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DIFFERENTIATION BETWEEN *Nocardia* spp. AND *Mycobacterium* spp.: CRITICAL ASPECTS FOR BACTERIOLOGICAL DIAGNOSIS

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SUMMARY

New methodologies were developed for the identification of Nocardia but the initial diagnosis still requires a fast and accurate method, mainly due to the similarity to Mycobacterium, both clinical and bacteriologically. Growth on Löwenstein-Jensen (LJ) medium, presence of acid-fast bacilli through Ziehl-Neelsen staining, and colony morphology can be confusing aspects between Nocardia and Mycobacterium. This study describes the occurrence of Nocardia spp. in a mycobacterial-reference laboratory, observing the main difficulties in differentiating Nocardia spp. from Mycobacterium spp., and correlating isolates with nocardiosis cases. Laboratory records for the period between 2008 and 2012 were analyzed, and the isolates identified as Nocardia sp. or as non-acid-fast filamentous bacilli were selected. Epidemiological and bacteriological data were analyzed as well. Thirty-three isolates identified as Nocardia sp. and 22 as non-acid-fast bacilli were selected for this study, and represented 0.12% of isolates during the study period. The presumptive identification was based on macroscopic and microscopic morphology, resistance to lysozyme and restriction profiles using the PRA-hsp65 method. Nocardia spp. can grow on media for mycobacteria isolation (LJ and BBL MGITTM) and microscopy and colony morphology are very similar to some mycobacteria species. Seventeen patients (54.8%) were reported and treated for tuberculosis, but presented signs and symptoms of nocardiosis. It was concluded that the occurrence of Nocardia sp. during the study period was 0.12%. Isolates with characteristics of filamentous bacilli, forming aerial hyphae, with colonies that may be pigmented, rough and without the BstEII digestion pattern in PRA-hsp65 method are suggestive of Nocardia spp. For a mycobacterial routine laboratory, a flow for the presumptive identification of Nocardia is essential, allowing the use of more accurate techniques for the correct identification, proper treatment and better quality of life for patients.

KEYWORDS: Nocardia; Mycobacteria; Diagnostic; Identification.

INTRODUCTION

Nocardia species are saprophyte aerobic actinomycetes, distributed in water, soil, dust, and decaying vegetation worldwide, which may occasionally cause disease^{1,24}. Immunosuppression is an important risk factor for human nocardial disease, and pulmonary nocardiosis is the most common presentation^{2,14}. Cutaneous disease also occurs, through which *Nocardia* may disseminate, mainly in AIDS patients, causing lung and central nervous system (CNS) diseases^{16,21}. The mortality rate in patients with disseminated disease is high, and alcoholism is a risk factor in the dissemination to the CNS¹⁷.

Diagnosis is challenging as clinical and radiographic findings are not specific and microbiological identification may be difficult²⁴. The laboratory diagnosis is based on microscopy and culture isolation, but *Nocardia* can be mistaken for *Mycobacterium*, as they not only grow in specific media for mycobacteria, but also form partially acidfast beaded branching filaments, similar to those formed by rapidly growing mycobacteria¹⁸. Only experts can differentiate *Nocardia* from Mycobacteria filamentous branches^{8,18}. Moreover, it is difficult to identify *Nocardia* only by phenotypic methods because recent taxonomic changes have not been included in the existing schemes^{5,22}. Therefore, molecular methods have been used to identify these agents. This study describes the occurrence of *Nocardia* spp. in a mycobacterial-reference laboratory, focusing on the main difficulties in differentiating *Nocardia* from *Mycobacterium*, and correlating isolates with nocardiosis cases.

MATERIAL AND METHODS

This study was conducted to analyze retrospectively the results from isolates sent to the Tuberculosis and Mycobacteriosis Branch of the Adolfo Lutz Institute (NTM-IAL), São Paulo, Brazil, for mycobacteria identification. NTM-IAL only receives isolates from the State of São Paulo. Records from 2008 to 2012 were analyzed and the data was stored in a Microsoft Excel spreadsheet. Isolates were classified into two groups, 1) isolates identified as *Nocardia* sp. and 2) isolates presumptively

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identified as *Nocardia*, based on the presence of non-acid-fast branching filamentous bacilli. Epidemiological data, such as date of notification, closing date, reason for closure, HIV testing results, comorbidities and clinical evaluation of patients were obtained from TBWEB, the electronic notification system of the Center for Epidemiological Surveillance of Tuberculosis of the State of São Paulo (CVE-TB).

Isolates were grown on Löwenstein-Jensen (LJ), Sabouraud agar, and BBL MGITTM (Becton Dickinson) media, and identified by colony morphology (pigment production and colony aspects), microscopic characteristics after Ziehl-Neelsen staining, and resistance to lysozyme. Lysozyme resistance was determined by inoculating a standardized culture suspension into two glycerol broth tubes, one with (test tube) and the other without (control tube) a 50 µg/mL lysozyme. Tubes were incubated at 35 °C and examined weekly for four weeks⁴. An isolate was considered as lysozyme resistant when equivalent growth appeared in both tubes. Isolates were also submitted to the PRA-hsp65 method as described by CHIMARA *et al.*¹¹.

This study was approved by the ethical committee of Adolfo Lutz Institute under number 072D-2010.

RESULTS

During the study period, NTM-IAL identified 27,804 isolates. Group 1 was composed of 33 (0.12%) *Nocardia* sp. isolates from 31 patients, obtained from sputum (20), corneal scrapings (2), abscess secretion (1), tracheal aspirate (1), pleural fluid (1), cerebrospinal fluid (1) and not discriminated (7). Group 2 was composed of 23 (0.08%) isolates, one per patient, obtained from sputum (17), blood (1) and not discriminated (5).

Microbiological aspects. In group 1, all isolates presented branching filaments (Fig. 2) and lysozyme resistance, wet colonies on LJ medium, showing a salmon (30 isolates) or yellow (3) color (Fig. 1A). Salmon-colored colonies presented a white aspect after three weeks on Sabouraud agar (Fig. 1B), very similar to fungi growth. Growth on BBL MGITTM media presented flocculated colonies similar to *M. tuberculosis*, but pigmented (Fig. 1C). The 33 isolates showed no restriction site for the BstEII enzyme and undescribed restriction patterns with the HaeIII by PRA-hsp65 method. In group 2, all isolates were shown to be filamentous bacilli on microscopy and none presented restriction site for the BstEII (Fig. 3). Lysozyme test was not performed on those isolates.

Clinical and epidemiological characteristics. In group 1, 25 (80.6%) of the 31 patients were male, mean age of 44.1 years (SD 14.8). All had multiple comorbidities such as arthritis, silicosis, chronic pulmonary obstructive disease, smoking, alcohol and some were illicit drug users. Seventeen patients (54.8%) were notified to CVE-TB as tuberculosis (TB) cases and have therefore been treated for TB. Two of them were HIV positive. Eleven patients (68.8%) were cured after drug treatment for TB, but three relapsed and three died. Among these patients, 10 underwent chest radiography and/or computed tomography. Three of them had cavities in the lung parenchyma; in the other seven, some suggestive signs of disease activity (consolidation, nodules, masses, bronchial wall thickening or "tree-in-bud" aspect) were reported. *Nocardia* sp. was isolated twice from two patients, one with aortic valve implantation history, and the other with a history of ocular surgery. During this study, CVE-TB intervened in the treatment of two patients who had

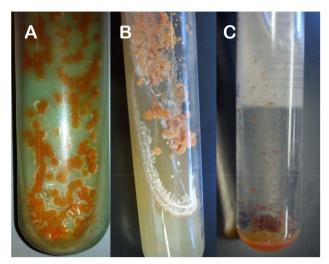


Fig. 1 - Macroscopic growth of *Nocardia* sp. in: A) Lowenstein Jensen medium, B) Agar Sabouraud and C) Bactec MGIT medium.

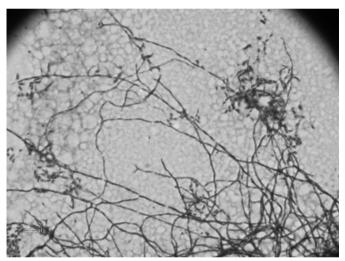


Fig. 2 - Microscopic appearance of *Nocardia* spp. isolates by Ziehl Neelsen staining. 100x increased.

been receiving antituberculosis drugs for six months. In the study period, *M. tuberculosis* from two patients and *M. kansasii* from one patient were also recovered in the same year as the *Nocardia* was isolated from them.

Among the 22 patients of group 2, 16 (72.7%) were male, mean age of 41.6 (SD 20.5). Of these, 14 (63.3%) were reported to CVE-TB and after TB treatment, nine (64.3%) were cured, two abandoned, one relapsed, one had the diagnosis changed, and one died. *M. tuberculosis* had previously been isolated from two patients, and from another individual, a *Nocardia* sp. isolate had also been recovered.

DISCUSSION

The initial diagnosis of nocardiosis requires fast and accurate methodology, mainly due to the clinical aspects and bacteriologic similarity to the genus *Mycobacterium*⁵. McNEIL & BROWN¹⁸ highlighted the importance of direct microscopic examination because, despite the new methodologies developed, there is no test replacing it.

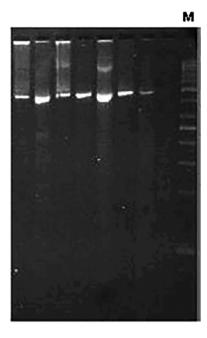


Fig. 3 - Molecular pattern of *Nocardia* spp. isolates digested by BstEII restriction enzyme. M: 50 bp molecular size marker.

In this study, 33 isolates and 17 cases of nocardiosis in a five-year period were found. Nocardia spp. isolation accounted for 0.12% of all isolates identified by NTM-IAL, from 2008 to 2011. HELAL et al.15 studied 200 sputum samples from patients diagnosed as pulmonary TB in Egypt, based on clinical symptoms, chest X-ray and direct smear test by acid-fast staining method. Four (2.0%) of these were PCR positive for 16S rRNA primer specific for Nocardia, a higher percentage than the one obtained in this study. Despite the presence of Nocardia detected by PCR, HELAL et al.¹⁵ could not obtain growth on LJ. Although the literature shows the need for special culture media¹⁷, isolates of Nocardia spp. received by NTM-IAL showed good growth on LJ medium, as well as on other media used for isolation of mycobacteria (Fig. 1). Colony morphology on LJ from many species of Nocardia may be confused with Mycobacterium species, making diagnosis difficult. Moreover, microscopic analysis should be carefully performed, because, as these two genera may present ramifications¹⁸, Nocardia may be misidentified as mycobacteria¹³ in the routine diagnosis of TB. The differentiation is performed by observation of aerial hyphae, produced only by species of the genus Nocardia¹⁸. Change in cell morphology of Nocardia depending on the age of the culture may also lead to misidentification. Young cultures exhibit branched-long filaments, while old cultures present bacilli and cocci originating from fragmentation of filaments, which can be considered mycobacteria bacilli for non-expert microscopists. Despite the Ziehl Neelsen method having been used for strong acid fast bacteria, such as the Mycobacterium species, and the modified Ziehl Neelsen method for weak acid fast bacteria, such as the Nocardia species, this study detected the Nocardia by the Ziehl Neelsen method used for mycobacteria. This fact shows the importance of an expertise reading, since both genera can be detected.

Many studies have shown that phenotypic methods are not accurate for *Nocardia* identification and the need for molecular methods is a reality^{1,5,19}. BAIO *et al.*³ identified seven isolates by 31 different phenotypic tests and analyzed profiles using six identification systems. Two isolates were identified as *Nocardia sp.*, because a common outcome was not found using different schemes. Only two isolates were correctly identified by phenotypic identification, compared to the MLSA results. Although phenotypic identification produces misidentifications, it can improve molecular identification since some molecular targets can present high sequence similarity¹².

PRA-hsp65 method¹³ is routinely used at NTM-IAL to amplify a 440 pb gene fragment that encodes a protein of 65 kDa for mycobacteria and actinomycetes⁹. As *Nocardia* lack the BstEII restriction site, this feature can be used in the presumptive identification of the genus. Additionally, *Nocardia* HaeIII patterns are different to the algorithm described by CHIMARA *et al.*¹¹.

This study showed three cases of mixed infection by different *Nocardia* and mycobacteria species in immunocompetent patients. A similar situation was described at the Baystate Medical Center, MA, USA, where bacteriological analysis revealed the presence of mixed cultures of *N. asteroides* and *M. tuberculosis*⁷ in two patients with clinical TB. The authors of this study showed the importance of bacteriological investigation in cases of treatment failure, because it may indicate that the patient harbors another agent that is not being addressed, also observed in the present study. Other cases of mixed infection by these two genera have been reported in immunocompromised patients^{10,20}.

The finding of *Nocardia* should not be considered contamination and the case should be investigated²³. In our study, 17 patients had some kind of comorbidity (silicosis, smoking, alcohol abuse) or immunosuppression, characterizing this group as high risk for nocardiosis²³.

Mycobacteriosis and nocardiosis treatments are different, so the correct identification of the agent is essential to avoid ineffective therapy¹⁵. Initially, all notified patients were treated as TB, but for 31.2% of them, treatment was not effective. In these cases, the failure of TB treatment was evaluated and new clinical, radiological and bacteriological exams were performed to establish the cause of failure. However, *Nocardia* isolation was not taken into consideration and some patients were characterized as TB relapse cases.

In Brazil, TB patients to be considered cured should have, at least, two negative smears during the follow-up phase and one more after treatment of the disease. In this survey, all patients with prior treatment for TB had this criterion⁶. Pulmonary nocardiosis should be considered as differential diagnosis when TB fails to respond with standard therapy, or if the patient's condition worsens despite adequate therapy.

The isolates of group 2 were only presumptively identified as *Nocardia* sp., a limitation of the study, and possibly they are *Nocardia* or even another actinomycete. An alternative to solve the identification can be sequencing the 16SrRNA region. BHARADWAJ *et al.*⁵, by sequencing organisms with ambiguous phenotypes, was able to identify different actinomycetes recovered from clinically significant specimens.

Therefore, patients with a history of comorbidities, immunosuppression, whose isolates present rough, pigmented colonies of filamentous aerial hyphae forming bacilli, resistant to lysozyme and MURICY, E.C.M.; LEMES, R.A.; BOMBARDA, S.; FERRAZOLI, L. & CHIMARA, E. - Differentiation between *Nocardia* spp. and *Mycobacterium* spp.: critical aspects for bacteriological diagnosis. Rev. Inst. Med. Trop. Sao Paulo, 56(5): 397-401, 2014.

without the restriction site for BstEII, are suggestive of having *Nocardia* spp. infection. In a mycobacterial-routine laboratory, an algorithm for genera and species identification of isolates suggestive of *Nocardia* spp. is essential for the correct diagnosis, which assists in providing adequate and early treatment for patients.

RESUMO

Diferenciação de *Nocardia* spp. e *Mycobacterium* spp.: aspectos críticos para o diagnóstico bacteriológico

Novas metodologias têm sido desenvolvidas para a identificação de Nocardia spp. mas o diagnóstico inicial ainda necessita de método rápido e preciso, principalmente devido à similaridade com o gênero Mycobacterium, clínica e bacteriologicamente. O crescimento em meio de Löwenstein Jensen (LJ), a presença de bacilos corados pela coloração de Ziehl Neelsen e colônias com características diferentes podem ser fatores de confusão entre nocardias e micobactérias. Este estudo descreve a ocorrência de Nocardia spp. em laboratório de referência em micobacteriologia, observando-se as principais dificuldades em diferenciar Nocardia spp. e Mycobacterium spp., correlacionando isolados com casos de nocardiose. Os registros laboratoriais dos anos 2008 a 2012 foram analisados e os isolados identificados como Nocardia sp. ou como bacilos não álcool - ácido resistentes (NBAAR) foram selecionados. Os dados epidemiológicos e bacteriológicos foram analisados. Trinta e três isolados identificados como Nocardia sp. e 22 como NBAAR foram selecionados para este estudo, perfazendo 0,12% do total de isolados identificados no período estudado. A identificação presuntiva foi baseada na morfologia macroscópica e microscópica, resistência à lisozima e perfis de restrição pelo método PRA-hsp65. Nocardia spp. pode crescer em meios de isolamento para micobactérias (LJ e BBL MGITTM) e microscopia de morfologia e as colônias são muito semelhantes a algumas espécies de micobactérias. Dezessete pacientes (54,8%) foram notificados e tratados para tuberculose, mas apresentaram sinais e sintomas para nocardiose. Concluimos que a ocorrência de Nocardia sp. no período estudado foi de 0,12%. Os isolados com características de bacilos filamentosos, formadores de hifas aéreas, com colônias que podem ter pigmento, rugosas e que não possuem padrão de digestão para BstEII no método PRA-hsp65 são sugestivos de Nocardia spp. Para um laboratório de rotina de Micobactérias, um fluxo de identificação presuntiva para Nocardia spp. é essencial para permitir que esses isolados sejam identificados com técnicas mais precisas, para que seja oferecido o tratamento adequado e qualidade de vida aos pacientes.

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