/RR-TB + SLI & FQ resistance, models were also adjusted for resistance to pyrazinamide and cycloserine (not shown for consistency with other subgroups). All models also adjusted for use of PAS. MDR: multidrug resistant tuberculosis; TB: tuberculosis; AFB: acid-fast bacillus; Amx-Clv: Amoxicillin-Clavulinic Acid

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the impact that provider belief and site convention have on duration, a problem which has not been previously described.

Our results produced correlates of individual treatment duration in MDR/RR-TB patients that may help identify patients who would benefit from shorter treatment. We found evidence

that certain patients with more extensive disease and drug resistance may benefit from shorter treatment and could be included in future treatment shortening trials.

Supporting information

S1 File. (DOCX)

Author Contributions

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References

- 1. World Health Organization. Global tuberculosis report 2022. Geneva, Switzerland: World Health Organization, 2022.
- Kwak N, Winters N, Campbell JR, et al. Changes in treatment for multidrug-resistant tuberculosis according to national income. Eur Respir J 2020; 56(5). https://doi.org/10.1183/13993003.01394-2020 PMID: 32586878
- 3. World Health Organization. WHO operational handbook on tuberculosis. Module 4: treatment—drugresistant tuberculosis treatment. Geneva: World Health Organization, 2022.
- 4. Nigam S, Sharma RK, Yadav R, et al. Experiences and needs of patients with MDR/XDR-TB: a qualitative study among Saharia tribe in Madhya Pradesh, Central India. BMJ Open 2021; 11(8): e044698.
- Deshmukh RD, Dhande DJ, Sachdeva KS, et al. Patient and Provider Reported Reasons for Lost to Follow Up in MDRTB Treatment: A Qualitative Study from a Drug Resistant TB Centre in India. PloS one 2015; 10(8): e0135802. https://doi.org/10.1371/journal.pone.0135802 PMID: 26301748
- Shringarpure KS, Isaakidis P, Sagili KD, Baxi RK, Das M, Daftary A. "When Treatment Is More Challenging than the Disease": A Qualitative Study of MDR/RR-TB Patient Retention. PloS one 2016; 11 (3): e0150849.
- 7. Kohler S, Sitali N, Achar J, Paul N. Programme costs of longer and shorter tuberculosis drug regimens and drug import: a modelling study for Karakalpakstan, Uzbekistan. ERJ Open Res 2022; 8(1).
- Van Deun A, Maug AK, Salim MA, et al. Short, highly effective, and inexpensive standardized treatment of multidrug-resistant tuberculosis. Am J Respir Crit Care Med 2010; 182(5): 684–92. https://doi.org/ 10.1164/rccm.201001-0077OC PMID: 20442432
- Nunn AJ, Phillips PPJ, Meredith SK, et al. A Trial of a Shorter Regimen for Rifampin-Resistant Tuberculosis. N Engl J Med 2019; 380(13): 1201–13. <u>https://doi.org/10.1056/NEJMoa1811867</u> PMID: 30865791
- Goodall RL, Meredith SK, Nunn AJ, et al. Evaluation of two short standardised regimens for the treatment of rifampicin-resistant tuberculosis (STREAM stage 2): an open-label, multicentre, randomised, non-inferiority trial. Lancet 2022; 400(10366): 1858–68. https://doi.org/10.1016/S0140-6736(22) 02078-5 PMID: 36368336
- Conradie F, Bagdasaryan TR, Borisov S, et al. Bedaquiline-Pretomanid-Linezolid Regimens for Drug-Resistant Tuberculosis. N Engl J Med 2022; 387(9): 810–23. https://doi.org/10.1056/NEJMoa2119430 PMID: 36053506

- Mok J, Lee M, Kim DK, et al. 9 months of delamanid, linezolid, levofloxacin, and pyrazinamide versus conventional therapy for treatment of fluoroquinolone-sensitive multidrug-resistant tuberculosis (MDR-END): a multicentre, randomised, open-label phase 2/3 non-inferiority trial in South Korea. Lancet 2022; 400(10362): 1522–30. https://doi.org/10.1016/S0140-6736(22)01883-9 PMID: 36522208
- Ahmad Khan F, Salim MAH, du Cros P, et al. Effectiveness and safety of standardised shorter regimens for multidrug-resistant tuberculosis: individual patient data and aggregate data meta-analyses. Eur Respir J 2017; 50(1). https://doi.org/10.1183/13993003.00061-2017 PMID: 28751411
- 14. World Health Organization. Consolidated guidelines on drug-resistant tuberculosis treatment. Geneva, Switzerland, 2019.
- Collaborative Group for the Meta-Analysis of Individual Patient Data in MDRTBt, Ahmad N, Ahuja SD, et al. Treatment correlates of successful outcomes in pulmonary multidrug-resistant tuberculosis: an individual patient data meta-analysis. Lancet 2018; 392(10150): 821–34.
- Bastos ML, Lan Z, Menzies D. An updated systematic review and meta-analysis for treatment of multidrug-resistant tuberculosis. Eur Respir J 2017; 49(3). https://doi.org/10.1183/13993003.00803-2016 PMID: 28331031
- Ahuja SD, Ashkin D, Avendano M, et al. Multidrug resistant pulmonary tuberculosis treatment regimens and patient outcomes: an individual patient data meta-analysis of 9,153 patients. PLoS medicine 2012; 9(8): e1001300. https://doi.org/10.1371/journal.pmed.1001300 PMID: 22952439
- World Health Organization. Public call for individual patient data on treatment of rifampicin and multidrug-resistant (MDR/RR-TB) tuberculosis.. 2018. https://www.who.int/news-room/articles-detail/publiccall-for-individual-patient-data-on-treatment-of-rifampicin-and-multidrug-resistant-(mdr-rr-tb)tuberculosis (accessed 10 Nov 2022.
- World Health Organization. Public call for individual patient data on treatment of drug-resistant tuberculosis. 2019. https://www.who.int/news/item/01-08-2019-public-call-for-individual-patient-data-ontreatment-of-drug-resistant-tuberculosis (accessed 10 Nov 2022.
- Bisson GP, Bastos M, Campbell JR, et al. Mortality in adults with multidrug-resistant tuberculosis and HIV by antiretroviral therapy and tuberculosis drug use: an individual patient data meta-analysis. Lancet 2020; 396(10248): 402–11. https://doi.org/10.1016/S0140-6736(20)31316-7 PMID: 32771107
- Campbell JR, Chan ED, Falzon D, et al. Low Body Mass Index at Treatment Initiation and Rifampicin-Resistant Tuberculosis Treatment Outcomes: An Individual Participant Data Meta-Analysis. Clin Infect Dis 2022. https://doi.org/10.1093/cid/ciac322 PMID: 35476134
- Laserson KF, Thorpe LE, Leimane V, et al. Speaking the same language: treatment outcome definitions for multidrug-resistant tuberculosis. The international journal of tuberculosis and lung disease: the official journal of the International Union against Tuberculosis and Lung Disease 2005; 9(6): 640–5. PMID: 15971391
- **23.** World Health Organization. Definitions and reporting framework for tuberculosis—2013 revision. Geneva: World Health Organization, 2013.
- van Buuren S, Groothuis-Oudshoorn K. mice: Multivariate Imputation by Chained Equations in R. Journal of Statistical Software 2011; 45(3): 1–67.
- 25. Zou H. The Adaptive Lasso and Its Oracle Properties. Journal of the American Statistical Association 2006; 101(476): 1418–29.
- VanderWeele TJ, Ding P. Sensitivity Analysis in Observational Research: Introducing the E-Value. Annals of internal medicine 2017; 167(4): 268–74. https://doi.org/10.7326/M16-2607 PMID: 28693043
- R Core Team (2020). R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria, URL https://wwwR-projectorg/.
- Ndjeka N, Campbell JR, Meintjes G, et al. Treatment outcomes 24 months after initiating short, all-oral bedaquiline-containing or injectable-containing rifampicin-resistant tuberculosis treatment regimens in South Africa: a retrospective cohort study. Lancet Infect Dis 2022; 22(7): 1042–51. <u>https://doi.org/10.1016/S1473-3099(21)00811-2 PMID</u>: 35512718
- 29. Conradie F, Diacon AH, Ngubane N, et al. Treatment of Highly Drug-Resistant Pulmonary Tuberculosis. N Engl J Med 2020; 382(10): 893–902. https://doi.org/10.1056/NEJMoa1901814 PMID: 32130813
- Yoshida K, Solomon DH, Kim SC. Active-comparator design and new-user design in observational studies. Nat Rev Rheumatol 2015; 11(7): 437–41. <u>https://doi.org/10.1038/nrrheum.2015.30</u> PMID: 25800216
- Leeb H, Pötscher BM. Model Selection and Inference: Facts and Fiction. Econometric Theory 2005; 21 (1): 21–59.
- 32. Acuna-Villaorduna C, Jacobson KR, Horsburgh CR, Canning M, Sinha P. Initial experience with BPaLbased regimens to treat multidrug-resistant TB. The international journal of tuberculosis and lung

disease: the official journal of the International Union against Tuberculosis and Lung Disease 2023; 27 (8): 649–50. https://doi.org/10.5588/ijtld.23.0185 PMID: 37491751

 Haley CA, Schechter MC, Ashkin D, et al. Implementation of BPaL in the United States: Experience using a novel all-oral treatment regimen for treatment of rifampin-resistant or rifampin-intolerant TB disease. Clin Infect Dis 2023. https://doi.org/10.1093/cid/ciad312 PMID: 37249079