Drug-susceptibility tests in tuberculosis: contradictory or neglected?

Testes de sensibilidade no contexto da tuberculose: contraditórios ou negligenciados?



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ABSTRACT

Tuberculosis continues to have a negative impact on public health, especially its multidrug-resistant variant. Consequently, rapid molecular tests are used to identify the genetic mutations responsible for resistance to the first- and secondline drugs used for treatment. However, these tests have detection limitations in some clinical cases. Therefore, all available information must be collected and validated. Here, we report a case where the laboratory results showed discrepancies between molecular and phenotypic methods—an aspect that had not been considered in the case's conclusion. This led to the discussion regarding noncompliance with established clinical care algorithms, which deserve to be revisited.

Headings: Tuberculosis, Multidrug-Resistant; Real-Time Polymerase Chain Reaction; Genetic Testing; Case Reports.

RESUMO

A tuberculose continua a impactar negativamente a saúde pública, especialmente quando multidrogarresistente. Uma das estratégias neste contexto é a utilização de testes rápidos que identificam, por meio da biologia molecular, as principais mutações genéticas responsáveis pela resistência aos fármacos da primeira e segunda linhas do seu tratamento específico. Existem situações clínicas, no entanto, em que estes exames possuem limitações para sua detecção, motivo pelo qual todas as informações disponíveis precisam ser adequadamente coletadas e checadas. Relatamos aqui um caso onde os resultados laboratoriais apresentaram divergências entre metodologias moleculares e fenotípicas, ao mesmo tempo em que o conjunto destas informações todas não foi levado em conta no encerramento do caso. Levantou-se, assim, a discussão prática sobre a inobservância dos algoritmos assistenciais já estabelecidos, os quais merecem ser revisitados.

Descritores: Tuberculose Resistente a Drogas; Reação em cadeia da polimerase em Tempo Real; Testes Genotípicos; Relato de Caso.

INTRODUCTION

Although Brazil is not considered a high-burden country for multidrug-resistant tuberculosis, the underdetection of the disease and the unsatisfactory outcomes of its treatment in recent years are a real threat to the disease's elimination in our country¹. Therefore, one of the goals to which Brazil committed at the first high-level meeting on tuberculosis held in 2018 by the World Health Organization (WHO) was to make new and better tests available for point-of-care diagnosis and to carry out drug sensitivity tests for the rapid diagnosis of cases with tuberculosis resistant to the primary drugs used in treatment¹. Consequently, molecular techniques have emerged, especially those based on polymerase chain reaction (PCR), to identify the DNA of the *Mycobacterium tuberculosis* complex. In addition to being a rapid molecular test (RMT) for the diagnosis of tuberculosis, it identifies mutations in the *rpoB* gene that confer resistance to rifampicin. Another test available is the Line Probe Assay (LPA), which detects mutations that confer resistance not only to rifampicin, but also to other drugs used in first- and second-line treatment²⁻⁴. The commercial tests most widely used in our country are the Xpert MTB/RIF[®] Ultra[®] (Cepheid) and GenoType MTBDRplus[®] (Hain Lifescience) tests.

However, there are rare mutations outside the target region of these two tests, which may remain undetected⁵⁻⁷. Considering this limitation, the Brazilian Ministry of Health recommends that the two tests be used for the rapid screening for rifampicin resistance, and mycobacteria cultures must be used as a complementary method, especially in patients living with HIV/AIDS (PLWHA) or those who are not progressing well in treatment^{4,6}.

We report a case where conflicting results between the different *M. tuberculosis* sensitivity tests were obtained and interpreted late, consequently revealing failure to comply with established clinical care algorithms, which deserve to be revisited.

CASE REPORT

The patient was a 27 year old transexual woman, a former sex worker who had been incarcerated for 8 years and was released about 30 days before our first appointment. Her medical history dated back 5 months when she had a cough with yellowish sputum, fever in the afternoon (not measured), night sweats, dyspnea on medium exertion, and unquantified weight loss. She was diagnosed with tuberculosis in the penal institution based on RMT, without detection of resistance to rifampicin. The date of this first exam was designated as "D1" in the chronological sequence that will be used henceforth. Following the initial tuberculosis diagnosis, the patient began treatment with RHZE (5 fixed-dose combination tablets of rifampicin 150 mg + isoniazid 75 mg + pyrazinamide 400 mg + ethambutol 275 mg) for 2 months. She proceeded to the second phase of treatment, with daily doses of rifampicin 750 mg and isoniazid 375 mg for another 4 months, completing a total of six months of treatment on D180, while still imprisoned. She progressed with weight gain and improved dyspnea, and ceased to have fever soon after; although, she remained with a productive cough throughout this period. She said he had only had a chest X-ray and a single HIV test (which was negative) at the beginning of the treatment. Moreover, we did not have access to her medical records from the prison system.

Thirty days after the end of the tuberculosis treatment and already released, the patient began to have episodes of afternoon fever (not measured), night sweats, and weight loss, and was referred from the basic health unit to our specialized outpatient department at D210. On reviewing the patient's tests available on the Laboratory System Manager (GAL) of the Central Public Health Laboratory (LACEN), we found that the same sample collected on D1 had been sent for LPA testing, the result of which was released on D73. The LPA test showed a mutation in the inhA gene and none in the katG and rpoB genes, and the report read "sensitive to rifampicin and resistant to isoniazid." The same sample from D1 was also used for culture in automated liquid culture medium (BD BACTEC Mycobacterium Growth Indicator Tube - MGIT 960[®]), which showed *M. tuberculosis* growth, and a sensitivity test (SIRE®) showed resistance to isoniazid and rifampicin and sensitivity to streptomycin, ethambutol, amikacin, levofloxacin, moxifloxacin, and bedaquiline. These last two results were released on D159.

Another important piece of information was obtained from the repetition of the HIV test at a primary care unit, because of the patient's new complaints, which came back positive. Thus, she was prescribed antiretroviral therapy with tenofovir, lamivudine, and dolutegravir. In a sample taken in the first visit, D210, the HIV viral load count was 29 copies/ml, CD4 T lymphocytes count was 1,360 cells/ mm³, and CD8 T lymphocytes count was 513 cells/ mm³. On that same day, the patient underwent a chest CT scan, which showed cavitations in both lung apices and areas with a tree-in-bud pattern (Figure 1). In view of the clinical picture, radiological images, and available laboratory test results, the diagnosis of active, multidrug-resistant tuberculosis (MDR-TB) to rifampicin and isoniazid was made. Consequently, the treatment consisted of bedaquiline (400 mg for 14 days followed by 200 mg thrice a week), levofloxacin (1,000 mg once a day), terizidone (750 mg once a day), and clofazimine (100 mg once a day), the latter replacing linezolid, which was in short supply in Brazil in 2024. The smear microscopy of a sputum sample collected on D210 was negative but the RMT



Figure 1. Tomographic sections performed on D210 showing thickwalled cavitations in both upper lobes (larger on the right), associated with consolidations. There are also some areas with centrilobular opacities, tree-in-bud infiltrates and bronchiectasis, more evident in the right lung.

was positive, showing no resistance to rifampicin (result released on D211). A second sample collected on D211 showed paucibacillary TB. There was no growth of mycobacteria in the culture of the sputum from D210, nor of the other two samples collected on D240 and D270 (whose smear microscopy results were also negative). Table 1 shows the chronological sequence of all the specific tests and their results, for educational purposes. The patient continues to be monitored on an outpatient basis with good adherence to treatment, no clinical complaints, and radiological improvement.

Table 1. Summary of tests carried out and respective dates of collection and release of results.

Date of	Clinical	Test	Technique or	Results	Release date
collection	sample	performed	Commercial kit		
D1	Sputum	qPCR	Xpert MTB/RIF Ultra [®] (Cepheid)	<i>M. tuberculosis</i> complex DNA detected, sensitive to rifampicin	D1
		LPA	GenoType MTBDRplus [®] (Hain Lifescience)	<i>inh</i> A gene: mutation detected <i>kat</i> G gene: mutation not detected <i>rpo</i> B gene: mutation not detected	D73
		culture	BD BACTEC <i>Mycobacterium</i> Growth Indicator Tube (MGIT) 960®	<i>M. tuberculosis</i> complex	D159
		Sensitivity Test	BD BACTEC MGIT 960® SIRE®	Sensitive to streptomycin, ethambutol, amikacin, levofloxacin, moxifloxacin, and bedaquiline. Resistant to rifampicin and isoniazid	D159
D210	Sputum	AFB test	Ziehl-Neelsen staining	Absence of AFB	D210
		qPCR	Xpert MTB/RIF Ultra® (Cepheid)	<i>M. tuberculosis</i> complex DNA detected, sensitive to rifampicin	D211
		culture	Ogawa-Kudoh solid medium	No growth	-
D211	Sputum	AFB test	Ziehl-Neelsen staining	Presence of AFB (1+/4+)	D211
D240	Sputum	AFB test	Ziehl-Neelsen staining	Absence of AFB	D240
		culture	Ogawa-Kudoh solid medium	No growth	-
D270	Sputum	AFB test	Ziehl-Neelsen staining	Absence of AFB	D270
		culture	BD BACTEC <i>Mycobacterium</i> Growth Indicator Tube (MGIT) 960®	No growth	-

Legend - D1 to D211: sequential days; qPCR: real-time polymerase chain reaction; LPA: line probe assay; AFB test: direct test for acid-fast bacilli;[®]: registered trademark.

DISCUSSION

Sensitivity tests (ST) are essential for detecting resistance to the drugs used to treat tuberculosis. These tests can be performed using phenotypic and genotypic methods, with the latter enabling a faster and more timely diagnosis because they can be performed directly on the initial sputum sample, provided that the sputum smear or RMT are positive^{3,4}. Conversely, all phenotypic methods (and genotypic methods for extrapulmonary or pulmonary samples other than sputum) can only be carried out using the culture isolate⁴.

The Xpert MTB/RIF® assay was the first RMT prequalified by the WHO for the diagnosis of resistance in tuberculosis and is based on the amplification of the "Rifampicin Resistance Determining Region" (RRDR) of the rpoB gene of M. tuberculosis. However, a lower sensitivity of the method has been described for paucibacillary samples in general and for detecting the C533G mutation of the aforementioned rpoB gene². The Ultra® version of Xpert MTB/RIF® was developed to overcome these limitations, as well as to identify the Q513Q and F514F mutations, which are silent mutations³. However, LPA makes it possible to detect or infer mutations in genes other than the rpoB gene, such as inhA and katG for first-line drugs (related to the of resistance primary drugs like rifampicin, isoniazid, and ethionamide) and the gyrA, gyrB, rrs, and eis genes for second-line drugs (related to the resistance to fluoroquinolones and aminoglycosides/ cyclic peptides)4.

In our case, although RMT and LPA did not detect resistance to rifampicin, the LPA identified the isoniazid resistance mutation-a result that was available in the first month of the second phase of basic tuberculosis treatment. Moroever, this would have been the usual timing during the treatment to have access to the results of the mycobacteria culture. However, unfortunately, they were released 5 months later. Despite this, the information on isoniazid resistance obtained in the third month of treatment was sufficient to refer the patient to our department, which is a reference center for cases of resistant tuberculosis in the state of Goiás. Additionally, the Brazilian Ministry of Health recommends that "if only an inferred mutation in the inhA gene is identified without a mutation in the katG gene, phenotypic ST for isoniazid and 2nd line ST should be carried out, in addition to starting the drug-resistance regimen according to the resistance profile"4.

We don't know if any control bacilloscopy was carried out (ideally in the second month of treatment) when the patient was still deprived of her liberty, and she told us that there was no repeat radiological exam. The absence of these two tests can influence cure outcomes. A study conducted in the Brazilian state of Paraíba showed that failure to achieve a cure and abandonment of tuberculosis treatment in the population deprived of liberty are mainly associated with not having a follow-up bacilloscopy and with acquired immunodeficiency syndrome8. These "small" failures, often underestimated, in the management of tuberculosis can have major repercussions on the chain of processes and practices related to patient care, impacting directly or indirectly on the health of the patient and on public health as a whole. In our case, the patient had a satisfactory progress (she gained weight, her dyspnea improved, and she no longer had a fever) despite the inadequate treatment, which is a fairly common situation; however, this does not justify noncompliance with the recommended algorithms⁶. Finally, the results of the culture and the respective phenotypic ST, although released late, would have been another reason to refer the patient to a reference center before the end of the treatment.

Although we are basically dealing with deficits in the flow of intersectoral communication (e.g., in the sharing of information between the laboratory and the point of care, in this case the prison system), we need to better understand the limitations of laboratory methods. Most of the mutations of interest for identifying tuberculostatic resistance occur in the codons covered by Xpert MTB/RIF® Ultra® and GenoType MTBDRplus®. However, there are other less frequent mutations that can occur outside of the RRDR⁴. Chan et al. described in 2023 a case of the 1572F mutation identified by genetic sequencing of the rpoB gene⁵, previously identified in Hong Kong, South Africa, and Eswatini (former Swaziland). In the latter country, it is believed that 30% of cases of resistant tuberculosis go undiagnosed because they are attributed to this mutation⁷. This condition has not been investigated or proven in our case. Genetic sequencing may even be justifiable for epidemiological purposes and for studying resistance profiles in certain countries, but the required time, infrastructure, and investment must be taken into account; ultimately, its cost-benefit is inferior to that achieved by using official algorithms^{4,6}. Moreover, culture for all tuberculosis cases confirmed by Xpert MTB/RIF® is part of these algorithms, especially in special populations like

PLWHA and those deprived of their liberty, for whom sending a sample for culture should be simultaneous with the RMT itself⁶.

It is to be expected that the management of tuberculosis in the prison system faces obstacles within this system, such as the difficulty of accessing tests in proper laboratories9 and outside it (accessibility to referrals and counter-referrals), or even in relation to the low completeness of compulsory notification data¹⁰. This reality alone needs to be addressed more effectively by the authorities to minimize the role of the prison environment as the main determinant of tuberculosis transmission, which has been estimated at 36.9% for prisoners >15 years of age in Brazil¹¹. But it also underlines the need for medical teams to focus on this vulnerable population that is more subject to aggravating factors, such as social, health, economic, behavioral, and cultural inequalities, often overlapped with greater organizational limitations on the health services offered to them¹.

CONCLUSION

This case report urges us to understand the importance of recognizing and interpreting the results of drug-susceptibility tests in tuberculosis. Additionally, it highlights the responsibility and commitment of all professionals involved in the diagnostic and care sectors to "always check" this information, which is often delayed by several months. Equally important is the need for professionals to understand and adhere to the protocols and algorithms proposed by the National Tuberculosis Control Program.

> "This case report deserved an official declaration of acknowledgement and ethical approval by its institution of origin and was peer-reviewed before publication, whilst the authors declare no fundings nor any conflicts of interest concerning this paper. It is noteworthy that case reports provide a valuable learning resource for the scientific community but should not be used in isolation to guide diagnostic or treatment choices in practical care or health policies. This Open Access article is distributed under the terms of the Creative Commons Attribution License (CC-BY), which allows immediate and free access to the work and permits users to read, download, copy, distribute, print, search, link and crawl it for indexing, or use it for any other lawful purpose without asking prior permission from the publisher or the author, provided the original work and authorship are properly cited."

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