Human and canine pulmonary *Mycobacterium bovis* infection in the same household: re-emergence of an old zoonotic threat?

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**ABSTRACT**

Bovine tuberculosis (bTB), caused by *Mycobacterium bovis*, remains a serious animal health problem in the UK, despite longstanding statutory surveillance and control measures. Endemic infection in the Eurasian badger population is thought to complicate bTB eradication efforts. Sporadic cases of *M. bovis* infection have also been reported in domestic animals other than cattle. Human *M. bovis* infection is extremely rare in the native UK population in the absence of unpasteurised milk consumption or residence abroad. Here, pulmonary TB infection in a UK born female and her pet dog is described, caused by an identical strain of *M. bovis*. Latent TB infection was also identified in a household contact. The potential routes of infection and implications of this case are discussed.

**CASE NO 1: HUMAN INFECTION**

A 42-year-old UK born, Caucasian female living in Cornwall (South West of England, UK) presented in August 2007 with a 1 month history of general malaise and mucopurulent cough. Clinical examination was unremarkable but chest x ray revealed right apical cavitation (fig 1A). Sputum was smear positive for acid fast bacilli and treatment was initiated with rifampicin, isoniazid, ethambutol and pyrazinamide. Subsequent culture grew *Mycobacterium bovis* and pyrazinamide was therefore withdrawn from the drug regimen. The isolated *M. bovis* strain underwent standard molecular typing at the Veterinary Laboratories Agency (VLA, Weybridge, UK) by spoligotyping and six loci variable number tandem repeats typing.1 The genotype recovered was SB0271: 7-4-5-4*-S-3-1; VLA genotype 12:a. The patient had no history of unpasteurised milk consumption, immunodeficiency or recent travel abroad. However, there was a history of potential occupational exposure to *M. bovis* when she worked as a veterinary nurse for two local practices in West Cornwall from 2000 to 2004. In particular, she assisted in tuberculin tests of cattle herds in the area and one farm that she visited had three skin test reactors in 2002, two of which were subsequently found to culture *M. bovis* at post mortem. The patient also recalled picking up an injured badger that had suffered a road traffic accident and transporting it to a veterinary surgery for treatment in 2004; this animal was euthanased a few days later without a postmortem examination. In addition, badgers were reportedly living in a sett at the bottom of the patient’s garden.

**DISCUSSION**

Despite the increasing incidence of bovine TB (bTB) in the UK since the mid-1980s,2 *M. bovis* infections account for less than 1% of the laboratory confirmed cases of human TB, in line with other industrialised countries.1,3,4 *M. bovis* infections in animals are transmissible to humans through inhalation of infectious aerosols, ingestion of unpasteurised dairy products or, less commonly, by contact with broken skin.3,4 Historically, consumption of unpasteurised cows’ milk accounted for the vast majority of human infection in the UK.2 However, with the almost universal adoption of milk pasteurisation, it has been suggested that inhalation, rather than ingestion, may now...
represent the predominant mode of *M. bovis* transmission from animals to humans. Although *M. bovis* has a broad host range, canine *M. bovis* infection is extremely rare in the UK, with only five cases confirmed by the VLA in the past 20 years. The genotype of *M. bovis* (12:a) recovered from case Nos 1 and 3 is relatively rare, occurring in less than 2% of the 50,000 isolates from cattle recorded in the VLA Spoligotype Database. Isolates of this strain are geographically localised to cattle in the Southwest of Cornwall (fig 1B) and have not been reported outside of Great Britain to the authors’ knowledge. All but five of the 46 badger samples with genotype 12:a recorded in the Spoligotype Database were also found in this immediate area. This indicates that the infections were, in all likelihood, locally acquired. As case No 1 was a middle-aged person and had no history of unpasteurised milk consumption, the evidence points to a recent, native *M. bovis* infection, probably as a consequence of airborne exposure to an infectious animal or its excretions. The original source of infection for the two confirmed cases (human and canine) may never be definitively elucidated, but it is reasonable to assume direct or indirect transmission from cattle or badgers.

There are several hypotheses as to the source of infection in the affected household. The chronology of clinical disease suggests that case No 1 may have been the index case, with subsequent airborne transmission of infection to her daughter and pet dog. Person to person transmission of *M. bovis* infection is extremely rare, but has been recently reported among immunocompromised and immunocompetent patients. Reverse zoonotic transmission of *M. bovis* and *M. tuberculosis* from humans to animals is also rare, but has been described previously. The source of infection for case No 1 may have been through previous occupational exposure to *M. bovis* infected animals while working as a veterinary nurse. Although there was no history of immunodeficiency to suggest reactivation of latent infection, interestingly one of the cattle isolates from cattle recorded in the VLA Spoligotype Database.

**Figure 1** (A) Case No 1: chest radiograph showing right apical cavitation. (B) Map showing the location of all genotyped strains of *Mycobacterium bovis* from cattle in England and Wales in 2007 (white circles). Localisation of genotype 12:a strains to the South West of Cornwall (UK) from cattle in 2007 is shown (red circles). Over 40 isolates of 12:a from cattle were located within the black square, which marks the location of this case.

**Figure 2** Case 3: canine postmortem specimen (see online data repository), trachea (black arrow), granulomatous lesion at tracheal bifurcation (black broken arrow), left lung (white arrow) and heart (white broken arrow).
herds that she helped to tuberculin test in 2002 was found to be infected with spoligotype 12. Unfortunately, no variable number tandem repeats data are available for this herd but spoligotype 12, when subtyped, is almost invariably of genotype 12:a, the local strain of *M. bovis*. We have no reports of TB being detected in either the attending veterinarian or the herd keeper, both of whom would have had much closer and prolonged contact with any infected cattle. Nevertheless, this infected cattle herd does represent a potential source of infection for case No 1. It is also possible that case 1 could have been infected through contact with badgers, both on her property and as part of her previous work, particularly as badgers have been identified as a maintenance host of *M. bovis*.

An alternative possibility is that the tuberculous pet dog acquired *M. bovis* infection from a local animal reservoir and then infected the human contacts through airborne transmission. *M. bovis* infection does occur sporadically in companion animals living in rural and suburban areas and this is thought to represent spillover of infection from cattle and wildlife reservoirs. However, even though the actual onset of the dog’s pulmonary infection cannot be reliably established, the fact that its clinical signs were noticed 4 months after the dog’s pulmonary infection cannot be reliably established, the low, but ongoing, public health risk. The potential for airborne transmission of *M. bovis*, particularly to segments of the rural population, highlights the need for vigilance and effective collaboration between clinicians, animal and public health professionals in the management of this often forgotten infection.

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**REFERENCES**

The Guideline Development Group regret that no advice was offered for the management of status epilepticus. As this is a life-threatening condition where a patient may suffer from cerebral hypoxia (and oximetry may not be possible), patients with status epilepticus should be treated in accordance with table 1 (reservoir mask until clinically stabilised). This advice also applies to other rare conditions that may cause life-threatening hypoxaemia that are not listed specifically in table 1.


The correct authors for this abstract are as follows: Roberts JA, Bakerly ND.


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This abstract has been withdrawn.


It has come to the attention of the Scientific Committee that this abstract was not seen by all the authors prior to submission and is therefore withdrawn.


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