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# Potential of Pan-Tuberculosis Treatment to Drive Emergence of Novel Resistance

## Appendix

### **Supplementary Methods**

We estimated the proportion durably cured by different regimens in a separate piece of work (T.S. Ryckman et al, unpub. data, 2024), which followed a similar approach to that of Ryckman et al. (1) and is reproduced in brief here and available at GitHub (https://github.com/rycktessman/pan-tb-modeling), with parameter values shown in Appendix Table 2. This captured regimen assignment and initiation, pre-treatment losses to follow-up which varied by regimen assignment, and outcomes of treatment, and ultimately resulted in an estimation of the proportion of diagnosed patients who achieved durable microbiological cure. The probability of durable cure depended on regimen efficacy, duration, ease of adherence, forgiveness (i.e., the extent to which durable cure can occur despite missed doses), and resistance. Efficacy was defined as the proportion of patients curable by the regimen under optimal conditions of perfect adherence, retention in care, and complete initial regimen susceptibility; for the standard of care, efficacy estimates were based on clinical trial data (2,3). Starting from each regimen's efficacy, the probability of cure was adjusted downward for resistance to drugs in the regimen, early treatment discontinuation, and missed doses while on treatment. Discontinuation and adherence with standard of care regimens were based on programmatic data and control groups in trials of adherence-improving interventions, respectively. The impact of adherence varied by regimen forgiveness (4); for more forgiving regimens, more doses could be intermittently missed without affecting the probability of cure.

The hypothetical pan-TB regimen was informed by the WHO's minimal target regimen profile and ongoing regimen development efforts (5). In particular, the oral regimen was modeled as easier to adhere to, at least as forgiving, of shorter duration (3.5 months), and

as efficacious and safe as HRZE – and consisting of drugs with a lower population-wide prevalence of resistance than rifampin.

Durable cure and resistance acquisition rates depended on the resistance phenotype given different regimens (Appendix Table 3). When resistance to all drugs was present, the potential for cure by individualized regimens was parameterized based on longer regimens used for RR-TB before use of bedaquiline, pretomanid, or linezolid ("conventional second-line regimen"). When susceptibility to either B or X was retained (i.e., for RR/BR-TB or RR/XR-TB), the inclusion of that drug in an "X-based" or "B-based" individualized regimen would restore half of the incremental potential for cure that BX would offer for fully-susceptible TB. For a B-based individualized regimen (and similarly for an X-based regimen), the probability of acquiring resistance to B was estimated as the mean of the risk of acquisition of resistance during treatment of (i) drug susceptible TB with BX and (ii) XR-TB with BX.

### **Supplementary Results**

Parameter values used for this illustration are the mean values we calculated previously (Appendix Table 1). For this set of parameter values, the pan-TB scenario resulted in fewer deaths, less treatment failure, and less drug resistance; however, in both scenarios, most TB that persisted after retreatment was DS-TB.

The likelihood of durable cure is higher under the pan-TB scenario irrespective of underlying prevalence of resistance in the population. This is a result of assumptions about the high rate of durable cure for the BX regimen, and the relatively minimal effect of existing resistance to regimen components on this. The likelihood of both increases as the prevalence of RR-TB increases (when RR-TB is likely to be undertreated in the standard of care scenario) and decreases as the prevalence of B resistance increases (when BR-TB is likely to be undertreated in the Pan-TB scenario).

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Appendix Table 1. Parameter values and sources.

Parameter	Description	Value*	Source
Regimen effectiveness			
Er	Proportion of DS-TB durably cured by rifamycin-	70.9% [58.5- 79.3%]	(T.S. Ryckman et al, unpub. data, 2024)
E <sub>BX</sub>	Proportion of DS-TB or RR-TB durably cured by	76.3% [68.8- 83.1%]	(T.S. Ryckman et al, unpub. data, 2024), based on a
Eind	Proportion of TB durably curred by individualized	43.9% [33.7- 53.6%]	(T.S. Ryckman et al, unpub. data, 2024)
CFR	Proportion of poor outcomes (i.e., no durable cure) that result in death	48.3% [40.3- 52.5%]	(6), uncertainty taken from regional variation
Impact of resistance on cure			
P <sub>R</sub>	Risk ratio of cure for rifamycin-based regimen given initial RR-TB	0.35 [0.22- 0.5]	(T.S. Ryckman et al, unpub. data, 2024) (7,8)
Рв	Risk ratio of cure for pan-TB regimen given	0.75 [0.54- 0.91]	(T.S. Ryckman et al, unpub. data, 2024) (9)
P <sub>X</sub>	Risk ratio of cure for pan-TB regimen given	0.75 [0.54- 0.91]	Assumed to be similar to $P_{B}$
P <sub>BX</sub>	Risk ratio of cure for pan-TB regimen given BR- and XR-TB	0.35 [0.22- 0.5]	Assumed to be similar to $P_R$
Resistance acquisition			
S <sub>R</sub>	Probability of acquired resistance to R after rifamycin-based treatment (if initially RS-TR)	0.6% [0.3- 1.2%]	(10,11)
S₅	Probability of acquired resistance to B after pan- TB treatment (if initially BS and XS)	1% [0.3%–2.3%]	(9,12–14), based on observational studies and trial results; point estimate reflects where these two intersect
Sx	Probability of acquired resistance to X after pan- TB treatment (if initially BS and XS)	1% [0.3%–2.3%]	Assumed to be similar to $S_B$ , based on (2)
Q	Risk ratio for B or X resistance acquisition given pre-existing X or B resistance respectively		Assumption based on (2,15) for other drugs
Drug susceptibility testing			
R_soc <sub>new</sub>	R DST for new patients, standard of care scenario	44.10% [33.1%–55.1%]	(6) weighted by the proportion with bacteriological confirmation with an assumed 25% uncertainty interval
BX_soc <sub>new</sub>	B and X DST coverage for new patients with known RB-TB, standard of care scenario	49.0% [36.8%–61.3%]	(6) using fluoroquinolone testing coverage with an
R_soc <sub>retR</sub>	R DST coverage for retreatment patients previously treated with R, standard of care scenario	80.0% [60.0%–100%]	Assumed higher than R_soc <sub>new</sub>
R_soc <sub>retBX</sub>	R DST coverage for retreatment patients previously treated with BX, standard of care scenario	100%	Assumption
BX_soc <sub>retR</sub>	B and X DST coverage for retreatment patients previously treated with R with known RR-TB, standard of care scenario	49.0% [36.8%–61.3%]	Assumed to be similar to new patients in the standard of care scenario
BX_soc <sub>retBX</sub>	B and X DST coverage for retreatment patients previously treated with BX with known RR-TB, standard of care scenario	60.0% [45.0%–75.0%]	Assumption, higher than retreatment patients in the standard of care scenario

Parameter	Description	Value*	Source		
R_pan	R DST coverage for retreatment patients, pan-	44.1% [33.1%–55.1%]	Assumed to be similar to new patients in the standard		
	TB scenario		of care scenario		
BX_pan	B and X DST coverage for retreatment patients	0% or 49.0% [36.8%–61.3%]	Assumed to be similar to new patients in the standard		
	with known RR-TB, pan-TB scenario		of care scenario		
Baseline prevalence of resistance					
prev <sub>DS</sub>	Initial prevalence of DS-TB	95.70%	(6)		
prev <sub>RR</sub>	Initial prevalence of RR-TB	4.20%	(6) weighting new and previously treated patients		
prev <sub>BR</sub>	Initial prevalence of BR-TB	0.20%	(9,16)		
prev <sub>XR</sub>	Initial prevalence of XR-TB	0.00%	Assumption		
prevrrbr	Initial prevalence of RR/BR-TB	0.09%	(12,17–19)		
prev <sub>RRXR</sub>	Initial prevalence of RR/XR-TB	0.00%	Assumption		
prev <sub>BRXR</sub>	Initial prevalence of BR/XR-TB	0.00%	Assumption		
prev <sub>RRBRXR</sub>	Initial prevalence of RR/BR/XR-TB	0.00%	Assumption		

\*All parameters were assumed to follow β distribution (fitted to the median and 95% uncertainty interval shown) except for the parameter Q, which followed a uniform distribution. DS-TB = drug susceptible tuberculosis, RR-TB = rifampin-resistant tuberculosis, BR-TB = diarylquinoline resistant tuberculosis, XR-TB = tuberculosis resistant to additional novel drug X.

#### Appendix Table 2. Parameters used to estimate durable cure rates shown in Appendix Table 1.

		Estimate [95%		
		uncertainty		
Parameter	Regimen	interval]	Distribution used for parameter sampling*	Sources/Notes
Pre-treatment LTFU if assigned	-	13% [8-19%]	Normal (mean 0.134, sdev 0.028)	Subbaraman et al. 2016 (20); Naidoo et al. 2017
to RS SOC or Pan-TB regimen				(21)
Additional pre-treatment LTFU if	_	16% [7-27%]	Beta (8, 42)	Based on Subbaraman et al. 2016 (20), Cox et
assigned to a separate care				al. 2017 (22), and WHO notifications data (6)
pathway (RR SOC or				
individualized regimen)				
Weekly probability of early	India	0.13% [0.08-	Beta (26, 19923)	WHO data from a range of countries (6)
discontinuation		0.18%]		Assumed to be constant over time based on
	South Africa	0.41% [0.31-	Beta (58, 14168)	Kruk et al. 2008 (23)
_		0.52%]		
	Philippines	0.14% [0.08-	Beta (20, 14824)	
		0.20%]		
Efficacy	RS-TB SOC	95% [93-97%]		Gegia et al. 2017 (2)
_	Pan-TB TRP	95% [93-97%]	Same as RS SOC, not	modeled independently
	Individualized	75% [67-83%]	Beta (75, 25)	Based on outcomes of MDR-TB patients pre-
				BPaL/BPaLM (24)
Duration	RS-TB SOC	24 weeks	NA – no uncertainty in duration was modeled	WHO guidelines (25)
	Pan-TB	14 weeks		WHO target regimen profile (minimal target) (5)
	Individualized	18 months		Based on RR regimens pre-BPaL/BPaLM
% patients w/ < 70% adherence	RS-TB SOC	38% [28-48%]	Multinomial (100, 0.379)	Median of control groups in 3 adherence-
				improving intervention studies (26–28)
	Pan-TB	14% [12-16%]	Multinomial (100, 0.14)	Best of control groups in 3 adherence-improving
				3 intervention studies (26–28), based on
				minimal TRP target of better tolerability than the
_				standard of care (5)
	Individualized	38% [28-48%]	Same as RR-TB	Assumed same as BPaL/BPaLM

		Estimate [95%		
	<b>—</b> ·	uncertainty		
Parameter	Regimen	interval	Distribution used for parameter sampling*	Sources/Notes
% patients w/ 70-90%	RS-TB SOC	31% [18-46%]	Multinomial (100, 0.312)	Median of control groups in 3 adherence-
adherence				improving intervention studies (26–28)
	Pan-TB	34% [31-38%]	Multinomial (100, 0.35)	Best of control groups in 3 adherence-improving
				3 intervention studies (26–28), based on
				minimal TRP target of better tolerability than the
				standard of care (5)
	Individualized	31% [18-46%]	Same as RR-TB	Assumed same as HRZE.
% patients w/ ≥ 90% adherence	RS-TB SOC	31% [22-40%]	Multinomial (100, 0.309)	Median of control groups in 3 adherence-
				improving intervention studies (26–28)
	Pan-TB	51% [47-55%]	Multinomial (100, 0.51)	Best of intervention groups in 3 adherence-
				improving 3 intervention studies (26–28), based
				on minimal TRP target of better tolerability than
				the standard of care (5)
	Individualized	31% [22-40%]	Same as RR-TB	Assumed same as HRZE.
Forgiveness (nonadherence	RS-TB SOC	10%	NA – no uncertainty was modeled in the	Imperial et al. (4)
threshold above which	Pan-TB oral	15%	forgiveness thresholds, but uncertainty was	Value from WHO minimal TRP (5)
probability of cure < efficacy)	Individualized	10%	included in the relative probability of cure above	Assumed same as HRZE.
			vs. below the forgiveness threshold (next row).	
Relative probability of cure if	All	82% [63-94%]	Simulated using the confidence intervals from	Imperial et al. (4)
missed doses exceed the			Imperial et al. (4), assuming Wald distributions.	
forgiveness threshold			-	

"RS" = rifampin-susceptible; "SOC" = standard of care; "HRZE" = 6 months of isoniazid, rifampin, pyrazinamide, and ethambutol (the standard of care regimen for RS-TB); "BPaLM" = 6 months of bedaquiline, pretomanid, linezolid, and moxifloxacin (the standard of care regimen for RR-TB). "TRP" = target regimen profile.

Both point estimates/means and uncertainty intervals have been estimated from the sources indicated in the "Sources/Notes" column, unless otherwise noted.

\*For the uncertainty distributions (column 4), the normal distribution is displayed with mean and standard deviation in parentheses, the beta distribution is displayed with alpha and beta in parentheses, where the mean of a beta distribution is equal to alpha divided by the sum of alpha and beta, the multinomial distribution is displayed with size and probability parameters in parentheses, and the gamma distribution is displayed with shape and scale parameters in parentheses.

Ap	pendix	Table 3	. Durable cure	and acquisit	on of resistance	for different resi	stance phenoty	pes aiven diffe	erent regimens	(assuming mean	parameter values fro	om Appendix Tat	ble 1).
										(			

	_	Resistance type							
Outcome	Regimen	DS-TB	RR-TB	BR-TB	XR-TB	RR+ BR-TB	RR+ XR-TB	BR+ XR-TB	RR+BR+ XR-TB
Durable cure	R	E <sub>R</sub> = 70.9%	E <sub>R</sub> ·P <sub>R</sub> = 24.8%	E <sub>R</sub> = 70.9%	E <sub>R</sub> = 70.9%	E <sub>R</sub> ·P <sub>R</sub> = 24.8%	E <sub>R</sub> ·P <sub>R</sub> = 24.8%	E <sub>R</sub> = 70.9%	E <sub>R</sub> ·P <sub>R</sub> = 24.8%
	BX	E <sub>BX</sub> = 79.7%	E <sub>BX</sub> = 79.7%	$E_{BX} \cdot P_{B} = 59.8\%$	E <sub>BX</sub> ·P <sub>X</sub> = 59.8%	E <sub>BX</sub> ·P <sub>B</sub> = 59.8%	E <sub>BX</sub> ·P <sub>X</sub> = 59.8%	E <sub>BX</sub> ·P <sub>BX</sub> = 27.9%	E <sub>BX</sub> ·P <sub>BX</sub> = 27.9%
	B-based	_	_	_	_	-	69.70%	_	_
	X-based	_	_	_	_	69.70%	_	_	_
	Conv 2nd- line	_	_	_	_	_	_	_	$E_{ind} = 43.9\%$
<b>RR</b> Acquisition	R	S <sub>R</sub> = 0.6%	_	S <sub>R</sub> = 0.6%	S <sub>R</sub> = 0.6%	-	-	S <sub>R</sub> = 0.6%	-
BR Acquisition	BX	S <sub>B</sub> = 1.0%	S <sub>B</sub> = 1.0%	_	S <sub>B</sub> ·Q = 7.5%	-	S <sub>B</sub> ·Q = 7.5%	-	-
	B- based	_	_	_	_	_	4.30%	_	-
XR Acquisition	BX	S <sub>X</sub> = 1.0%	S <sub>X</sub> = 1.0%	S <sub>X</sub> ·Q = 7.5%	-	S <sub>X</sub> ·Q = 7.5%	-	_	-
	X- based	_	_	_	_	4.30%	_	_	_

\*TB = tuberculosis, DS-TB = drug-susceptible tuberculosis, RR-TB = rifampin-resistant tuberculosis, BR-TB = diarylquinoline resistant tuberculosis, XR-TB = tuberculosis resistant to additional novel drug X, ER = proportion of DS-TB durably cured by rifamycin-based regimen, E<sub>BX</sub> = proportion of DS-TB durably cured by pan-TB regimen, E<sub>md</sub> = proportion of TB durably cured by individualized regimen, P<sub>R</sub> = Risk ratio of cure for rifamycin-based regimen given initial RR-TB, P<sub>B</sub> = Risk ratio of cure for pan-TB regimen given initial RR-TB, P<sub>B</sub> = Risk ratio of cure for pan-TB regimen given initial RR-TB, P<sub>B</sub> = Risk ratio of cure for pan-TB regimen given initial RR-TB, P<sub>B</sub> = Risk ratio of adquired RR-TB after rifamycin-based treatment (if initially RS-TB), S<sub>B</sub> = Probability of acquired BR-TB after pan-TB treatment (if initially BS and XS), S<sub>X</sub> = Probability of acquired XR-TB after pan-TB treatment (if initially BS and XS), Q = Risk ratio for D or X resistance acquision given pre-existing X or B resistance, respectively.



**Appendix Figure 1.** Treatment pathways for previously treated patients, comparing the Pan-TB scenario (left) with standard of care (right). TB = tuberculosis, RS-TB = rifampin-susceptible tuberculosis, RR-TB = rifampin-resistant tuberculosis, BR-TB = diarylquinoline resistant tuberculosis, XR-TB = tuberculosis resistant to additional novel drug X, R DST = rifampin drug susceptibility testing, B/X DST = diarylquinoline and other novel drug(s) susceptibility testing, R = rifampin-based regimen, BX = pan-TB regimen.



**Appendix Figure 2.** Sankey diagram using mean parameter values for the (a) standard of care and (b) pan-TB scenario. Colors indicate the final treatment outcome. DS = drug susceptible tuberculosis, RR = rifampin-resistant tuberculosis, Novel resistance = tuberculosis resistant to a diarylquinoline and/or novel drug X.



**Appendix Figure 3.** Probability that pan-TB leads to a higher durable cure rate than the standard of care after 1 cohort of patients for varying initial prevalence of resistance. Red indicates where pan-TB TB performs better, blue where SoC performs better. Both RR-TB and BR-TB are varied as a proportion of all TB, where RR+BR-TB is the product of both. No other forms of resistance are initially present. X indicates current estimated prevalence of resistance globally.



**Appendix Figure 4.** Effect of difference in cure rates on relative performance of regimens for varying prevalence of B resistance.



Risk ratio of cure given resistance for novel drugs

**Appendix Figure 5.** Probability that the pan-TB scenario leads to higher durable cure compared to the standard of care after 10 cohorts. Red indicates where pan-TB TB performs better, blue where SoC performs better. Note parameters values (resistance acquisition rate and risk ratio of cure) are varied for both novel drug types simultaneously.



**Appendix Figure 6.** Univariate sensitivity analysis, sampling a parameter set and fixing each parameter in turn at the extremes of its 95% uncertainty interval, where the mean estimate and upper bound of the uncertainty interval for acquisition of resistance to novel drugs have been increased. We based these new estimates on high rates of resistance acquisition seen under programmatic

conditions (e.g., in centers that were not green-light approved), such that the probability of acquired resistance to B after pan-TB treatment (if initially BS and XS) = 2.3% [0.3%–8%], and the probability of acquired resistance to X after pan-TB treatment (if initially BS and XS) = 1% [0.3%–8%]. Comparing likelihood over that durable cure in the pan-TB scenario is greater than the standard of care scenario after (a) one cohort of treatment (b) ten cohorts. Blue circles represent low parameter values, red circles high parameter values.



**Appendix Figure 7.** Prevalence of "complex" resistance to both R and either B and/or X over multiple cohorts where (a) B/X DST availability for retreatment patients with known RR-TB is zero, or (b) R DST availability for retreatment patients is 100%. Shaded areas indicate 95% uncertainty intervals. DS-TB = drug susceptible tuberculosis, RR-TB = rifampin-resistant tuberculosis, BR-TB = diarylquinoline resistant tuberculosis, XR-TB = tuberculosis resistant to novel drug X.



**Appendix Figure 8.** Prevalence of "complex" resistance both to R and to either B and/or X after 10 cohorts for (a) standard of care and (b) pan-TB.