

Reply to "Breakpoints and Drug Exposure Are Inevitably Closely Linked"

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We thank Alffenaar and colleagues on their very insightful comments regarding the new breakpoints for multidrugresistant tuberculosis (MDR TB) (1). We agree that the impact of the definition goes beyond statistical increases in the prevalence of MDR TB and has a direct impact on who should be treated with first-line regimens. Revision of breakpoints implies changing the regimens on which patients are started. Patients may be started on less effective second-line drugs with more toxicity. However, a preferred solution is exactly what Alffenaar and colleagues suggest, i.e., to use pharmacokinetics/ pharmacodynamics (PK/PD) to define the drug doses necessary to overcome the high MICs.

In the early PK/PD work on rifampin and isoniazid, the efficacies of these drugs were found to be area under the concentration-time curve (AUC)/MIC driven, while resistance suppression and postantibiotic effect were found to be related to the peak concentrations/MIC (2-5). This implies that if the MIC rises, the effect can be compensated for by increased doses. The dependence of sterilizing effect and resistance suppression on peak drug concentration and AUC, and indeed on AUC/MIC and peak concentration/MIC, has recently been demonstrated in patients for all the first-line antituberculosis drugs (6, 7). Both AUC and peak concentration are increased by higher doses. Thus, as Alffenaar et al. suggest, increased doses will lower the breakpoints. In the original PK/PD-based derivation of the new breakpoints 5 years ago and in the clinical validation of these breakpoints, we emphasized that breakpoints are dependent on the dose being administered (8–10). Thus, for those drugs, such as rifampin, isoniazid, pyrazinamide, and ethambutol (and quinolones), for which the maximum tolerated doses are far above what we currently administer, it is a good solution indeed to increase the dose. Higher doses of these compounds are likely to be well tolerated based on current clinical trials (as Alffenaar et al. point out) and also based on our reanalysis of earlier clinical trials. Increasing the AUC/MIC and peak/MIC ratios will extend the efficacy of the regimen against organisms with a wider range of MICs and the number of patients with favorable responses, thereby changing the breakpoint MIC, as we have pointed out especially for rifampin and pyrazinamide (4, 9). Therefore, we agree that increasing the dose may obviate the need to change the regimen to a second-line regimen.

On the other hand, clinicians are often reluctant to increase doses. Where there is concern that doses high enough to be effective may be toxic, replacement of the drug deemed to have an MIC indicative of drug resistance with a fluoroquinolone, such as moxifloxacin, gatifloxacin, or levofloxacin, at the correct dose may be a good solution. An alternative solution would also be to measure the drug concentrations achieved in patients and, if these are low, to increase the dose of the particular drug by using a Bayesian approach. The last approach has the virtue of allaying the fears of those who worry about concentration-driven toxicity, since the concentration of the drug to be dosed higher is low to begin with. Individual dose adjustment based on drug concentration measurement and MIC determination may even reduce toxicity.

REFERENCES

- Alffenaar JWC, Akkerman OW, Bolhuis MS, de Lange WCM, van der Werf TS. 2015. Breakpoints and drug exposure are inevitably closely linked. Antimicrob Agents Chemother 59:1384. http://dx.doi.org/10.1128 /AAC.04485-14.
- Jayaram R, Gaonkar S, Kaur P, Suresh BL, Mahesh BN, Jayashree R, Nandi V, Bharat S, Shandil RK, Kantharaj E, Balasubramanian V. 2003. Pharmacokinetics-pharmacodynamics of rifampin in an aerosol infection model of tuberculosis. Antimicrob Agents Chemother 47:2118–2124. http://dx.doi.org/10.1128/AAC.47.7.2118-2124.2003.
- Jayaram R, Shandil RK, Gaonkar S, Kaur P, Suresh BL, Mahesh BN, Jayashree R, Nandi V, Bharath S, Kantharaj E, Balasubramanian V. 2004. Isoniazid pharmacokinetics-pharmacodynamics in an aerosol infection model of tuberculosis. Antimicrob Agents Chemother 48:2951– 2957. http://dx.doi.org/10.1128/AAC.48.8.2951-2957.2004.
- Gumbo T, Louie A, Deziel MR, Liu W, Parsons LM, Salfinger M, Drusano GL. 2007. Concentration-dependent *Mycobacterium tuberculo*sis killing and prevention of resistance by rifampin. Antimicrob Agents Chemother 51:3781–3788. http://dx.doi.org/10.1128/AAC.01533-06.
- Gumbo T, Louie A, Liu W, Brown D, Ambrose PG, Bhavnani SM, Drusano GL. 2007. Isoniazid bactericidal activity and resistance emergence: integrating pharmacodynamics and pharmacogenomics to predict efficacy in different ethnic populations. Antimicrob Agents Chemother 51:2329–2336. http://dx.doi.org/10.1128/AAC.00185-07.
- Chigutsa E, Pasipanodya JG, Visser ME, van Helden PD, Smith PJ, Sirgel FA, Gumbo T, McIlleron H. 2015. Impact of nonlinear interactions of pharmacokinetics and MICs on sputum bacillary kill rates as a marker of sterilizing effect in tuberculosis. Antimicrob Agents Chemother 59:38–45. http://dx.doi.org/10.1128/AAC.03931-14.
- Pasipanodya JG, McIlleron H, Burger A, Wash PA, Smith P, Gumbo T. 2013. Serum drug concentrations predictive of pulmonary tuberculosis outcomes. J Infect Dis 208:1464–1473. http://dx.doi.org/10.1093/infdis/jit352.
- Gumbo T. 2010. New susceptibility breakpoints for first-line antituberculosis drugs based on antimicrobial pharmacokinetic/pharmacodynamic science and population pharmacokinetic variability. Antimicrob Agents Chemother 54:1484–1491. http://dx.doi.org/10.1128/AAC.01474-09.
- 9. Gumbo T, Chigutsa E, Pasipanodya J, Visser M, van Helden PD, Sirgel FA, McIlleron H. 2014. The pyrazinamide susceptibility breakpoint above which combination therapy fails. J Antimicrob Chemother 69: 2420–2425. http://dx.doi.org/10.1093/jac/dku136.
- Gumbo T, Pasipanodya JG, Wash P, Burger A, McIlleron H. 2014. Redefining multidrug-resistant tuberculosis based on clinical response to combination therapy. Antimicrob Agents Chemother 58:6111–6115. http://dx.doi.org/10.1128/AAC.03549-14.

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