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# Changes in weight-for-age z-score predict effectiveness of childhood tuberculosis therapy

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Key words: multidrug-resistant tuberculosis, treatment outcome, nutritional status, growth and development, Peru

Running title: Early weight change predicts TB outcome

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Summary: In a cohort of children treated for tuberculosis disease, the degree of weight change early in therapy predicted final treatment outcome. This finding can lead to improved management and clinical trial design in childhood tuberculosis therapy.

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# ABSTRACT

<u>Background</u>: International guidelines recommend monitoring weight as an indicator of therapeutic response in childhood tuberculosis (TB) disease. This recommendation is based on observations in adults. In this study, we evaluated the association between weight change and treatment outcome, the accuracy of using weight change to predict regimen efficacy, and whether successfully treated children achieve catch-up weight gain.

<u>Methods</u>: We enrolled children treated for drug-susceptible TB disease (Group 1) and multidrugresistant TB disease (Group 2) in Peru. We calculated the change in weight-for-age z-score ( $\Delta$ WAZ) between baseline and the end of treatment months 2-5 for Group 1, and between baseline and months 2-8 for Group 2. We used logistic regression and generalized estimating equation (GEE) models to evaluate the relationship between  $\Delta$ WAZ and outcome. We plotted receiver operating characteristic curves to determine the accuracy of  $\Delta$ WAZ for predicting treatment failure/death.

<u>Results</u>: Groups 1 and 2 included 100 and 94 children, respectively. In logistic regression, lower  $\Delta$ WAZ in months 3-5 and month 7 was associated with treatment failure/death in Groups 1 and 2, respectively. In GEE models, children in both group who experienced treatment failure or death had lower  $\Delta$ WAZ than successfully treated children.  $\Delta$ WAZ predicted treatment failure/death with 60-90% sensitivity and 60-86% specificity in months 2-5 for Group 1 and months 7-8 for Group 2. All successfully treated children—except Group 2 subjects with unknown microbiologic confirmation status—achieved catch-up weight gain.

Conclusions: Weight change early in therapy can predict outcome of childhood TB treatment.

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#### INTRODUCTION

Each year, an estimated approximately one million children become sick with develop tuberculosis (TB) disease [1, 2]. Failure to thrive—which refers to i.e., weight loss or lack of expected weight gain—is a well-recognized complication of TB disease [3]. International guidelines recommend using weight gain as an indicator of good therapeutic response in childhood TB [3-5]\_; however, tThis recommendation is based on expert opinion. While studies in adults have shown associations between weight gain and good therapeutic response in adults with TB [6-13], and few studies have described weight trends in children on TB treatment [14, 15].

From a clinical perspective, the early identification of children who are failing therapy allows for timely intervention, such as adherence support or regimen modification. From a research perspective, a well-validated surrogate endpoint would facilitate clinical trials. For adults with TB disease, sputum smear and/or culture conversion are used to evaluate regimen efficacy [16]. However, <30% of children have microbiologically confirmed disease [17]. Resolution of radiographic abnormalities and clinical symptoms are suboptimal markers of treatment response: radiographic abnormalities lag behind clinical response, and <u>symptom</u> reporting of symptoms-is subjective. Therefore, there is a need to evaluate other objective markers—like weight change— of treatment response in children.

Another important knowledge gap is whether children who are successfully treated for TB disease achieve catch-up weight gain. Characterizing weight reconstitution among TB-affected

children could identify needs for nutritional supplementation and help clinicians provide anticipatory guidance to families.

We addressed these knowledge gaps using data from children treated for TB disease in Lima, Peru. Our cohort included children treated for drug-susceptible TB and children treated for multidrug-resistant (MDR)-TB, caused by strains of *Mycobacterium tuberculosis* resistant to at least isoniazid and rifampicin. We had three objectivesWe aimed to: (1) to evaluate the association between weight change early in therapy and treatment outcome; (2) to determine the accuracy of weight change as a predictor of regimen efficacy; and (3) to assess whether successfully treated children achieve catch-up weight gain.

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# METHODS

#### Setting and study population

This study took place in Lima, Peru's capital and home to 30% of its population of 32 million [18]. Lima accounts for 54% of the 31,000 TB cases notified annually in the country [19]. Peru has a human immunodeficiency virus (HIV) prevalence of 0.3% among 15-49 year-olds [20].

Peru's National TB Program (NTP) provides free <u>TB</u> treatment to <u>TB patients</u> at public health centers, where clinical charts for these patients are kept. Additionally, the NTP maintains medical records of all patients treated for MDR-TB in Peru.

We enrolled children treated for TB that was microbiologically confirmed or clinically diagnosed according toper NTP guidelines [21, 22]. We enrolled all children  $\leq$ 17 years of age who began treatment for drug-susceptible TB between 2012 and 2014 at eight health centers with high volumes of TB patients and longstanding collaborations with our research group. We also enrolled all children  $\leq$ 15 years of age who initiated MDR-TB therapy anywhere in the Lima metropolitan area between 2005 and 2009. We did not include 16- and 17-year-olds who initiated MDR-TB therapy because their clinical data were not available to us. We divided the participants into two groups. Group 1 consisted of children initially treated for drug-susceptible TB, which included subjects who failed treatment for drug-susceptible TB disease and were subsequently diagnosed with MDR-TB. Group 2 consisted of participants with confirmed MDR-TB (i.e. they had their own *M. tuberculosis* isolate with drug susceptibility test (DST)

confirm<u>eding</u> resistance to at least isoniazid and rifampin) or probable MDR-TB (i.e. they lacked their own <u>DST-isolate</u> but had a presumed source case with confirmed MDR-TB) [23]. Some children with MDR-TB had extensively drug-resistant (XDR)-TB, caused by MDR strains of *M. tuberculosis* with additional resistance to a fluoroquinolone and a second-line injectable agent. During the study period, DSTs were performed with the proportional method, microscopic observation drug susceptibility [24], and/or the nitrate reduction assay [25].

Weight-for-age Z-scores

Weight-for-age z-score (WAZ) is a measurecompares of how a child's weight relates to children of the same age and sex from a healthy reference population. A-Ddistributions of weight-for-age measures is-are created for each sex over the spectrum of ages, with; the z-score being is the number of standard deviations (SDs) from the mean of that distributionfrom. WAZ of 0 represents the age- and sex-appropriate mean. The change in WAZ between two time points, or  $\Delta$ WAZ, can be used to evaluatedenotes weight change in comparison to the reference population.  $\Delta$ WAZ of 0 represents age- and sex-appropriate weight gain.  $\Delta$ WAZ>0 indicates cCatch-up weight gain, or weight attainment above normal limits following a period of slowed development, is represented by  $\Delta$ WAZ>0. Since the World Health Organization (WHO) does not publish WAZ standards for children >10 -years of age, we used the reference from the U.S. Centers for Disease Control and Prevention (CDC) to calculate WAZ scores [26].

Definitions

 We defined microbiological confirmation as the visualization of bacilli on acid-fast smear or the isolation of *M. tuberculosis* from culture. Following Wiseman *et al.*, we classified participants into those with severe and non-severe disease [27].

At the end of therapy, participants were <u>classified intoassigned</u> treatment outcomes by a clinician according to NTP guidelines:  $cure_{a;}$  treatment completion<sub>a;</sub> loss to follow- $up_{a;}$  transfer<sub>a;</sub> treatment failure<sub>a;</sub> or death [21, 22]. For this analysis, we grouped cure and treatment completion together under treatment success. In addition, wWe classified participants as having an unknown outcome if no treatment outcome was recorded in the medical record; tThis category is distinct\_differs from loss to follow-up, which is an outcome recorded in the patient's chart.

Health center staff measured each TB patientparticipant's weight at the end of every treatment month, defined as 25 days of supervised doses. We defined baseline weights as those taken  $\leq$ *within*-30 days of the initiation of the first-line regimen for Group 1 and a second-line regimen for Group 2; end-of-therapy weights as those taken in the final treatment month of successful therapy (month 6 for most Group 1 subjects, month 12 or later for Group 2 subjects); and interim weights as all other weights taken during therapy. A child's baseline WAZ (WAZ<sub>baseline</sub>) and WAZ<sub>final</sub> refers to the z-scores calculated from the baseline and end-of-therapy weights.

### Clinical care

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Group 1 received first-line regimens—isoniazid, rifampin, ethambutol, and pyrazinamide daily for the first two months, followed by isoniazid and rifampin thrice weekly for  $\geq$ 4 at least four more months. Group 2 received second-line regimens that were tailored to the DST result of the participant or his/her presumed source case, contained  $\geq$ 4 at least four drugs to which the *M*. *tuberculosis* strain was susceptible, and lasted  $\geq$ at least 12 months. Health workers administered directly observed therapy (DOT) at the health center six days a week. No supervised or selfadministered doses were given on Sundays. Subjects did not receive nutritional supplements as part of TB therapy.

#### Data collection

At the eight health centers, we collected data on children who received only first-line regimens. At NTP headquarters, we collected data on children who received any second-line regimen; this group included children who experienced treatment failure on the first-line regimen prior to being diagnosed with MDR-TB. From each subject's chart, we abstracted date of birth; sex; HIV status; anatomic site of TB disease; smear, culture, DST, and radiograph results; antibiotics received; treatment outcome; and all weights.

#### Exclusion criteria

We excluded children who lacked baseline and/or interim weights, were lost to follow-up, <u>or</u> had an unknown outcome, <u>or whose WAZ<sub>baseline</sub> was more than three SDs from the mean</u>. <u>Suspecting</u> <u>errors in data entry, we excluded children whose WAZ scores were >6 SDs from the mean</u>. Some

participants in Group 1 were classified as having failed treatment if their DST or a source case's DST demonstrated resistance to isoniazid and rifampin. We included these children in Group 1 only if they had documentation of smear or culture positivity and/or radiological worsening after  $\geq 2$  at least two-months of treatment, despite documented good adherence [3]. If they lacked this documentation, we excluded them. Participants who had received first-line therapy in the six months preceding their second-line regimen were excluded from Group 2.

#### Analysis

We used SAS version 9.4 (SAS Institute, Cary, NC) to perform analyses. To check for selection and loss-to-follow-up biases, we used the chi-square or Fisher's exact test to compare characteristics between included and excluded subjects.

To examine the relationship between weight change and treatment outcome, we used  $\Delta$ WAZ as the measurement because it accounts for age- and sex-specific differences in normal childhood growth. We examined  $\Delta$ WAZ through treatment month 5 for Group 1 since the standard firstline regimen lasts six months, and through month 8 for Group 2 since eight months typically is the maximum duration of the intensive phase of MDR-TB therapy. We calculated  $\Delta$ WAZ using the following formula:  $\Delta$ WAZ<sub>x</sub> = WAZ<sub>x</sub> - WAZ<sub>baseline</sub>, where x represents the month of interest.

We performed two separate analyses to address the first aim of this study. First, using logistic regression, we evaluated the associations of the following independent variables and treatment outcome (success vs. failure/death) for Groups 1 and 2 separately:  $WAZ_{baseline}$ ,  $\Delta WAZ_2$ ,  $\Delta WAZ_3$ ,

 $\Delta$ WAZ<sub>4</sub>,  $\Delta$ WAZ<sub>5</sub>, age (0-4 vs. 5-9 vs.  $\geq$ 10), sex, microbiologic confirmation, and disease severity. For Group 2, we assessed additional independent variables:  $\Delta$ WAZ<sub>6</sub>,  $\Delta$ WAZ<sub>7</sub>,  $\Delta$ WAZ<sub>8</sub>, and resistance pattern (MDR vs. XDR).

Second, we used generalized estimating equation (GEE) models using the first-order autoregressive correlation structure to detect differences in  $\Delta$ WAZs between successfully treated children vs. those with failed treatment/died. To determine which covariates (age, sex, microbiologic confirmation, disease severity, resistance pattern) to include in the GEE models, we performed stepwise selection and maintained covariates with *p*-value <0.05.

To address the second aim of this study, we plotted receiver operating characteristic (ROC) curves and calculated area under the curve (AUC), sensitivity, specificity, positive predictive value (PPV), and negative predictive values (NPV) to determine the accuracy of  $\Delta$ WAZ at our time points of interest for predicting treatment failure/death.

Finally, we assessed catch-up weight gain for successfully treated subjects by comparing  $\Delta$ WAZ between WAZ<sub>baseline</sub> and WAZ<sub>final</sub>. We used the Student's t-test and analysis of variance with Bonferroni post-hoc tests to compare  $\Delta$ WAZ between subgroups stratified by age, sex, microbiologic confirmation, and disease severity.

Ethics

The Institutional Review Boards (IRBs) of Peru's National Institute of Health, Harvard Medical School, and Baylor College of Medicine approved this study and waived informed consent.

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RESULTS

We excluded 71 of 171 subjects from Group 1 and 138 of 232 subjects from Group 2 (Figure 1).

Table 1 compares included and excluded subjectsparticipants. Median ages for Groups 1 and 2

were 14.4 (interquartile range [IQR] 12.3-16.65) years and 9.5 (IQR 3.49-13.73) years,

respectively. In Group 1, the 74 successfully treated subjects had a median treatment durations of were 6.2-4 (SD: 1.8IQR 6.0-6.9) months in the 74 successfully treated subjects ; the 26 who died or failed therapy had a mean treatment duration of and 4.82 (SD: 1.5IQR 3.6-5.7) months in the 26 subjects in whom therapy failed. No participants in Group 1 died. In Group 2, median treatment durations were the 83 successfully treated subjects had a mean treatment duration of 20.05 (SD: 9.0IQR 18.0-23.8) months for the 83 successfully treated subjects and ; the 11 who died or failed therapy had a mean treatment duration of 12.714.1 (SD: 8.3IQR 5.9-21.9) months for the 10 subjects in whom therapy failed. The one subject in Group 2 who died received treatment for 7.8 months.

Association between  $\Delta WAZ$  and treatment outcome

In logistic regression, female sex and lower  $\Delta WAZ_3$ ,  $\Delta WAZ_4$ , and  $\Delta WAZ_5$  were associated with treatment failure/death in Group 1. Lower  $\Delta WAZ_7$  was associated with treatment failure/death in Group 2 (Table 2).

In the GEE model for Group 1, after adjusting for WAZ<sub>baseline</sub> and age group, children in whom treatment failed or who died had significantly lower  $\Delta$ WAZ<sub>2</sub>,  $\Delta$ WAZ<sub>3</sub>,  $\Delta$ WAZ<sub>4</sub>, and  $\Delta$ WAZ<sub>5</sub>

values than successfully treated children. For Group 2, the GEE model was not adjusted because no covariates were significant in stepwise selection; children in whom treatment failed or who died had lower  $\Delta WAZ_6$ ,  $\Delta WAZ_7$ , and  $\Delta WAZ_8$  values compared to successfully treated children (Figure 2a). When we repeated the GEE model for participants with microbiologically confirmed TB, the findings for Group 1 did not change. In Group 2, children in whom treatment failed or who died had lower  $\Delta WAZ_7$  and  $\Delta WAZ_8$  values compared to successfully treated children (Figure 2b).

Accuracy of  $\Delta WAZ$  as a predictor of treatment outcome

The AUC of  $\Delta$ WAZ as a predictor of treatment failure/death increased as therapy progressed for Group 1. The selected cut-off values had sensitivities of 60-90% and specificities of 60-85% at the end of months 2-5. For Group 2, the AUC of  $\Delta$ WAZ as a predictor of treatment failure/death was highest at the end of months 7-8. Before month 7, the selected  $\Delta$ WAZ cut-off values had <43% sensitivity for predicting treatment failure/death (Figure 3, Table 3).

Catch-up weight gain among successfully treated children

In Groups 1 and 2, children with severe TB started therapy with lower WAZ<sub>baseline</sub> compared to children with non-severe TB: -0.97 for severe disease vs. -0.37 for non-severe disease (p=0.02) in Group 1, and -0.82 for severe disease vs. -0.31 for non-severe disease (p=0.04) in Group 2 (Supplementary Table 1). At the end of successful therapy, there were no significant differences in WAZ<sub>final</sub> for children with severe vs. non-severe TB in Group 1 or 2 (Supplementary Table 2).

There were no significant differences in  $WAZ_{baseline}$  or  $WAZ_{final}$  between subjects stratified by sex, age group, microbiologic confirmation, or resistance pattern in Group 1 or 2.

All successfully treated children achieved catch-up weight gain, except for Group 2 subjects with unknown microbiologic confirmation-status. In Group 1, children with microbiologically confirmed or severe TB experienced more weight gain than those with unconfirmed or non-severe TB, respectively (Figure 4).

# DISCUSSION

In this study, we found an association between weight gain during childhood TB therapy and treatment success. that cChildren who ultimately have had treatment successful outcomes had different weight trajectories compared to those who experienced treatment failure or death. We further found thatFurthermore,  $\Delta WAZ$  early in therapy may serve as a moderately sensitive predictor of treatment failure/death for children on first-line TB regimens.  $\Delta WAZ$  lacks sensitivity for predicting unsuccessful MDR-TB treatment, but its high specificity means that clinicians should strongly consider regimen modification for children with  $\Delta WAZ$  values below the cut-off value.

For children  $\leq 10$  years, clinicians can calculate WAZ using WHO Anthro, which is available for free download onto computers and mobile devices (www.who.int/childgrowth/software/en/). PediTools (https://peditools.org) and the Children's Hospital of Philadelphia (https://zscore.research.chop.edu) provide links to CDC's WAZ calculator for children ages 2-20. Manual calculation of  $\Delta$ WAZ is likely to be inaccurate, since growth charts do not capture sufficiently small units of age and weight. Therefore, clinicians in the resource-constrained settings, where these technologies may be unavailable, may have difficulty using this tool.

The accuracy of  $\Delta$ WAZ for predicting pediatric TB treatment outcome should be evaluated in other groups of children. In clinical trials and prospective cohorts of children with TB, it would be useful to collect weights, heights, and mid-upper arm circumference (MUAC) of children and evaluate  $\Delta$ WAZ and other nutritional measures as predictors of treatment outcome. If validated,

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 $\Delta$ WAZ could serve as a red flag to prompt further evaluation of children on TB therapy who may need regimen modification. From a research perspective,  $\Delta$ WAZ could serve as a surrogate endpoint for clinical trials, reducing cost and duration and facilitating the use of novel trial designs, such as the multi-stage, multi-arm trial.

In this cohort, we found that all groups of children with successfully treated TB disease achieved catch-up weight gain. In Group 1, children with microbiologically confirmed or severe disease had more weight gain than children with non-microbiologically confirmed or non-severe disease, respectively. This finding suggests that successful treatment of TB disease leads to weight recovery; this improvement is most pronounced for the sickest children, who likely begin treatment with a greater weight deficit.

Previous work has shown that children gain weight on TB therapy [14, 15], but our study is the first to demonstrate different weight trends in children who are successfully treated and those who experienced treatment failure or death. This finding is consistent with reported associations between weight gain and treatment outcome in adults on TB treatment [6, 8, 9]. Studies of HIV-infected children on antiretroviral therapy also have shown associations between good treatment response and catch-up growth [28, 29]. Our study is the first to quantify the degree of weight gain in early TB therapy that predicts ultimate outcome.

This analysis had several limitations. The data, which were obtained under programmatic conditions, had missing values. Twenty-seven percent of children in Group 1 and 16% of children in Group 2 were excluded from the analysis due to missing weights; the exclusion of

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these subjects may have led to a selection bias. In Group 2, there were higher proportions of microbiologically confirmed and severe disease among excluded subjects compared to included subjects. This exclusion may have minimized the difference in WAZ trajectories between successfully treated children and those who experienced treatment failure or death, considering our finding that children in Group 1 with microbiologically confirmed or severe TB gained more weight than those with non-confirmed or non-severe TB. Second, because few subjects had documented heights and no subjects had MUACs, we could not undertake a more complete evaluation of nutritional recovery, including weight adjusted for height and linear growth. This limitation may be problematic for adolescents, who undergo pubertal growth spurts at different times. Adolescents with earlier-than-average puberty will have relatively higher WAZ, while body mass index-for-age z-scores may track more consistently. Despite the fact that our sample consisted mostly of adolescents, we were still able to demonstrate a clear difference in the weight trajectories of successfully treated subjects vs. those who experienced treatment failure or death.

An important limitation specific to Group 2 was the heterogeneity of second-line regimens, which were individualized to DST profiles. Additionally, some regimens were and adjusted modified during therapy because of new DST results, adverse drug events, or stock-outs. Regimen heterogeneity may have impacted our results since different medications have distinct effects on appetite; moreover, ethionamide, a second-line drug used in Peru, may affect thyroid function. Nonetheless, we can still conclude from this analysis that children successfully treated for MDR-TB achieve catch-up weight gain and achieved more early weight gain compared to those who experienced treatment failure or death.

This analysis also had strengths. The enrollment of consecutive children treated for TB disease reduces the potential for selection bias. These data were collected as part of routine care delivered at public health centers; as a result, the data reflect clinical practice and treatment response in a "real-world" setting. Our study also evaluates weight change in children being treated for drug-susceptible and MDR-TB; this analysis has not been undertaken previously in adults or children.

In conclusion, our study establishes the potential for using weight change early in therapy to predict childhood TB treatment outcome and provides reassurance that successfully treated children achieve catch-up weight gain. If validated in other cohorts, the use of weight change to predict treatment outcome can lead to better outcomes for children with TB disease and more efficient clinical trial design in childhood TB therapy.

# AUTHOR CONTRIBUTIONS

SSC, JFF, JAS, and MCB conceptualized the study. SSC, EW, HDC, and LL collected and managed the data. SP performed the analyses. SSC, SP, JFF, ATC, JAS, and MCB interpreted the results of the analyses. SSC, SP, and EW drafted the manuscript, tables, and figures. All authors revised and approved the final version of this work for publication.

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		Gr	Group 1 (n=171)			Group 2 (n=232)		
		Included	Excluded	p-value	Included	Excluded	p-value	
		(n=100)	(n=71)		(n=94)	(n=138)		
Age				0.55			0.025	
٠	0-4	6 (6.0%)	9 (12.7%)		35 (37.2%)	29 (21.0%)		
•	5-9	6 (6.0%)	5 (7.0%)		12 (12.8%)	23 (16.7%)		
•	≥10	88 (88.0%)	57 (80.3%)		47 (50.0%)	86 (62.3%)		
Sex				0.32			0.41	
•	Female	43 (43.0%)	36 (50.7%)		48 (51.1%)	78 (56.5%)		
٠	Male	57 (57.0%)	35 (49.3%)		46 (48.9%)	60 (43.5%)		
Microb	oiologic confirmatio	n		0.78			0.011	
٠	Confirmed	59 (59.0%)	45 (63.4%)		54 (57.4%)	99 (71.7%)		
•	Unconfirmed	13 (13.0%)	7 (9.9%)		28 (29.8%)	19 (13.8%)		
٠	Unknown	28 (28.0%)	19 (26.8%)		12 (12.8%)	20 (14.5%)		
Disease	e severity			0.07			0.021	
•	Severe	24 (24.0%)	26 (36.6%)		25 (26.6%)	57 (41.3%)		
٠	Non-severe	76 (76.0%)	45 (63.4%)		69 (73.4%)	81 (58.7%)		
Resista	nce pattern						0.06	
٠	MDR		0		88 (93.6%)	136 (98.6%)		
•	XDR				6 (6.4%)	2 (1.4%)		

Data are presented as n (% of total).

Abbreviations: MDR, multi-drug resistant; WAZ, weight-for-age z-score; XDR, extensively-drug resistant.

	(	Group 1 (n=100	))		Group 2 (n=94)			
	Treatment	Treatment	OR	Treatment	Treatment	OR		
	success	failure/death	(95% CI)	success	failure/death	(95% CI)		
Mean WAZ <sub>baseline</sub>	-0.44 (SD:	-0.72 (SD:	0.79 (0.52-	-0.44 (SD:	-0.56 (SD:	0.90 (0.50		
	1.18)	0.83)	1.19)	0.98)	1.76)	1.60)		
Mean ∆WAZ <sub>2</sub>	0.22 (SD:	0.01 (SD:	0.35 (0.11-	0.18 (SD:	-0.08 (SD:	0.19 (0.01		
	0.47)	0.37)	1.12)	0.49)	0.26)	2.83)		
Mean ∆WAZ <sub>3</sub>	0.41 (SD:	-0.08 (SD:	0.25 (0.08-	0.18 (SD:	0.00 (SD:	0.46 (0.10		
	0.88)	0.47)	$0.78)^{*}$	0.52)	0.27)	2.18)		
Mean ∆WAZ₄	0.32 (SD:	-0.07 (SD:	0.33 (0.11-	0.25 (SD:	0.10 (SD:	0.62 (0.18		
	0.58)	0.41)	$1.00)^{*}$	0.60)	0.34)	2.17)		
Mean ∆WAZ <sub>5</sub>	0.40 (SD:	-0.20 (SD:	0.02 (0.00-	0.2 <u>8</u> 2 (SD:	-0.21 (SD:	0. <del>58-<u>23</u></del>		
	0.41)	0.53)	0.29)**	0. <u>55</u> 7 <del>6</del> )	0.79)	(0. <del>27<u>06</u>-</del>		
						<u>1.240.90</u> )		
Mean ∆WAZ <sub>6</sub>				0.20 (SD:	-0.12 (SD:	0.45 (0.13		
				0.65)	0.45)	1.54)		
Mean <b>AWAZ</b> 7				0.25 (SD:	-0.25 (SD:	0.22 (0.05		
				0.59)	0.48)	0.92)*		
Mean ∆WAZ <sub>8</sub>				0.30 (SD:	-0.27 (SD:	0.20 (0.04		
				0.63)	0.55)	1.05)		
Age								
• 0-4 years	6 (100%)	0 (0%)	Reference	28 (96.6%)	1 (3.5%)	Reference		
• 5-9 years	6 (100%)	0 (0%)		16 (94.1 %)	1 (5.9%)	1.75 (0.10		
						29.92)		
• $\geq 10$ years	62 (70.5%)	26 (29.5%)	4	39 (81.3%)	9 (18.8%)	6.46 (0.77		
						53.95)		
Sex								
• Male	47 (82.5%)	10 (17.5%)	Reference	41 (89.1%)	5 (10.9%)	Reference		
• Female	27 (62.8%)	16 (37.2%)	2.79 (1.11-	42 (87.5%)	6 (12.5%)	1.17 (0.33		
			$7.00)^{*}$			4.14)		
Microbiologic confi	irmation							
<ul> <li>Unconfirmed</li> </ul>	13 (100%)	0 (0%)	Reference	28 (100%)	0 (0%)	Reference		
<ul> <li>Confirmed</li> </ul>	35 (59.3%)	24 (40.7%)		44 (81.5%)	10 (18.5%)			
• Unknown	26 (92.9%)	2 (7.1%)		11 (91.7%)	1 (8.3%)			
Disease severity								
Non-severe	55 (72.4%)	21 (27.6%)	Reference	61 (89.7%)	7 (10.3%)	Reference		
• Severe	19 (79.2%)	5 (20.8%)	0.69 (0.23-	22 (84.6%)	4 (15.4%)	1.58 (0.42		
	. ,	. /	2.08)		. /	5.94)		
Resistance pattern								
• MDR				77 (87.5%)	11 (12.5%)	Reference		
• XDR				6 (100%)	0 (0%)			

Abbreviations: MDR, multidrug-resistant; N/A, not applicable; SD, standard deviation; WAZ, weight-for-age z-score; XDR, extensively-drug resistant.

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to predict treatment	t failure/death					
	AUC	∆WAZ	Sensitivity	Specificity (%)	PPV (%)	NPV (%)
Group 1		cut on	(/*)	(, , ,		
• $\Delta WAZ_2$	0.68	0.142	60.0	73.1	85.7	40.4
• $\Delta WAZ_3$	0.73	0.029	81.5	60.0	86.9	50.0
• $\Delta WAZ_4$	0.78	0.153	72.3	84.6	95.9	37.9
• $\Delta WAZ_5$	0.83	-0.024	89.8	80.0	96.4	57.1
Group 2						
• $\Delta WAZ_2$	0.64	0.151	42.2	100	100	11.9
• $\Delta WAZ_3$	0.61	0.342	28.6	100	100	15.3
• $\Delta WAZ_4$	0.57	0.303	42.9	88.9	96.8	16.7
• $\Delta WAZ_5$	0. <del>68<u>69</u></del>	0.425	40. <mark>0<u>6</u></mark>	100	100	16. <del>0</del> 3
• $\Delta WAZ_6$	0.67	0.320	40.3	100	100	14.9
• $\Delta WAZ_7$	0.75	0.145	62.3	85.7	97.7	18.8
<ul> <li>ΔWAZ<sub>8</sub></li> </ul>	0.76	0.129	64.6	80.0	97.7	14.8

 Table 3: Receiver operating characteristic curves characteristics for using change in weight-for-age z-score to predict treatment failure/death

Abbreviations: AUC, area under the curve; NPV, negative predictive value; PPV, positive predictive value; WAZ, weight-for-age z-score;  $\Delta$ WAZ<sub>x</sub>, change in WAZ between baseline and the end of treatment month x.

			Group 1			Group 2	
		n	Mean (SD)	p-value	n	Mean (SD)	p-value
All	participants	100	-0.51 (1.10)		94	-0.45 (1.09)	
Ag	e						
•	0-4 years	6	-0.69 (0.70)	0.74	29	-0.44 (0.80)	0.09
•	5-9 years	6	-0.21 (1.05)	_	17	0.04 (1.45)	
•	≥10 years	88	-0.52 (1.13)	_	48	-0.64 (1.06)	
Sez	x						
•	Male	57	-0.49 (1.21)	0.76	46	-0.38 (1.11)	0.51
•	Female	43	-0.55 (0.95)	_	48	-0.53 (1.07)	
Mi	crobiologic confirmation						
•	Unconfirmed	13	-0.19 (1.47)	0.14	28	-0.35 (0.91)	0.12
•	Confirmed	59	-0.69 (1.07)	_	54	-0.62 (1.08)	
•	Unknown	28	-0.29 (0.93)	_	12	0.05 (1.39)	
Dis	sease severity						
•	Non-severe	76	-0.37 (1.02)	0.019	68	-0.31 (1.13)	0.044
•	Severe	24	-0.97 (1.25)	_	26	-0.82 (0.89)	
Re	sistance pattern		7				
•	MDR		A		88	-0.45 (1.07)	0.83
•	XDR				6	-0.54 (1.46)	
Tr	eatment outcome						
•	Treatment success	74	-0.44 (1.18)	0.19	83	-0.44 (0.98)	0.82
-	Treatment failure or death	26	-0.72 (0.83)	-	11	-0.56 (1.76)	

Abbreviations: IQR, interquartile range; MDR, multi-drug resistant; WAZ, weight-for-age z-score; XDR, extensively-drug resistant.

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	Supplement	aly radic 2. $WAL_{i}$	final of Study	Subject	13	
		Group 1				
	n	Mean (SD)	p-value	n	Mean (SD)	p-value
All participants	100	-0.08 (1.05)		94	-0.04 (1.01)	
Age						
• 0-4 years	6	-0.26 (0.87)	0.78	29	0.34 (0.76)	0.10
• 5-9 years	6	0.18 (0.94)		17	0.06 (1.35)	
• ≥10 years	88	-0.10 (1.08)		48	-0.28 (0.95)	
Sex						
• Male	57	-0.05 (1.13)	0.72	46	0.06 (1.09)	0.41
• Female	43	-0.15 (0.88)		48	-0.15 (0.93)	
Microbiologic confirmation	on					
• Unconfirmed	13	-0.03 (1.12)	0.63	28	0.19 (0.74)	0.31
Confirmed	59	0.02 (1.13)		54	-0.21 (1.03)	
• Unknown	28	-0.26 (0.88)		12	0.13 (1.55)	
Disease severity						
• Non-severe	76	0.05 (1.03)	0.10	68	0.10 (1.00)	0.08
• Severe	24	-0.45 (1.05)		26	-0.39 (0.98)	
Resistance pattern						
• MDR				88	-0.05 (1.03)	0.71
• XDR				6	0.17 (0.78)	

Abbreviations: IQR, interquartile range; MDR, multi-drug resistant; WAZ, weight-for-age z-score; XDR, extensively-drug resistant.

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3	1	FIGURE LEGENDS
4	2	
5	$\frac{2}{3}$	Figure 1
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7	- - -	Abbraviations: MDP TP multidrug registent tuberculogis: TP tuberculogis: WA 7 baseline, weight for age 7 secre
8	5	Addreviations. MDR-1D, multiding-resistant tuderculosis, 1D, tuderculosis, WAZDasenne, weight-101-age Z-score
9	07	within 50 days of the start of tuberculosis therapy. The start of tuberculosis therapy fefers to the start of a first-fine
10	/	regimen for Group 1 and a second-line regimen for Group 2.
11	0	
12	10	Figure 2
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13		———— Achieved treatment success
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15	11	Failed treatment of died
16	12	Abbraviations: AWAZ delta (change in) weight for age z score
17	12	Abbreviations. A w AZ, detta (change in) weight-toi-age 2-score.
18	13	*p-value < 0.05.
19	14	**p-value <0.01.
20	15	***p-value <0.001.
21	16	Values represent least square mean $\pm$ 95% confidence interval for each time point using generalized estimating
22	17	equation models. After stenwise selection we adjusted for WAZ
23	17	equation models. After stepwise selection, we adjusted for wAZ <sub>baseline</sub> and microbiologic committation in Group 1,
23	18	but did not adjust for any covariates in Group 2.
24	19	The numbers of participants in Group 1 contributing to the <u>overall</u> analysis were as follows: 96 for month 2, 85 for
25	20	month 3, 78 for month 4, and 69 for month 5. In Group 2, the numbers were 69 for month 2, 79 for month 3, 79 for
20	21	month 4, 78 for month 5, 74 for month 6, 76 for month 7, and 70 for month 8. The numbers of participants in Group
27	22	1 contributing to the sensitivity analysis were as follows: 58 for month 2, 52 for month 3, 46 for month 4, and 39 for
28	22	<u>i contributing to the sensitivity analysis were as follows: 56 for month 2, 32 for month 5, 40 for month 4, 42 for worth 5, 41 for</u>
29	25	month 5. In Group 2, the numbers were 39 for month 2, 46 for month 3, 43 for month 4, 43 for month 5, 41 for
30	24	month 6, 43 for month 7, and 38 for month 8.
31	25	
32	26	Figure 3
33	27	
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35		Group 1 Group 2
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37		End of treatment month 2 End of treatment month 5
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<u>/0</u>		End of treatment month 4 End of treatment month 7
40		End of treatment month 5 End of treatment month 8
41	20	
42	28	
43	29	See Table 3 for area under the curve (AUC) values.
44	30	
45	31	Figure 4
46	32	
47	33	Abbreviations: AWAZ delta (change in) weight-for-age z-score
48	31	This analysis avaluates study subjects without decumented and of therapy weights. Number of subjects in each
49	25	This analysis excludes study subjects without documented end-of-therapy weights. Number of subjects in each
50	30	subgroup appears in parentheses below subgroup name. The only significant differences among subgroups are
51	36	indicated in the figure.
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#### Figure 1: Included and excluded subjects

# Group 1



# Group 1



Group 2

# Group 2



# 

# 1 Figure 2<u>a</u>: Adjusted ΔWAZ for children who were successfully treated vs. children who failed treatment/died

# 2 (microbiologically confirmed and clinically diagnosed cases)









treatment failure/death





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Changes in weight-for-age z-score predict effectiveness of childhood tuberculosis therapy Silvia S. Chiang,<sup>1,2</sup> Sangshin Park,<sup>1,2</sup> Emily I. White,<sup>1</sup> Jennifer F. Friedman,<sup>1,2</sup> Andrea T. Cruz,<sup>3</sup> Hernán Del Castillo,<sup>4</sup> Leonid Lecca,<sup>5,6</sup> Mercedes C. Becerra,<sup>5,6\*</sup> James A. Seddon<sup>7\*</sup> 1. Department of Pediatrics, Alpert Medical School of Brown University, Providence, U.S.A. 2. Center for International Health Research, Rhode Island Hospital, Providence, U.S.A. 3. Department of Pediatrics, Baylor College of Medicine, Houston, U.S.A. 4. Instituto Nacional de Salud del Niño, Lima, Peru 5. Socios En Salud Sucursal Perú, Lima, Peru 6. Department of Global Health and Social Medicine, Harvard Medical School, Boston, U.S.A. 7. Centre for International Child Health, Imperial College London, London, U.K. \*These authors contributed equally to this work. Key words: multidrug-resistant tuberculosis, treatment outcome, nutritional status, growth and development, Peru Running title: Early weight change predicts TB outcome Corresponding author: Dr. Silvia S. Chiang, Center for International Health Research, 55 Claverick St., Ste. 101, Providence, RI 02906, U.S.A.

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Summary: In a cohort of children treated for tuberculosis disease, the degree of weight change early in therapy predicted final treatment outcome. This finding can lead to improved management and clinical trial design in childhood tuberculosis therapy.

<text>

#### 

# ABSTRACT

<u>Background</u>: International guidelines recommend monitoring weight as an indicator of therapeutic response in childhood tuberculosis (TB) disease. This recommendation is based on observations in adults. In this study, we evaluated the association between weight change and treatment outcome, the accuracy of using weight change to predict regimen efficacy, and whether successfully treated children achieve catch-up weight gain.

<u>Methods</u>: We enrolled children treated for drug-susceptible TB disease (Group 1) and multidrugresistant TB disease (Group 2) in Peru. We calculated the change in weight-for-age z-score ( $\Delta$ WAZ) between baseline and the end of treatment months 2-5 for Group 1, and between baseline and months 2-8 for Group 2. We used logistic regression and generalized estimating equation (GEE) models to evaluate the relationship between  $\Delta$ WAZ and outcome. We plotted receiver operating characteristic curves to determine the accuracy of  $\Delta$ WAZ for predicting treatment failure/death.

<u>Results</u>: Groups 1 and 2 included 100 and 94 children, respectively. In logistic regression, lower  $\Delta$ WAZ in months 3-5 and month 7 was associated with treatment failure/death in Groups 1 and 2, respectively. In GEE models, children in both group who experienced treatment failure or death had lower  $\Delta$ WAZ than successfully treated children.  $\Delta$ WAZ predicted treatment failure/death with 60-90% sensitivity and 60-86% specificity in months 2-5 for Group 1 and months 7-8 for Group 2. All successfully treated children—except Group 2 subjects with unknown microbiologic confirmation status—achieved catch-up weight gain.

Conclusions: Weight change early in therapy can predict outcome of childhood TB treatment.

tor Review Only

#### INTRODUCTION

Each year, approximately one million children develop tuberculosis (TB) disease [1, 2]. Failure to thrive—i.e., weight loss or lack of expected weight gain—is a complication of TB disease [3]. International guidelines recommend using weight gain as an indicator of good therapeutic response in childhood TB [3-5]. This recommendation is based on expert opinion. While studies have shown associations between weight gain and good therapeutic response in adults with TB [6-13], few studies have described weight trends in children on TB treatment [14, 15].

From a clinical perspective, the early identification of children who are failing therapy allows for timely intervention, such as adherence support or regimen modification. From a research perspective, a well-validated surrogate endpoint would facilitate clinical trials. For adults with TB disease, sputum smear and/or culture conversion are used to evaluate regimen efficacy [16]. However, <30% of children have microbiologically confirmed disease [17]. Resolution of radiographic abnormalities and clinical symptoms are suboptimal markers of treatment response: radiographic abnormalities lag behind clinical response, and symptom reporting is subjective. Therefore, there is a need to evaluate other markers—like weight change—of treatment response in children.

Another knowledge gap is whether children who are successfully treated for TB disease achieve catch-up weight gain. Characterizing weight reconstitution among TB-affected children could identify needs for nutritional supplementation and help clinicians provide anticipatory guidance to families.

We addressed these knowledge gaps using data from children treated for TB disease in Lima, Peru. Our cohort included children treated for drug-susceptible TB and children treated for multidrug-resistant (MDR)-TB, caused by strains of Mycobacterium tuberculosis resistant to at least isoniazid and rifampicin. We aimed to (1) evaluate the association between weight change early in therapy and treatment outcome; (2) determine the accuracy of weight change as a predictor of regimen efficacy; and (3) assess whether successfully treated children achieve catch-up weight gain.

# METHODS

#### Setting and study population

This study took place in Lima, Peru's capital and home to 30% of its population of 32 million [18]. Lima accounts for 54% of the 31,000 TB cases notified annually in the country [19]. Peru has a human immunodeficiency virus (HIV) prevalence of 0.3% among 15-49 year-olds [20].

Peru's National TB Program (NTP) provides free TB treatment at public health centers, where clinical charts for these patients are kept. Additionally, the NTP maintains medical records of all patients treated for MDR-TB in Peru.

We enrolled children treated for TB that was microbiologically confirmed or clinically diagnosed per NTP guidelines [21, 22]. We enrolled all children  $\leq$ 17 years of age who began treatment for drug-susceptible TB between 2012 and 2014 at eight health centers with high volumes of TB patients and longstanding collaborations with our research group. We also enrolled all children  $\leq$ 15 years of age who initiated MDR-TB therapy anywhere in the Lima metropolitan area between 2005 and 2009. We did not include 16- and 17-year-olds who initiated MDR-TB therapy because their clinical data were not available to us. We divided participants into two groups. Group 1 consisted of children initially treated for drug-susceptible TB, which included subjects who failed treatment for drug-susceptible TB disease and were subsequently diagnosed with MDR-TB. Group 2 consisted of participants with confirmed MDR-TB (i.e. they had their own *M. tuberculosis* isolate with confirmed resistance to at least isoniazid and rifampin) or

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probable MDR-TB (i.e. they lacked their own isolate but had a presumed source case with confirmed MDR-TB) [23]. Some children with MDR-TB had extensively drug-resistant (XDR)-TB, caused by MDR strains of *M. tuberculosis* with additional resistance to a fluoroquinolone and a second-line injectable agent. During the study period, DSTs were performed with the proportional method, microscopic observation drug susceptibility [24], and/or the nitrate reduction assay [25].

Weight-for-age Z-scores

Weight-for-age z-score (WAZ) compares a child's weight to children of the same age and sex from a healthy reference population. Distributions of weight-for-age measures are created for each sex over the spectrum of ages; the z-score is the number of standard deviations (SDs) from the age- and sex-appropriate mean. The change in WAZ between two time points, or  $\Delta$ WAZ, denotes weight change in comparison to the reference population.  $\Delta$ WAZ of 0 represents ageand sex-appropriate weight gain.  $\Delta$ WAZ>0 indicates catch-up weight gain, or weight attainment above normal limits following a period of slowed development. Since the World Health Organization (WHO) does not publish WAZ standards for children >10 years, we used the reference from the U.S. Centers for Disease Control and Prevention (CDC) to calculate WAZ scores [26].

#### Definitions

We defined microbiologic confirmation as the visualization of bacilli on acid-fast smear or the isolation of *M. tuberculosis* from culture. Following Wiseman *et al.*, we classified participants into those with severe and non-severe disease [27].

At the end of therapy, participants were assigned treatment outcomes by a clinician according to NTP guidelines: cure, treatment completion, loss to follow-up, transfer, treatment failure, or death [21, 22]. For this analysis, we grouped cure and treatment completion together under treatment success. We classified participants as having an unknown outcome if no treatment outcome was recorded in the medical record; this category differs from loss to follow-up, which is an outcome recorded in the patient's chart.

Health center staff measured each participant's weight at the end of every treatment month, defined as 25 days of supervised doses. We defined baseline weights as those taken  $\leq$ 30 days of the initiation of the first-line regimen for Group 1 and a second-line regimen for Group 2; endof-therapy weights as those taken in the final treatment month of successful therapy (month 6 for most Group 1 subjects, month 12 or later for Group 2 subjects); and interim weights as all other weights taken during therapy. WAZ<sub>baseline</sub> and WAZ<sub>final</sub> refer to z-scores calculated from baseline and end-of-therapy weights, respectively.

#### Clinical care

Group 1 received first-line regimens—isoniazid, rifampin, ethambutol, and pyrazinamide daily for the first two months, followed by isoniazid and rifampin thrice weekly for  $\geq$ 4 more months.

Group 2 received second-line regimens that were tailored to the DST result of the participant or his/her presumed source case, contained  $\geq$ 4 drugs to which the *M. tuberculosis* strain was susceptible, and lasted  $\geq$ 12 months. Health workers administered directly observed therapy (DOT) at the health center six days a week. No supervised or self-administered doses were given on Sundays. Subjects did not receive nutritional supplements as part of TB therapy.

# Data collection

At the eight health centers, we collected data on children who received only first-line regimens. At NTP headquarters, we collected data on children who received any second-line regimen; this group included children who experienced treatment failure on the first-line regimen prior to being diagnosed with MDR-TB. From each subject's chart, we abstracted date of birth; sex; HIV status; anatomic site of TB disease; smear, culture, DST, and radiograph results; antibiotics received; treatment outcome; and all weights.

#### Exclusion criteria

We excluded children who lacked baseline and/or interim weights, were lost to follow-up, or had an unknown outcome. Suspecting errors in data entry, we excluded children whose WAZ scores were >6 SDs from the mean. Some participants in Group 1 were classified as having failed treatment if their DST or a source case's DST demonstrated resistance to isoniazid and rifampin. We included these children in Group 1 only if they had documentation of smear or culture positivity and/or radiological worsening after  $\geq$ 2 months of treatment, despite documented good

adherence [3]. Participants who had received first-line therapy in the six months preceding their second-line regimen were excluded from Group 2.

#### Analysis

We used SAS version 9.4 (SAS Institute, Cary, NC) to perform analyses. To check for selection and loss-to-follow-up biases, we used Fisher's exact test to compare characteristics between included and excluded subjects.

To examine the relationship between weight change and treatment outcome, we used  $\Delta$ WAZ because it accounts for age- and sex-specific differences in normal childhood growth. We examined  $\Delta$ WAZ through treatment month 5 for Group 1 since the standard first-line regimen lasts six months, and through month 8 for Group 2 since eight months typically is the maximum duration of the intensive phase of MDR-TB therapy. We calculated  $\Delta$ WAZ using the following formula:  $\Delta$ WAZ<sub>x</sub> = WAZ<sub>x</sub> - WAZ<sub>baseline</sub>, where x represents the month of interest.

We performed two separate analyses to address the first aim of this study. First, using logistic regression, we evaluated the associations of the following independent variables and treatment outcome (success vs. failure/death) for Groups 1 and 2 separately:  $WAZ_{baseline}$ ,  $\Delta WAZ_2$ ,  $\Delta WAZ_2$ ,  $\Delta WAZ_3$ ,  $\Delta WAZ_4$ ,  $\Delta WAZ_5$ , age (0-4 vs. 5-9 vs.  $\geq$ 10), sex, microbiologic confirmation, and disease severity. For Group 2, we assessed additional independent variables:  $\Delta WAZ_6$ ,  $\Delta WAZ_7$ ,  $\Delta WAZ_8$ , and resistance pattern (MDR vs. XDR).

Second, we used generalized estimating equation (GEE) models using the first-order autoregressive correlation structure to detect differences in  $\Delta$ WAZs between successfully treated children vs. those with failed treatment/died. To determine which covariates (age, sex, microbiologic confirmation, disease severity, resistance pattern) to include in the GEE models, we performed stepwise selection and maintained covariates with *p*-value <0.05.

To address the second aim of this study, we plotted receiver operating characteristic (ROC) curves and calculated area under the curve (AUC), sensitivity, specificity, positive predictive value (PPV), and negative predictive values (NPV) to determine the accuracy of  $\Delta$ WAZ at our time points of interest for predicting treatment failure/death.

Finally, we assessed catch-up weight gain for successfully treated subjects by comparing  $\Delta$ WAZ between WAZ<sub>baseline</sub> and WAZ<sub>final</sub>. We used the Student's t-test and analysis of variance with Bonferroni post-hoc tests to compare  $\Delta$ WAZ between subgroups stratified by age, sex, microbiologic confirmation, and disease severity.

#### **Ethics**

The Institutional Review Boards (IRBs) of Peru's National Institute of Health, Harvard Medical School, and Baylor College of Medicine approved this study and waived informed consent.

RESULTS

#### 

# We excluded 71 of 171 subjects from Group 1 and 138 of 232 subjects from Group 2 (Figure 1). Table 1 compares included and excluded participants. Median ages for Groups 1 and 2 were 14.4 (interquartile range [IQR] 12.3-16.65) years and 9.5 (IQR 3.49-13.73) years, respectively. In Group 1, median treatment durations were 6.4 (IQR 6.0-6.9) months in the 74 successfully treated subjects and 4.8 (IQR 3.6-5.7) months in the 26 subjects in whom therapy failed. No participants in Group 1 died. In Group 2, median treatment durations were 20.0 (IQR 18.0-23.8) months for the 83 successfully treated subjects and 14.1 (IQR 5.9-21.9) months for the 10 subjects in whom therapy failed. The one subject in Group 2 who died received treatment for 7.8 months.

Association between  $\Delta WAZ$  and treatment outcome

In logistic regression, female sex and lower  $\Delta WAZ_3$ ,  $\Delta WAZ_4$ , and  $\Delta WAZ_5$  were associated with treatment failure in Group 1. Lower  $\Delta WAZ_7$  was associated with treatment failure/death in Group 2 (Table 2).

In the GEE model for Group 1, after adjusting for WAZ<sub>baseline</sub> and age group, children in whom treatment failed had significantly lower  $\Delta$ WAZ<sub>2</sub>,  $\Delta$ WAZ<sub>3</sub>,  $\Delta$ WAZ<sub>4</sub>, and  $\Delta$ WAZ<sub>5</sub> than successfully treated children. For Group 2, the GEE model was not adjusted because no covariates were significant in stepwise selection; children in whom treatment failed or who died had lower  $\Delta$ WAZ<sub>6</sub>,  $\Delta$ WAZ<sub>7</sub>, and  $\Delta$ WAZ<sub>8</sub> compared to successfully treated children (Figure 2a).

When we repeated the GEE model for participants with microbiologically confirmed TB, the findings for Group 1 did not change. In Group 2, children in whom treatment failed or who died had lower  $\Delta WAZ_7$  and  $\Delta WAZ_8$  values compared to successfully treated children (Figure 2b).

Accuracy of  $\Delta WAZ$  as a predictor of treatment outcome

The AUC of  $\Delta$ WAZ as a predictor of treatment failure increased as therapy progressed for Group 1. The selected cut-off values had sensitivities of 60-90% and specificities of 60-85% at the end of months 2-5. For Group 2, the AUC of  $\Delta$ WAZ as a predictor of treatment failure/death was highest at the end of months 7-8. Before month 7, the selected  $\Delta$ WAZ cut-off values had <43% sensitivity for predicting treatment failure/death (Figure 3, Table 3).

#### Catch-up weight gain among successfully treated children

In Groups 1 and 2, children with severe TB started therapy with lower WAZ<sub>baseline</sub> compared to children with non-severe TB: -0.97 for severe disease vs. -0.37 for non-severe disease (p=0.02) in Group 1, and -0.82 for severe disease vs. -0.31 for non-severe disease (p=0.04) in Group 2 (Supplementary Table 1). At the end of successful therapy, there were no significant differences in WAZ<sub>final</sub> for children with severe vs. non-severe TB in Group 1 or 2 (Supplementary Table 2). There were no significant differences in WAZ<sub>baseline</sub> or WAZ<sub>final</sub> between subjects stratified by sex, age group, microbiologic confirmation, or resistance pattern in Group 1 or 2.

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All successfully treated children achieved catch-up weight gain, except for Group 2 subjects with unknown microbiologic confirmation. In Group 1, children with microbiologically confirmed or severe TB experienced more weight gain than those with unconfirmed or non-severe TB, respectively (Figure 4).

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#### DISCUSSION

In this study, we found that children who ultimately had treatment success had different weight trajectories compared to those who experienced treatment failure or death. Furthermore,  $\Delta WAZ$  early in therapy may serve as a moderately sensitive predictor of treatment failure/death for children on first-line TB regimens.  $\Delta WAZ$  lacks sensitivity for predicting unsuccessful MDR-TB treatment, but its high specificity means that clinicians should strongly consider regimen modification for children with  $\Delta WAZ$  values below the cut-off value.

For children  $\leq 10$  years, clinicians can calculate WAZ using WHO Anthro (www.who.int/childgrowth/software/en/). PediTools (https://peditools.org) and the Children's Hospital of Philadelphia (https://zscore.research.chop.edu) provide links to CDC's WAZ calculator for children ages 2-20. Manual calculation of  $\Delta$ WAZ is likely to be inaccurate, since growth charts do not capture sufficiently small units of age and weight. Therefore, clinicians in resource-constrained settings, where these technologies may be unavailable, may have difficulty using this tool.

The accuracy of  $\Delta$ WAZ for predicting pediatric TB treatment outcome should be evaluated in other groups of children. In clinical trials and prospective cohorts of children with TB, it would be useful to collect weights, heights, and mid-upper arm circumference (MUAC) of children and evaluate  $\Delta$ WAZ and other nutritional measures as predictors of treatment outcome. If validated,  $\Delta$ WAZ could serve as a red flag to prompt further evaluation of children on TB therapy who may need regimen modification. From a research perspective,  $\Delta$ WAZ could serve as a surrogate

endpoint for clinical trials, reducing cost and duration and facilitating the use of novel trial designs, such as the multi-stage, multi-arm trial.

In this cohort, we found that all groups of children with successfully treated TB disease achieved catch-up weight gain. In Group 1, children with microbiologically confirmed or severe disease had more weight gain than children with non-microbiologically confirmed or non-severe disease. This finding suggests that successful treatment of TB disease leads to weight recovery; this improvement is most pronounced for the sickest children, who likely begin treatment with a greater weight deficit.

Previous work has shown that children gain weight on TB therapy [14, 15], but our study is the first to demonstrate different weight trends in children who are successfully treated and those who experienced treatment failure or death. This finding is consistent with reported associations between weight gain and treatment outcome in adults on TB treatment [6, 8, 9]. Studies of HIV-infected children on antiretroviral therapy also have shown associations between good treatment response and catch-up growth [28, 29]. Our study is the first to quantify the degree of weight gain in early TB therapy that predicts ultimate outcome.

This analysis had several limitations. The data, which were obtained under programmatic conditions, had missing values. Twenty-seven percent of children in Group 1 and 16% of children in Group 2 were excluded from the analysis due to missing weights; the exclusion of these subjects may have led to a selection bias. In Group 2, there were higher proportions of microbiologically confirmed and severe disease among excluded subjects compared to included

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subjects. This exclusion may have minimized the difference in WAZ trajectories between successfully treated children and those who experienced treatment failure or death, considering our finding that children in Group 1 with microbiologically confirmed or severe TB gained more weight than those with non-confirmed or non-severe TB. Second, because few subjects had documented heights and no subjects had MUACs, we could not undertake a more complete evaluation of nutritional recovery, including weight adjusted for height and linear growth. This limitation may be problematic for adolescents, who undergo pubertal growth spurts at different times. Adolescents with earlier-than-average puberty will have relatively higher WAZ, while body mass index-for-age z-scores may track more consistently. Despite the fact that our sample consisted mostly of adolescents, we were still able to demonstrate a clear difference in the weight trajectories of successfully treated subjects vs. those who experienced treatment failure or death.

An important limitation specific to Group 2 was the heterogeneity of second-line regimens, which were individualized to DST profiles and adjusted during therapy because of new DST results, adverse drug events, or stock-outs. Regimen heterogeneity may have impacted our results since different medications have distinct effects on appetite; moreover, ethionamide, a second-line drug used in Peru, may affect thyroid function. Nonetheless, we can still conclude from this analysis that children successfully treated for MDR-TB achieve catch-up weight gain and achieved more early weight gain compared to those who experienced treatment failure or death.

This analysis also had strengths. The enrollment of consecutive children treated for TB disease reduces the potential for selection bias. These data were collected as part of routine care

delivered at public health centers; as a result, the data reflect clinical practice and treatment response in a "real-world" setting. Our study also evaluates weight change in children being treated for drug-susceptible and MDR-TB; this analysis has not been undertaken previously in adults or children.

In conclusion, our study establishes the potential for using weight change early in therapy to predict childhood TB treatment outcome and provides reassurance that successfully treated children achieve catch-up weight gain. If validated in other cohorts, the use of weight change to predict treatment outcome can lead to better outcomes for children with TB disease and more efficient clinical trial design in childhood TB therapy.

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# AUTHOR CONTRIBUTIONS

SSC, JFF, JAS, and MCB conceptualized the study. SSC, EW, HDC, and LL collected and managed the data. SP performed the analyses. SSC, SP, JFF, ATC, JAS, and MCB interpreted the results of the analyses. SSC, SP, and EW drafted the manuscript, tables, and figures. All authors revised and approved the final version of this work for publication.

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		Group 1 (n=171)			Gi		
		Included	Excluded	p-value	Included	Excluded	p-valu
		(n=100)	(n=71)		(n=94)	(n=138)	
Age				0.55			0.025
•	0-4	6 (6.0%)	9 (12.7%)		35 (37.2%)	29 (21.0%)	
٠	5-9	6 (6.0%)	5 (7.0%)		12 (12.8%)	23 (16.7%)	
٠	≥10	88 (88.0%)	57 (80.3%)		47 (50.0%)	86 (62.3%)	
Sex				0.32			0.41
٠	Female	43 (43.0%)	36 (50.7%)		48 (51.1%)	78 (56.5%)	
٠	Male	57 (57.0%)	35 (49.3%)		46 (48.9%)	60 (43.5%)	
Microb	iologic confirmation			0.78			0.011
٠	Confirmed	59 (59.0%)	45 (63.4%)		54 (57.4%)	99 (71.7%)	
٠	Unconfirmed	13 (13.0%)	7 (9.9%)		28 (29.8%)	19 (13.8%)	
•	Unknown	28 (28.0%)	19 (26.8%)		12 (12.8%)	20 (14.5%)	
Disease	e severity			0.07			0.021
٠	Severe	24 (24.0%)	26 (36.6%)		25 (26.6%)	57 (41.3%)	
٠	Non-severe	76 (76.0%)	45 (63.4%)		69 (73.4%)	81 (58.7%)	
Resista	nce pattern						0.06
٠	MDR		-		88 (93.6%)	136 (98.6%)	
٠	XDR				6 (6.4%)	2 (1.4%)	
Data are Abbrev	e presented as n (% of t iations: MDR, multi-dr	total). Tug resistant; WA	AZ, weight-for-a	age z-score	; XDR, extensi	vely-drug resista	int.

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			Group 1 (n=100	))		Group 2 (n=94)			
		Treatment	Treatment	OR	Treatment	Treatment	OR		
		success	failure/death	(95% CI)	success	failure/death	(95% CI)		
Me	an WAZ <sub>baseline</sub>	-0.44 (SD:	-0.72 (SD:	0.79 (0.52-	-0.44 (SD:	-0.56 (SD:	0.90 (0.50		
		1.18)	0.83)	1.19)	0.98)	1.76)	1.60)		
Me	an <b>AWAZ</b> <sub>2</sub>	0.22 (SD:	0.01 (SD:	0.35 (0.11-	0.18 (SD:	-0.08 (SD:	0.19 (0.01		
	_	0.47)	0.37)	1.12)	0.49)	0.26)	2.83)		
Me	an <b>AWAZ</b> 3	0.41 (SD:	-0.08 (SD:	0.25 (0.08-	0.18 (SD:	0.00 (SD:	0.46 (0.10		
		0.88)	0.47)	$0.78)^{*}$	0.52)	0.27)	2.18)		
Me	an ∆WAZ <sub>4</sub>	0.32 (SD:	-0.07 (SD:	0.33 (0.11-	0.25 (SD:	0.10 (SD:	0.62 (0.18		
		0.58)	0.41)	1.00)*	0.60)	0.34)	2.17)		
Me	an <b>AWAZ</b> 5	0.40 (SD:	-0.20 (SD:	0.02 (0.00-	0.28 (SD:	-0.21 (SD:	0.23 (0.06		
		0.41)	0.53)	0.29)**	0.55)	0.79)	0.90)*		
Me	an ∆WAZ <sub>6</sub>	🔨			0.20 (SD:	-0.12 (SD:	0.45 (0.13		
					0.65)	0.45)	1.54)		
Me	an ∆WAZ <sub>7</sub>	(			0.25 (SD:	-0.25 (SD:	0.22 (0.05		
					0.59)	0.48)	$0.92)^{*}$		
Me	an ∆WAZ <sub>8</sub>				0.30 (SD:	-0.27 (SD:	0.20 (0.04		
					0.63)	0.55)	1.05)		
Age	e		<b>N</b>						
•	0-4 years	6 (100%)	0 (0%)	Reference	28 (96.6%)	1 (3.5%)	Reference		
•	5-9 years	6 (100%)	0 (0%)		16 (94.1 %)	1 (5.9%)	1.75 (0.10		
							29.92)		
•	≥10 years	62 (70.5%)	26 (29.5%)		39 (81.3%)	9 (18.8%)	6.46 (0.77		
							53.95)		
Sex	[								
•	Male	47 (82.5%)	10 (17.5%)	Reference	41 (89.1%)	5 (10.9%)	Reference		
•	Female	27 (62.8%)	16 (37.2%)	2.79 (1.11-	42 (87.5%)	6 (12.5%)	1.17 (0.33		
				7.00)*			4.14)		
Mi	crobiologic conf	irmation							
•	Unconfirmed	13 (100%)	0 (0%)	Reference	28 (100%)	0 (0%)	Reference		
•	Confirmed	35 (59.3%)	24 (40.7%)		44 (81.5%)	10 (18.5%)			
•	Unknown	26 (92.9%)	2 (7.1%)		11 (91.7%)	1 (8.3%)			
Dis	ease severity								
•	Non-severe	55 (72.4%)	21 (27.6%)	Reference	61 (89.7%)	7 (10.3%)	Reference		
•	Severe	19 (79.2%)	5 (20.8%)	0.69 (0.23-	22 (84.6%)	4 (15.4%)	1.58 (0.42		
				2.08)	<pre></pre>	× · · ·	5.94)		
Res	sistance pattern			,			,		
•	MDR				77 (87.5%)	11 (12.5%)	Reference		
	VDD				6 (1000/)	0 (00/)			

\*p<0.05, \*\*p<0.01 Abbreviations: MDR, multidrug-resistant; N/A, not applicable; SD, standard deviation; WAZ, weight-for-age z-

score; XDR, extensively-drug resistant.

Table 3: Receiver operating characteristic curves characteristics for using change in weight-for-age z-score

to predict treatment fa	ilure/death					
	AUC	ΔWAZ cut-off	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)
Group 1						
• $\Delta WAZ_2$	0.68	0.142	60.0	73.1	85.7	40.4
• $\Delta WAZ_3$	0.73	0.029	81.5	60.0	86.9	50.0
• $\Delta WAZ_4$	0.78	0.153	72.3	84.6	95.9	37.9
• $\Delta WAZ_5$	0.83	-0.024	89.8	80.0	96.4	57.1
Group 2						
• $\Delta WAZ_2$	0.64	0.151	42.2	100	100	11.9
• $\Delta WAZ_3$	0.61	0.342	28.6	100	100	15.3
• $\Delta WAZ_4$	0.57	0.303	42.9	88.9	96.8	16.7
• $\Delta WAZ_5$	0.69	0.425	40.6	100	100	16.3
• $\Delta WAZ_6$	0.67	0.320	40.3	100	100	14.9
• $\Delta WAZ_7$	0.75	0.145	62.3	85.7	97.7	18.8
<ul> <li>ΔWAZ<sub>8</sub></li> </ul>	0.76	0.129	64.6	80.0	97.7	14.8

Abbreviations: AUC, area under the curve; NPV, negative predictive value; PPV, positive predictive value; WAZ, weight-for-age z-score;  $\Delta$ WAZ<sub>x</sub>, change in WAZ between baseline and the end of treatment month x.

Su	pplementar	y Table 1: WAZ <sub>b</sub>	aseline of Stud	ly Subje	cts	
		Group 1			Group 2	
	n	Mean (SD)	p-value	n	Mean (SD)	p-value
All participants	100	-0.51 (1.10)		94	-0.45 (1.09)	
Age						
• 0-4 years	6	-0.69 (0.70)	0.74	29	-0.44 (0.80)	0.09
• 5-9 years	6	-0.21 (1.05)		17	0.04 (1.45)	
• $\geq 10$ years	88	-0.52 (1.13)		48	-0.64 (1.06)	
Sex						
• Male	57	-0.49 (1.21)	0.76	46	-0.38 (1.11)	0.51
• Female	43	-0.55 (0.95)		48	-0.53 (1.07)	_
Microbiologic confirmation						
• Unconfirmed	13	-0.19 (1.47)	0.14	28	-0.35 (0.91)	0.12
Confirmed	59	-0.69 (1.07)		54	-0.62 (1.08)	_
• Unknown	28	-0.29 (0.93)		12	0.05 (1.39)	
Disease severity						
• Non-severe	76	-0.37 (1.02)	0.019	68	-0.31 (1.13)	0.044
• Severe	24	-0.97 (1.25)		26	-0.82 (0.89)	
Resistance pattern		Y				
• MDR		A		88	-0.45 (1.07)	0.83
• XDR				6	-0.54 (1.46)	
Treatment outcome						
Treatment success	74	-0.44 (1.18)	0.19	83	-0.44 (0.98)	0.82
• Treatment failure or death	26	-0.72 (0.83)		11	-0.56 (1.76)	

Abbreviations: IQR, interquartile range; MDR, multi-drug resistant; WAZ, weight-for-age z-score; XDR, extensively-drug resistant.

			Group 2					
		n	Mean (SD)	p-value	n	Mean (SD)	p-valu	
All participant	s	100	-0.08 (1.05)		94	-0.04 (1.01)		
Age								
0-4 years		6	-0.26 (0.87)	0.78	29	0.34 (0.76)	0.10	
5-9 years		6	0.18 (0.94)	_	17	0.06 (1.35)		
$\geq 10$ years		88	-0.10 (1.08)		48	-0.28 (0.95)		
Sex								
Male		57	-0.05 (1.13)	0.72	46	0.06 (1.09)	0.41	
Female		43	-0.15 (0.88)	_	48	-0.15 (0.93)		
Microbiologic (	confirmation							
Unconfirme	ed	13	-0.03 (1.12)	0.63	28	0.19 (0.74)	0.31	
Confirmed		59	0.02 (1.13)	_	54	-0.21 (1.03)		
Unknown		28	-0.26 (0.88)	_	12	0.13 (1.55)		
Disease severit	у		~					
Non-severe	;	76	0.05 (1.03)	0.10	68	0.10 (1.00)	0.08	
Severe		24	-0.45 (1.05)	_	26	-0.39 (0.98)		
Resistance patt	tern							
MDR					88	-0.05 (1.03)	0.71	
XDR				_	6	0.17 (0.78)		
Abbreviations: I extensively-drug	IQR, interquartile g resistant.	range; MDR	, multi-drug resi	stant; WAZ,	weight-f	for-age z-score; XD	DR,	
extensively-drug	g resistant.							

FIGURE LEGENDS

#### Figure 1

Abbreviations: MDR-TB, multidrug-resistant tuberculosis; TB, tuberculosis; WAZbaseline, weight-for-age z-score within 30 days of the start of tuberculosis therapy. The start of tuberculosis therapy refers to the start of a first-line regimen for Group 1 and a second-line regimen for Group 2.

#### Figure 2

----- Achieved treatment success

----- Failed treatment or died

 $\begin{array}{l} 11\\ 12 \end{array} \quad \text{Abbreviations: } \Delta \text{WAZ, delta (change in) weight-for-age z-score.} \end{array}$ 

- 13 \*p-value <0.05.
- 14 \*\*p-value <0.01.
- **\*\*\***p-value <0.001.

16 Values represent least square mean +/- 95% confidence interval for each time point using generalized estimating

equation models. After stepwise selection, we adjusted for WAZ<sub>baseline</sub> and microbiologic confirmation in Group 1,
but did not adjust for any covariates in Group 2.

The numbers of participants in Group 1 contributing to the overall analysis were as follows: 96 for month 2, 85 for month 3, 78 for month 4, and 69 for month 5. In Group 2, the numbers were 69 for month 2, 79 for month 3, 79 for month 4, 78 for month 5, 74 for month 6, 76 for month 7, and 70 for month 8. The numbers of participants in Group 1 contributing to the sensitivity analysis were as follows: 58 for month 2, 52 for month 3, 46 for month 4, and 39 for month 5. In Group 2, the numbers were 39 for month 2, 46 for month 3, 43 for month 4, 43 for month 5, 41 for month 6, 43 for month 7, and 38 for month 8.

#### Figure 3

27		
	Group 1	Group 2
	End of treatment month 2	End of treatment month 5
	End of treatment month 3	End of treatment month 6
	End of treatment month 4	End of treatment month 7
	End of treatment month 5	End of treatment month 8

See Table 3 for area under the curve (AUC) values.

#### 31 Figure 4

33 Abbreviations:  $\Delta$ WAZ, delta (change in) weight-for-age z-score.

34 This analysis excludes study subjects without documented end-of-therapy weights. Number of subjects in each 35 subgroup appears in parentheses below subgroup name. The only significant differences among subgroups are

36 indicated in the figure.

# Figure 1: Included and excluded subjects

# Group 1







2 (microbiologically confirmed and clinically diagnosed cases)



4 Figure 2b: Adjusted ΔWAZ for children who were successfully treated vs. children who failed treatment/died



Group 1









