

Short Communication

## Comparison of resazurin microtiter assay performance and BACTEC MGIT 960 in the susceptibility testing of Brazilian clinical isolates of *Mycobacterium tuberculosis* to four first-line drugs

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Submitted: May 24, 2011; Approved: July 2, 2012.

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### Abstract

We assessed the performance of REMA in comparison with BACTEC MGIT 960 in the susceptibility testing of 80 *Mycobacterium tuberculosis* clinical isolates from Clemente Ferreira Institute against four drugs. REMA proved to be a rapid and accurate method, providing excellent correlation with BACTEC MGIT 960, with the exception of results for the ethambutol drug.

**Key words:** *Mycobacterium tuberculosis*, tuberculosis, drug susceptibility testing.

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Tuberculosis (TB) remains a serious health problem, especially in underdeveloped and developing countries (Sanchotene *et al.*, 2008). The rising incidence of drug resistant TB gives cause for concern around the world (WHO, 2010). Currently, Brazil occupies the 19th position among the 22 countries with the most cases of TB prevalence (WHO, 2010). The initial treatment for TB involves four drugs (MS/SVS, 2009): isoniazid (INH), rifampicin (RMP), pyrazinamide (PZA) and ethambutol (EMB). Big cities such as São Paulo have high population densities and the high human immunodeficiency virus (HIV) infection rates and large numbers of institutionalized and homeless people contribute to the high prevalence of TB (Ferrari *et al.*, 2010). In this population, other factors contribute to the number of TB cases, such as irregular supply of drugs, unsuitable medical prescriptions and abandonment of treatment (Vareldzis *et al.*, 1994). Low-cost, sustainable and accessible methods (Ferrari *et al.*, 2010) that also exhibit high sensitivity, specificity and accuracy are indispensable in the TB control program, even as to achieve rapid detection of *Mycobacterium tuberculosis* in patients and characterization of drug resistance in the clinical isolates.

BACTEC MGIT 960 TB System is considered a simple and fast *in-vitro* diagnostic instrument, designed and optimized for the fast detection of mycobacteria from clinical specimens (Somoskövi *et al.*, 2000). Resazurin Microtiter Assay (REMA) is a simple and inexpensive method utilizing resazurin, a redox indicator (Franzblau *et al.*, 1998), to test the antimicrobial activity of drugs against clinical isolates of *M. tuberculosis* in low-income countries (Palomino *et al.*, 2002).

We aimed to assess the performance of the REMA method in determining patterns of susceptibility to four first-line drugs in *M. tuberculosis* clinical isolates from Brazil, by comparison with BACTEC MGIT 960, taken as a gold standard employing Minimum Inhibition Concentration (MIC) estimation and cut-off values for each drug.

Eighty *M. tuberculosis* clinical isolates from Clemente Ferreira Institute located in São Paulo city were analyzed by the BACTEC MGIT 960 kit and REMA method to determine their susceptibility patterns to the drugs INH, RMP, streptomycin (STR) and EMB. Multi-drug resistance (MDR) was taken to be resistance to at least INH and RMP. Reference susceptibility patterns were obtained with

BACTEC MGIT 960, performed at the Clemente Ferreira Institute, the TB reference center for São Paulo (SP) state, utilizing commercial kits supplied with fixed concentrations of 0.1 µg/mL, 1.0 µg/mL, 1.0 µg/mL and 5.0 µg/mL, for INH, RMP, STR and EMB, respectively (Siddiqi *et al.*, 2006).

REMA (Palomino *et al.*, 2002) was carried out with a standardized bacterial inoculum, 96-well plate (Nunc Thermo Fisher Scientific, Waltham, MA) and resazurin (Sigma-Aldrich, Steinheim, Germany) as developing agent, to reveal bacterial viability and growth. Each test was carried out in triplicate. MIC was defined as the lowest concentration that inhibited 90% of *M. tuberculosis* growth (Palomino *et al.*, 2002). Comparative analysis between the BACTEC MGIT 960 kit and REMA data allowed the breakpoint concentration to be determined with the MedCalc Software (Mariakerke, Belgium), by means of the Receiver Operating Characteristic (ROC) curve. Accuracy of REMA was evaluated from the area under the curve (AUC): 1 - 0.9 = excellent; 0.9 - 0.8 = good; 0.8 - 0.7 = moderate and < 0.7 weak correlation.

Among the eighty *M. tuberculosis* clinical isolates, BACTEC MGIT 960 detected 57 (71.2%) INH-resistant isolates, 49 (61.2%) RMP-resistant isolates, 29 (36.2%) STR-resistant isolates, 21 (26.2%) EMB-resistant isolates and 48 (60%) MDR isolates (Table 1). REMA determined the MIC values and the MedCalc Software determined cut-off values for each drug, based on the ROC curve concept (Table 1). ROC curve analysis resulted in sensitivity of 100%, 97.2%, 92.6% and 89.5%, specificity of 100%, 100%, 90% and 56.4% and accuracy levels of excellent, excellent, good and reasonable, with cut-off values of 0.0625 µg/mL, 0.125 µg/mL, 0.25 µg/mL and 8 µg/mL for INH, RMP, STR and EMB, respectively. Once the values had been fixed, all clinical isolates were classified as sensitive or resistant, according to their MIC values. REMA results demonstrated that, among the 80 *M. tuberculosis* clinical isolates, 63 (75%) were INH-resistant, 41 (51.2%) were RMP-resistant, 43 (53.7%) were STR-resistant and 47 (58.7%) were EMB-resistant. Simultaneous resistance to INH and RMP was found in 39 (48.7%) isolates, thus considered as MDR (Table 1). Some authors (5, 3, 8) based on these observations have proposed the existence of three categories of susceptibility: susceptible, partially resistant (isolates with MICs close to the cut-off value) and resistant (Mengatto *et al.*, 2006).

Based on a sub-classification (Palomino *et al.*, 2002, Tudó *et al.*, 2010) of resistant isolates by REMA, a division was made between high and low resistance, with cut-off values of 1 µg/mL, 2 µg/mL, 1 µg/mL and 16 µg/mL, for INH, RMP, STR and EMB, respectively. As result of this sub-classification, the resistant isolates were divided into: 53 (84.1%) INH high-resistance and 10 (15.8%) INH low-resistance (Figure 1a); 30 (73.1%) RMP high-resistance and 11 (26.8%) RMP low-resistance (Figure 1b); 19

**Table 1** - Results for 80 *M. tuberculosis* clinical isolates analyzed by BACTEC MGIT 960 and REMA.

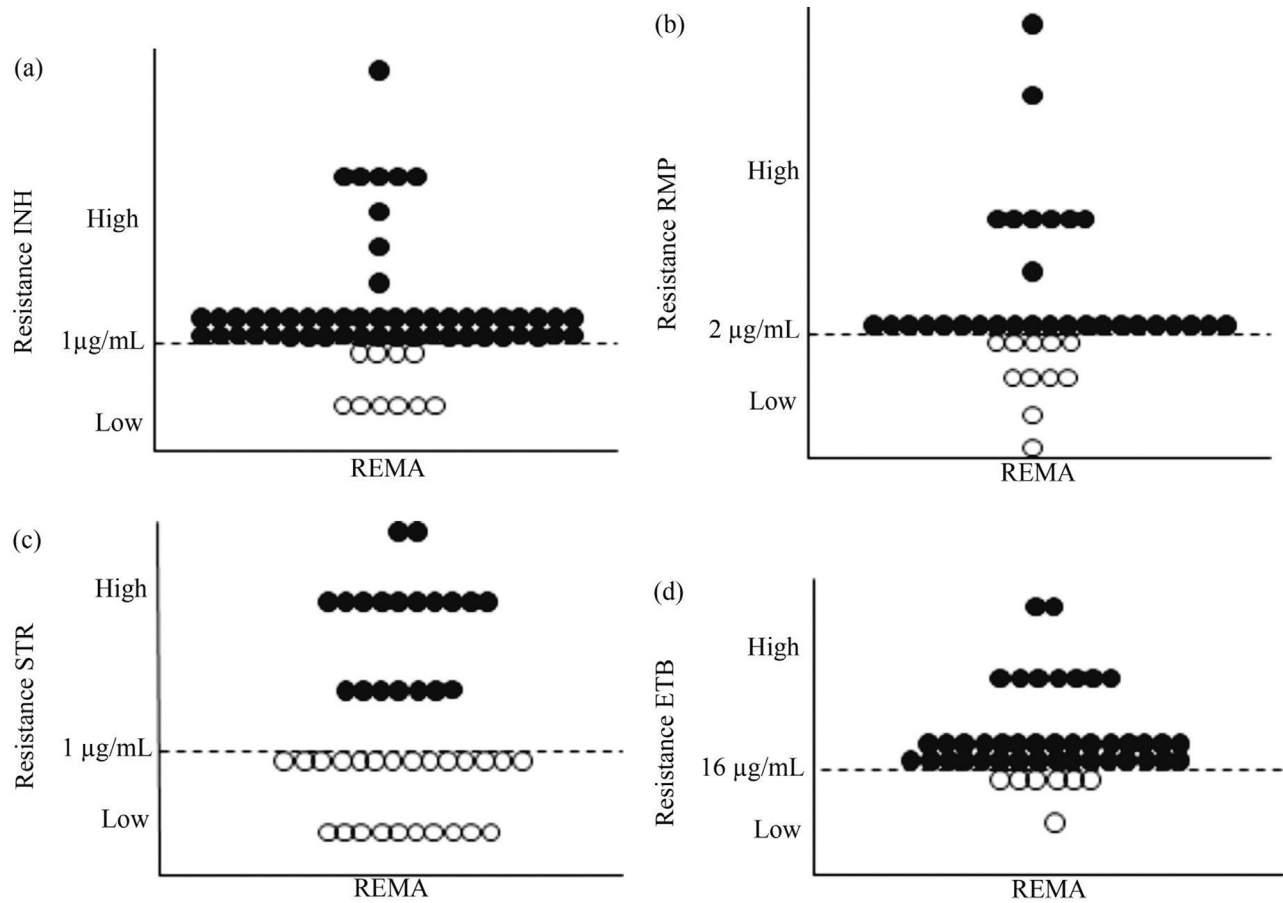
Drug	BACTEC MGIT 960 resistance	REMA resistance	Statistical analysis cut-off value (µg/mL)
INH	57 (71.2%)	63 (75%)	> 0.0625
RMP	49 (61.2%)	41 (51.2%)	> 0.125
STR	29 (36.2%)	43 (53.7%)	> 0.25
EMB	21 (26.2%)	47 (58.7%)	> 8
MDR	48 (60%)	39 (48.7%)	

(44.1%) STR high-resistance and 24 (55.8%) STR low-resistance (Figure 1c) and 40 (85.1%) EMB high-resistance and 7 (14.8%) EMB low-resistance (Figure 1d).

The high percentage of MDR isolates found in this study was probably related to the main objective of the Clemente Ferreira Institute, to assist TB patients with a difficulty history of TB treatment. Studies performed in the period from 1995 to 1998 also revealed a high prevalence of MDR cases (Melo *et al.*, 2003), characterized mainly by treatment failure and abandoned and inappropriate prescriptions (Jardim *et al.*, 2001).

Drug susceptibility patterns of *M. tuberculosis* clinical isolates were tested by REMA because this methodology enables the resistance level to be assessed with more precision in terms of MIC values (Heifets, 1988). The correct evaluation of the MIC is important because if resistant strains are classified as sensitive, the treatment will not reduce the patient's suffering and this resistant strain will spread to other people. Besides that, if sensitive strains are classified as resistant, the patient will be treated with unnecessarily toxic and expensive drugs, resulting in overload of the health system (Ahmad *et al.*, 2007) and patients with collateral effects. Cut-off values found in this paper were lower than those reported by other authors (Montoro *et al.*, 2005, Jadaun *et al.*, 2007). Such differences probably arise bacterial populations coming from different sources, with different susceptibility patterns and proportions of resistant isolates in each study, resulting in different cut-off values for each drug. Despite the discrepancies found, the authors agree that REMA is a fast method for MDR strain detection (Montoro *et al.*, 2005, Jadaun *et al.*, 2007, Rivoire *et al.*, 2007).

For INH, there were no discordant results between REMA and BACTEC MGIT 960 (Table 2) and the sensitivity and specificity of REMA were excellent. Similar results are found in the literature (Palomino *et al.*, 2002, Luna-Herrera *et al.*, 2003, Montoro *et al.*, 2005, Mengatto *et al.*, 2006, Nateche *et al.*, 2006, Rivoire *et al.*, 2007), indicating the viability of REMA in testing patterns of susceptibility to INH, even though cut-off values range between 0.0625 and 0.225 µg/mL and different results for sensitivity and specificity are found in the literature. RMP showed one discrepancy (1 false sensitive) (Table 2), resulting in a



**Figure 1** - Sub-classification within resistant isolates by REMA. (a) INH-resistant isolates. (b) RMP-resistant isolates. (c) STR-resistant isolates. (d) EMB-resistant isolates. Full circles represent high-resistant isolates and empty circles represent low-resistant isolates.

lower sensitivity than that of INH, but with the same specificity and also excellent correlation, with similar results for sensitivity (95%) and specificity (100%) in the literature (Palomino *et al.*, 2002, Luna-Herrera *et al.*, 2003, Montoro *et al.*, 2005, Nateche *et al.*, 2006, Rivoire *et al.*, 2007), as well as excellent correlation (Palomino *et al.*, 2002, Luna-Herrera *et al.*, 2003, Mengatto *et al.*, 2006). For STR, there were 4 discordant results (1 false resistant and 3 false sensitive) (Table 2), resulting in lower values of both sensitivity and specificity and excellent correlation, with similar published results ranging between 88.5 and 100% (Palomino *et al.*, 2002, Luna-Herrera *et al.*, 2003, Montoro *et al.*, 2005, Nateche *et al.*, 2006, Rivoire *et al.*, 2007), as well as an excel-

lent correlation (Luna-Herrera *et al.*, 2003). EMB showed 26 discordant results (3 false resistant and 23 false sensitive) (Table 2), resulting in a lower value of sensitivity, low value of specificity (Montoro *et al.*, 2005) and reasonable correlation; previous sensitivity results ranged between 92 and 98% (6, 8, 13), specificity results between 98 and 100% (Luna-Herrera *et al.*, 2003, Jadaun *et al.*, 2007) and correlation was reasonable (Madison *et al.*, 2002). According to the literature, the INH high-resistance isolates showed similar results to those of Palomino *et al.* (Palomino *et al.*, 2002), RMP and STR high-resistance isolates showed worse results than Palomino *et al.* (2002) and Tudó *et al.* (2010), respectively, and EMB high-resistance isolates showed similar results to those of Siddiqi *et al.* (1985).

STR and EMB, but especially EMB, have important factors that may have contributed to the discrepancies in the results: (i) a small proportion of drug-resistant strains in the population studied, (ii) substantial differences in the volume of medium utilized by the two methods, and (iii) different periods of incubation, which may result in different degrees of degradation of the drugs (Mengatto *et al.*, 2006). Partially resistant isolates with borderline MICs were not found in this study, so that MIC values were well defined as susceptible or resistant. Furthermore, some authors have

**Table 2** - Results from 80 *M. tuberculosis* clinical isolates analyzed by BACTEC MGIT 960 and REMA showing numbers of agreements and disagreements results.

REMA	INH		RMP		STR		EMB	
	R	S	R	S	R	S	R	S
R	41	0	35	0	26	3	13	23
S	0	16	1	21	1	27	3	18
Total	41	16	36	21	27	30	16	41

shown that these “borderline” cultures, more frequently associated with EMB resistance, have classically caused problems in the interpretation of EMB susceptibility testing (Siddiqi *et al.*, 1985), adversely affecting the overall performance of the assay. On the other hand, for STR, a low level of sensitivity was obtained.

Finally, our results demonstrate that REMA is a rapid method for the determination of the susceptibility to drugs of *M. tuberculosis* clinical isolates in reference laboratories such as Clemente Ferreira Institute, which shows a high level of accuracy when compared with BACTEC MGIT 960, except in tests with the drug EMB.

## Acknowledgments

We thank CNPQ and FAPESP for financial support and the Clemente Ferreira Institute for *M. tuberculosis* clinical isolates.

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