

# Month 2 culture and duration as predictors of TB relapse

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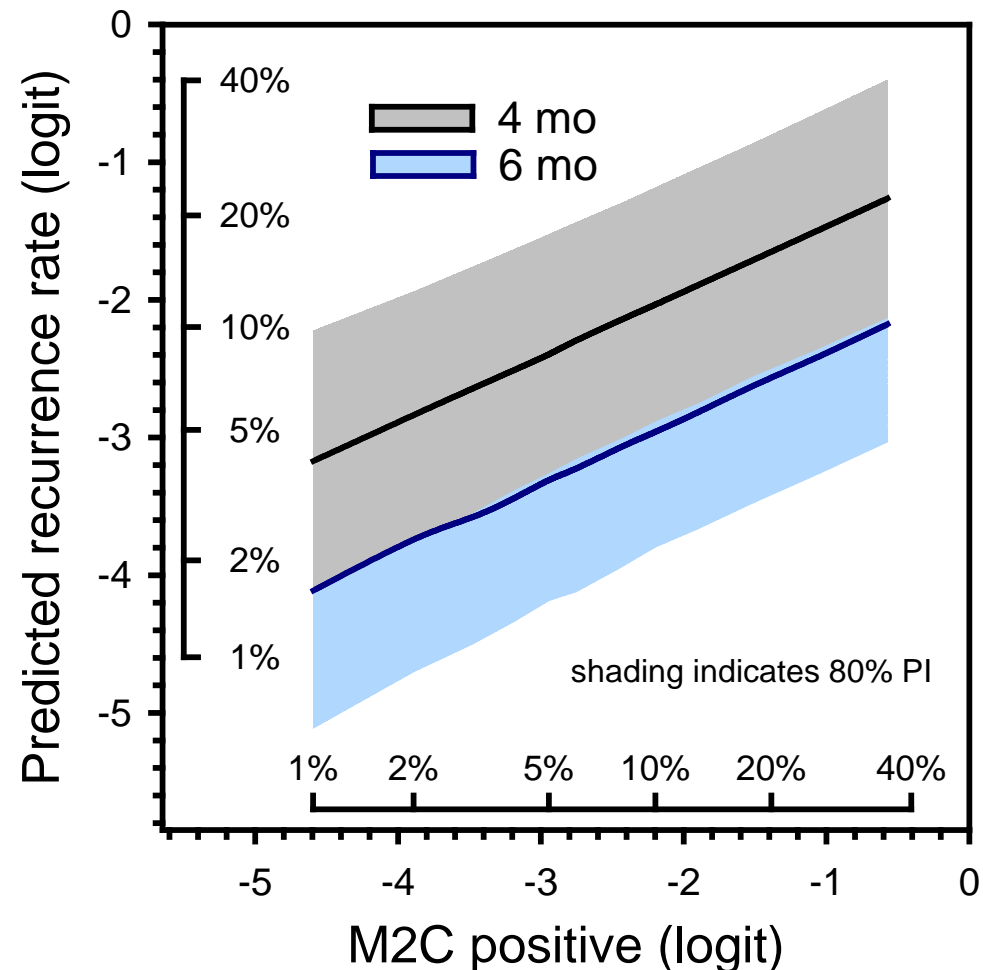
FDA TB workshop  
Jul 2017



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# M2C and duration as predictors of relapse

- A 2013 analysis of 24 trials published from 1973 to 1997 of 58 regimens studied in 7793 patients identified positive month 2 culture status using solid medium (M2C) and treatment duration as independent predictors of relapse.
- Predictions took the form:  $\text{logit}(\text{relapse}) = a + bx + cy$ , where  $x = \text{logit}(\text{M2C})$  and  $y = \log(\text{duration})$
- The model was subsequently validated using independent data from 6 studies of 12 regimens involving 3907 patients.

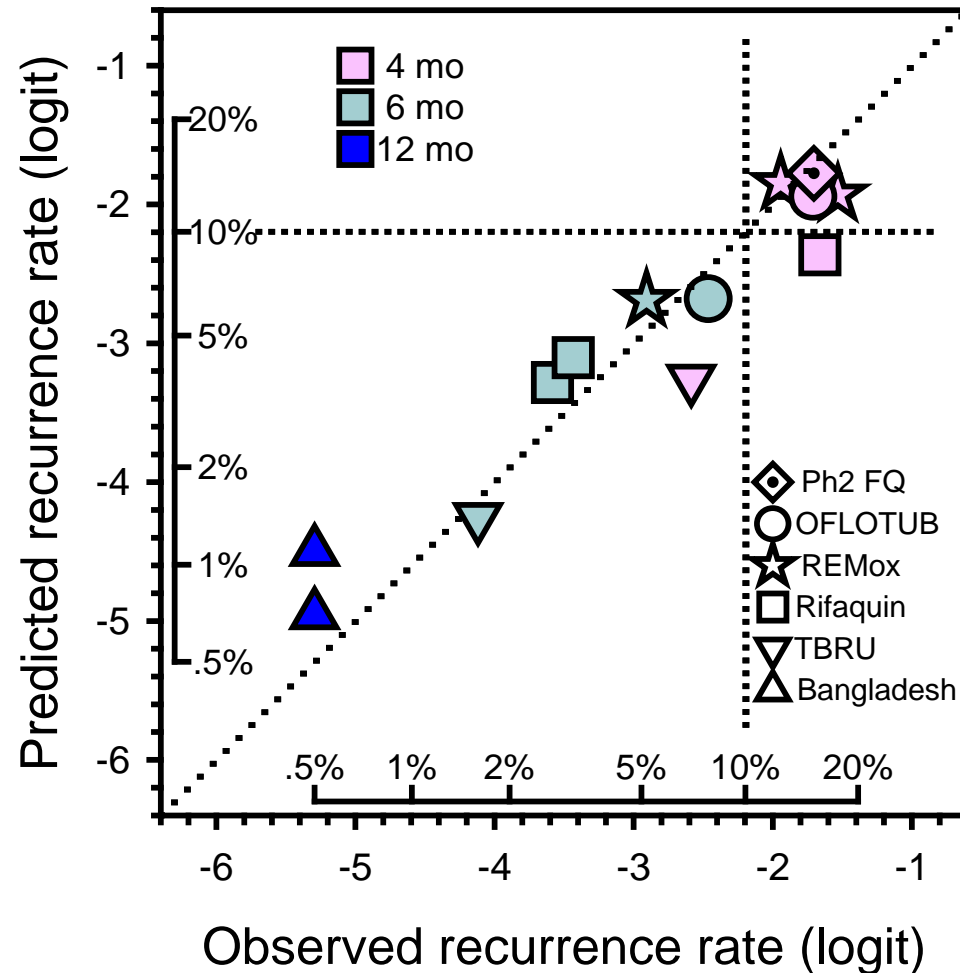


Wallis,  
*PLoS One*  
2013



# M2C and duration as predictors of relapse

- RESEARCH OBJECTIVE AND DESIGN:**  
 200 HIV-2 patients were enrolled into a vertically site-specific based cohort data from 5 of a 12 trials (10/16 female-dominant settings) with 644 patients total treated. M2C positively predicted relapse rates in all 8 arms of these trials where 6% and relapse rate in 3 trials (3388 patients) were later predicted based on M2C and post-treatment. No relapses were predicted based on M2C and duration of each arm, at  $R^2 = .86$  that predicted.
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Wallis,  
**PLoS One**  
 2015



# M2C and duration as predictors of relapse

## Database characteristics

	Total	Duration			Location	
		<i>3-5mo</i>	<i>6mo</i>	<i>7-18mo</i>	<i>Africa</i>	<i>Asia/Pacific</i>
Regimens	70	11	48	11	28	50
Subjects	11790	2824	7435	1531	7296	7814

## Model parameters

	Estimate	SE	P
Intercept	2.529	0.478	0.00002
Ln duration	-2.502	0.221	<.00001
Logit M2C+	0.440	0.010	0.00008

Online calculator at [rswallis.com](http://rswallis.com)



# M2C and duration as predictors of relapse

- The model is accurate ( $R^2 > .9$ ) and generalizable (remaining accurate under previously untested conditions):
  - FQ results were predicted w/o FQ data
  - TBRU results were predicted w/o host data
  - MDR results were predicted w/o MDR or clofazimine data
- None of the studies in the training dataset and few in the validation dataset excluded recurrence due to reinfection
  - Presumably noise would be reduced and accuracy increased had they done so
- Data are insufficient to create a similar model using time to culture conversion in liquid medium, or one including baseline parameters
  - Collecting such data should be a research priority

