Identification of non-tuberculous mycobacteria isolated from opossum (*Didelphis virginiana*) lymph nodes and characterisation of lesions

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ABSTRACT. The aim of this study was to investigate the presence of NTM in the lymph nodes of opossums (*D. virginiana*) and to characterise the microscopic changes in affected tissue. Retropharyngeal and tracheobronchial lymph nodes were collected postmortem from 18 opossums in the state of Colima, Mexico in 2013. The lymph nodes were also cultured for mycobacterial organisms and processed for histopathological examination. Bacteriological cultures yielded 5/18X100 (28%) isolates of NTM, which were subsequently identified as *M. terrae*, *M. szulgai*, *M. gastri* and *M. asiaticum*. Microscopic examination of the affected nodes revealed a necrotic granulomatous lymphadenitis (3/60%) composed of histiocytes, epithelioid cells and giant cells with intralymphal alcohol-resistant acid bacteria. An association between the sex of the opossum and the presence of NTM was observed. To our knowledge, this is the first report of NTM isolation in opossums with granulomatous lymphadenitis in Mexico.

Key words: mycobacteria, tuberculosis, opossums, lesions.

RESUMEN. El objetivo de este estudio fue investigar la presencia de MNT en los nódulos linfáticos de los tlacuaches (*D. virginiana*) y caracterizar los cambios microscópicos en el tejido afectado. Se recolectaron 18 *D. virginiana* en el estado de Colima, México, en 2013. Los nódulos linfáticos retrofaríngeos y traqueobronquial fueron procesados para aislamiento de micobacterias e histopatología. Se obtuvieron 5 aislamientos de MNT (5/18X100=28%), de entre estos destacan las especies de *M. terrae*, *M. szulgai*, *M. asiaticum* y *M. gastri*. El examen microscópico de los nodos afectados reveló una linfadenitis necrótica granulomatosa (3/60%) compuesta de histiociotos, células epiteloides y células gigantes con bacterias ácido alcohol resistente intralymphal. Se encontró una relación entre las MNT y el sexo de los tlacuaches. Este es el primer informe de aislamiento de MNT en los tlacuaches con linfadenitis granulomatosa en México.

Palabras clave: mycobacteria, tuberculosis, tlacuache, lesión.

INTRODUCTION

There are more than 130 species of nontuberculous mycobacteria (NTM), also known as atypical or environmental mycobacteria. These opportunistic bacteria are regular inhabitants of soil, water, earth and inanimate objects. They are well distributed in our environment and have great pathogenic potential, with a clinical profile or lesions similar to those caused by tuberculosis (Casal and Casal 2007). The aim of this study was to investigate the presence of NTM in the lymph nodes of opossums (*D. virginiana*) and to characterise the microscopic changes in affected tissue. Retropharyngeal and tracheobronchial lymph nodes were collected postmortem from 18 opossums in the state of Colima, Mexico in 2013. The lymph nodes were also cultured for mycobacterial organisms and processed for histopathological examination. Bacteriological cultures yielded 5/18X100 (28%) isolates of NTM, which were subsequently identified as *M. terrae*, *M. szulgai*, *M. asiaticum* and *M. gastri*. Microscopic examination of the affected nodes revealed a necrotic granulomatous lymphadenitis (3/60%) composed of histiocytes, epithelioid cells and giant cells with intralymphal alcohol-resistant acid bacteria. An association between the sex of the opossum and the presence of NTM was observed. To our knowledge, this is the first report of NTM isolation in opossums with granulomatous lymphadenitis in Mexico.

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MATERIAL AND METHODS

HOST COLLECTION AND NECROPSY

During 2013, eighteen road-killed *D. virginiana* were collected in the state of Colima, Mexico. The site was located
at 19°41’-19°16’ N, 102°46’-103°47’ W, with an altitude of 500 meters (INEGI 2010). A necropsy identified lesions suggestive of tuberculosis. Homologous tissue sections collected from the tracheobronchial and retropharyngeal lymph nodes were preserved in a 6% borate solution (Pronabive-SAGARPA) and 10% buffered formalin.

**BACTERIOLOGY**

After the sodium borate and the excess fat were removed, the lymphoid tissue was cut into small pieces in a sterile mortar; sterile sand was added, and the tissue was manually macerated. The liquid was decanted in a 1:1 proportion into a 4% sodium hydroxide solution with phenol red, and then it was incubated at 37 °C for 20 minutes, after which it was centrifuged at 3000 x g for 20 minutes. The sediment was deposited in a sodium hypochlorite solution, which was then neutralised with a 1 N hydrochloric acid solution until a shift in the color (from pink to yellow) was observed.

Three cultures were seeded in two tubes of Stonebrink (Pronabive) culture medium and one tube of Lowenstein-Jensen medium (Pronabive), which were then incubated at 37 °C for 24 hours. After 24 hours, they were incubated at 37 °C under a 5% CO₂ atmosphere for 9 weeks. The growth obtained was verified using the Ziehl-Neelsen stain.

The isolates were identified based on their characteristics and growth morphology as well as metabolic activity, specifically, niacin production, nitrate reduction, catalase activity at 22 °C and 68 °C, urea activity, TWEEN 80 hydrolysis, growth on MacConkey agar without crystal violet and 5% NaCl tolerance. All isolates were seeded in four replicates. The four replicates were incubated for 30 days under different conditions: one culture was incubated at 22 °C, another at 45 °C and two at 37 °C under a 5% CO₂ atmosphere, one in the absence of light and another in the presence of light.

**HISTOLOGY**

The lymph nodes were assessed for the presence of NTM. They were embedded in paraffin; 4.0-micron-thick sections were cut and stained with hematoxylin-eosin (H and E), and Ziehl-Neelsen (ZN) (Prophet et al 1995).

**STATISTICAL ANALYSIS**

A chi-squared test was performed to evaluate the association between the sex of the opossum and the presence of NTM, and the prevalence rates were expressed as percentages.

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The work was subject to the provisions of the Official Mexican Standard: NOM-033- ZOO-1995 y NOM-051-ZOO-1995.

**RESULTS AND DISCUSSION**

Eighteen *D. virginiana* were collected, of which 7/18 (39%) were female and 11/18 (61%) were male. Five out of the 18 animals (28%) were positive for the isolation of NTM, and three of them presented granulomatous lesions in their lymph nodes. The NTM isolated corresponded to the species *Mycobacterium szulgai*, *M. terrae*, *M. asiaticum* and *M. gastri* (table 1 and figure 1). An association was found between the sex of the animal and its NTM status: infection was more frequent in males of *D. virginiana* (table 2). The distribution of NTM in the environment and their importance as opportunistic pathogenic organisms or even strict pathogens in humans, domestic animals and wild animals have been reported as sources of the disease called mycobacteriosis (García and García 2012). The role of NTM does not appear anywhere in the research literature about *D. virginiana*; however, life habits imply a strong interaction between them and their environment due to their varied diet and their behaviour and distribution patterns. These habitat interactions facilitate the presence of NTM in *D. virginiana*, which could eventually become a wild reservoir of the bacteria, hindering the microbiological isolation of the complex of mycobacteria when mixed infections exist. It has also been proved that some NTM interfere with today’s official diagnostic tests for bovine tuberculosis due to cross-reactivity (Jaroso et al 2010).

The problem with this interference in the epidemiological study of bovine tuberculosis in Mexico is that despite 21 years of efforts to eradicate the disease, only 82% of the national territory has less than 0.5% prevalence of bovine tuberculosis (NOM-031-ZOO-1995). The association between sex and the presence of NTM is due to *D. virginiana* having a sex ratio of 1:1, which is considered an ideal condition for population growth. However, in polygamous species such as *D. virginiana*, there tends to be a larger number of males, especially in zones where resources are more available, and competition is far less stringent. In these zones, it is believed that the sex ratio observed could be associated with the size of the population and the distribution patterns of the species, which are determined primarily by competence, predation, availability of food resources, shelter, immigration, dispersion and extinction (Cruz et al 2014). The wild boar is considered the most

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Important reservoir of tuberculosis, and it is primarily accountable for transmitting the disease to red deer (Cervus elaphus) and fallow deer (Dama dama) (García et al 2013). In Spain, diversity and co-infections of NTM and M. bovis in wild boar, fallow deer, red deer and cattle (in the south) have been described (Gortázar et al 2011). The NTM M. chelonae and M. avium have been isolated from wild boar (García et al 2015). Wild animals infected with Mycobacterium spp. may, under certain conditions, become reservoirs of such agents. In the wild, the badger, deer, opossum, ferret and buffalo are recognised as reservoirs of bovine tuberculosis (Machackova et al 2003).

In this study, we present the finding of four NTM species that nevertheless produce similar granulomatous lesions. Therefore, the opossum D. virginiana could become a reservoir of Mycobacterium spp. for wild and domestic animals.
The risk represented by *D. virginiana* is determined by its interaction with the environment due to its habitat, eating habits and behavioral patterns that expose it to mycobacteria present in environments such as water, soil, plants and animals; their coexistence with domestic animals and even with humans can spread these mycobacterial infections. This could facilitate a high presence of NTM in the lymph nodes of these animals, so it is useful to verify the presence, prevalence and variety of NTM in opossums in areas where bovine tuberculosis is endemic. These cases indicate that NTM can cause tuberculosis-like disease in *D. virginiana* and illustrates the importance of identifying causal agents of tuberculosis-like diseases in wildlife.

The lesions in the lymph nodes consisted of granulomas characterised by histiocytes, epithelial cells and giant cells; in two of the animals, acid-fast bacteria (ZN+) were observed (table 1 and figure 2). Granulomatous lesions associated with NTM have been described in black-tailed deer (Hall *et al* 2005), and granulomas in the lymph nodes of *D. virginiana* caused by *M. bovis* have been described in Mexico (Acosta *et al* 2012). Two types of lesions, pyo-granulomatous and granulomatous, have been classified in the lymph nodes of bovines in Mexico where NTM were isolated (Hernández 2014). Depending on the susceptibility of the host, granulomas caused by mycobacteria in the wild can vary in size and may or may not be well delimited by connective tissue. The literature contains a great variety of descriptions of the types of lesions (macroscopic) and granulomas (microscopic) observed in wild fauna (Zanella *et al* 2008, García *et al* 2012). Granulomatous lesions in the lymph nodes and lungs of *D. virginiana* caused by mycobacteria could be confused with lipid pneumonia, as it manifests with whitish multifocal lesions on the lung parenchyma, which are common in opossums with the lung parasite *Didelphostrongylus hayesi* (García *et al* 2012).

Another parasite similar to *Paragonimus mexicanus* causes granulomatous pneumonia in opossums (García *et al* 2010).

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**REFERENCES**


Hernández JA. 2014. Identificación de micobacterias no tuberculosas aisladas de nódulos linfáticos de bovinos y caracterización de lesiones asociadas. Tesis Maestría en Ciencias Veterinarias y Zootécnicas, Facultad de Medicina Veterinaria y Zootecnia, Universidad Autónoma de Tamaulipas, México.


