



Enfermedades Infecciosas y Microbiología Clínica

www.elsevier.es/eimc



Original article

Human *Mycobacterium bovis* infection in Castile and León (Spain), 2006–2015[☆]



Teresa Nebreda-Mayoral^{a,*}, M. Fé Brezmes-Valdivieso^b, Nieves Gutiérrez-Zufiaurre^c, Susana García-de Cruz^d, Cristina Labayru-Echeverría^e, Ramiro López-Medrano^f, Luis López-Urrutia-Lorente^g, Almudena Tinajas-Puertas^h, Octavio Rivero-Lezcanoⁱ

^a Servicio de Microbiología, Complejo Asistencial Universitario de León (CAULE), León, Spain

^b Sección de Microbiología, Complejo Hospitalario de Zamora, Zamora, Spain

^c Servicio de Microbiología, Hospital Universitario de Salamanca, Salamanca, Spain

^d Sección de Microbiología, Complejo Asistencial de Soria, Soria, Spain

^e Sección de Microbiología, Complejo Asistencial de Burgos, Burgos, Spain

^f Unidad de Microbiología, Hospital El Bierzo, Ponferrada, León, Spain

^g Sección de Microbiología, Hospital Universitario Río Hortega, Valladolid, Spain

^h Sección de Microbiología, Hospital de Palencia, Palencia, Spain

ⁱ Unidad de Investigación, Complejo Asistencial Universitario de León (CAULE), León, Spain

ARTICLE INFO

Article history:

Received 19 August 2017

Accepted 10 November 2017

Available online 23 November 2018

Keywords:

Mycobacterium bovis

Pulmonary tuberculosis

Extrapulmonary tuberculosis

Incidence

Risk factors

ABSTRACT

Introduction: The annual incidence of tuberculosis (TB) from *Mycobacterium bovis* in humans has considerably declined in industrialised countries since the early twentieth century. The objective of this study was to determine the epidemiological, clinical and microbiological characteristics of patients with this illness in Castile and León (CyL).

Methods: Retrospective study of all *M. bovis* TB cases in CyL over a 10-year period, comparing the risk factors, the epidemiology and the clinical course between pulmonary (PTB) and extrapulmonary TB (EPTB).

Results: 75 cases of TB were due to *M. bovis*: 45 PTB and 31 EPTB. The annual incidence of TB due to *M. bovis* was 0.3 cases per 100,000. It remained stable between the first and second five-year period (0.27 vs. 0.33, $p = 0.656$). However, the overall incidence of TB fell in both five-year periods (13.58 vs. 10.71, $p < 0.0001$). The mean age was 66.2 ± 21.3 years, mainly men (63%) and Spanish patients (92%). PTB was significantly more frequent in men, aged over 66 years, with immunosuppressive conditions or who were smokers. Mortality was 9%, associated with higher age, immunosuppression or treatment different from that recommended by the WHO.

Conclusions: The incidence of *M. bovis* TB in CyL was higher than that for Spain and for other European countries, and remained stable despite the decreased the TB due to MTC. It affected mostly Spanish-born patients who lived in rural areas and with a high mean age.

© 2017 Elsevier España, S.L.U. and Sociedad Española de Enfermedades Infecciosas y Microbiología Clínica. All rights reserved.

DOI of original article: <https://doi.org/10.1016/j.eimc.2017.11.018>

[☆] Please cite this article as: Nebreda-Mayoral T, Brezmes-Valdivieso MF, Gutiérrez-Zufiaurre N, García-de Cruz S, Labayru-Echeverría C, López-Medrano R, et al. Tuberculosis por *Mycobacterium bovis* en la población de Castilla y León (España), 2006–2015. Enferm Infecc Microbiol Clin. 2019;37:19–24.

* Corresponding author.

E-mail address: tnebreda@saludcastillayleon.es (T. Nebreda-Mayoral).

Palabras clave:*Mycobacterium bovis*

Tuberculosis pulmonar

Tuberculosis extrapulmonar

Incidencia

Factores de riesgo

Tuberculosis por *Mycobacterium bovis* en la población de Castilla y León (España), 2006–2015**R E S U M E N**

Introducción: La incidencia anual de tuberculosis (TB) humana por *Mycobacterium bovis* ha disminuido considerablemente en los países industrializados desde inicios del siglo XX. El objetivo de este estudio fue conocer las características epidemiológicas, clínicas y microbiológicas de esta enfermedad en Castilla y León (CyL).

Métodos: Estudio retrospectivo de los casos de TB por *M. bovis* de CyL en un periodo de 10 años, comparando la epidemiología, los factores de riesgo y la evolución entre las formas pulmonares (TBP) y extrapulmonares (TBEP).

Resultados: Se recopilaron 75 casos de TB por *M. bovis*: 45 TBP y 31 TBEP. La incidencia acumulada de TB por *M. bovis* fue de 0,3 casos por 100.000 habitantes. Se mantuvo estable entre el primer y el segundo quinquenio (0,27 vs. 0,33, $p = 0,656$), a pesar del descenso de la incidencia global de la TB (13,58 vs. 10,71, $p < 0,0001$). La edad media fue de 66,2 ± 21,3 años, principalmente varones (63%) y nacidos en España (92%). TBP fue significativamente más frecuente en varones, mayores de 66 años, con inmunosupresión o fumadores. La mortalidad fue del 9%, asociada a la edad, a la inmunosupresión o a un tratamiento diferente al recomendado por la OMS.

Conclusiones: La incidencia de TB por *M. bovis* en CyL es superior a la de España y otros países europeos, y se mantuvo estable a pesar del descenso de la TB por MTC. Afectó mayoritariamente a población nacida en España que vivía en zonas rurales y con elevada media de edad.

© 2017 Elsevier España, S.L.U. y Sociedad Española de Enfermedades Infecciosas y Microbiología Clínica.
Todos los derechos reservados.

Introduction

Mycobacterium bovis, a member of the *Mycobacterium tuberculosis* complex (MTC), is a zoonotic pathogen that primarily infects cattle. Humans are infected by inhaling aerosols contaminated with the mycobacterium, consuming unpasteurised dairy products or due to continuous and close contact with infected animals.¹ Human-to-human transmission has occasionally been described both in immunocompetent and immunosuppressed patients.^{2,3} The MTC species are grouped into eight closely phylogenetically related lines. The clinical, radiological and pathological features of the main MTC species which cause tuberculosis (TB) in humans—*M. tuberculosis*, *Mycobacterium africanum* and *M. bovis*—are indistinguishable.¹ Nevertheless, the differentiation thereof is of great epidemiological and therapeutic interest. Historically, taxonomic separation was based on phenotype characteristics, antibiotic resistance, geographical distribution and host affinity,¹ but these have been replaced with faster, more accurate and more specific molecular methods.^{4,5}

The annual incidence of *M. bovis* cases in humans has decreased considerably in industrialised countries since the beginning of the 20th century, as a result of eradication campaigns in animals. Currently, it represents less than 2% of all TB cases, although there may be areas and populations in which this prevalence is greater.^{6–8} Most of the cases diagnosed are attributed to the reactivation of a latent infection acquired either before the pasteurisation era or in non-industrialised countries.^{6–11}

Bovine TB was once endemic in the autonomous community of Castile and León (CyL) but is now being eradicated thanks to the implementation of a nationwide bovine TB eradication programme since 1985.¹² The prevalence of infected herds has been reduced from 5.11% in 2006 to 1.93% in 2015.¹³ However, there is limited information on the significance and evolution of human TB caused by *M. bovis* in this community. This study's endpoint was to determine the microbiological, clinical and epidemiological characteristics of *M. bovis* infection in CyL based on pulmonary and extrapulmonary locations, as well as to compare it with epidemiological reports of TB due to MTC in this autonomous community.

Materials and methods

A retrospective study of all TB cases due to *M. bovis* collected in the laboratory information systems (LIS) of 14 public hospitals in CyL (autonomous community comprising 9 provinces and 14 Health Areas) between 2006 and 2015, both inclusive. One isolation was considered per patient. Patients with *M. bovis* BCG TB were excluded. Sample cultures were performed at each hospital. Phenotype and/or genotype identification and studies on antibiotic sensitivity to first-line anti-tuberculosis drugs were carried out in the reference laboratories of each Health Area. Fifteen strains (20%) were identified using GenoType MTBC hybridisation probes (HAIN Lifescience, Germany). Forty-four strains (59%) were identified at the Spanish National Mycobacteria Centre (Instituto de Salud Carlos III, Majadahonda, Madrid, Spain) using the genotype method described by Rodríguez et al.,⁷ based on the amplification of the *gyrB* gene and the detection of region of difference 1 (RD1), as well as the detection of the C169G mutation (H57D) in the *pncA* gene as an additional differentiating element for *M. bovis* and *Mycobacterium caprae*. In 16 strains (21%), mono-resistance to pyrazinamide (PZA) was considered the main identification criterion for *M. bovis*, alongside other morphological and biochemical characteristics unique to the species.¹

The medical histories of all patients with TB due to *M. bovis* were reviewed. Demographic, bacteriological and clinical data were gathered, as well as data related to risk factors, treatment and outcome, in a database designed for this study. Moreover, the total number of confirmed TB cases was quantified by Health Area in order to estimate the proportion of TB due to *M. bovis*.

Patients were grouped into two categories according to disease location: pulmonary (PTB) and extrapulmonary (EPTB). The latter category included patients with or without concomitant pulmonary involvement and pleural TB.

The risk factors included in the study were: (a) rural life, patients living in populations of less than 2000 inhabitants or who worked in agriculture; (b) animal contact, people in direct contact with cattle; (c) travel abroad, patients who come from or who have travelled to a TB endemic country; (d) immunosuppressive conditions, patients with end-stage kidney disease or who have received organ transplants or immunosuppressant treatments or who have other

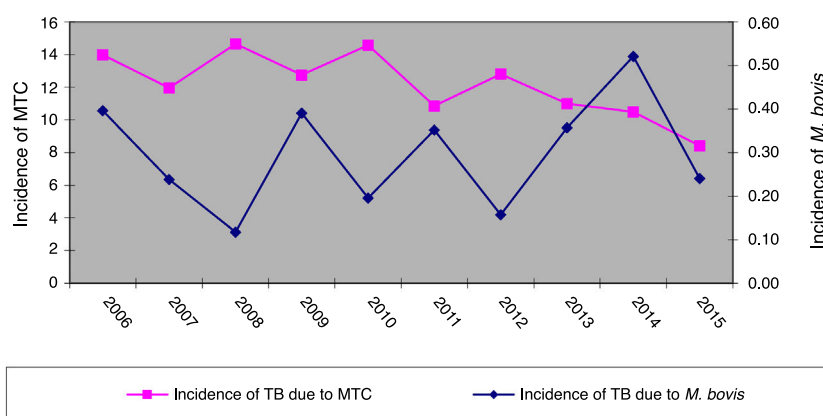


Fig. 1. Incidence of the *Mycobacterium tuberculosis* complex and *Mycobacterium bovis* in Castile and León, 2006–2015.

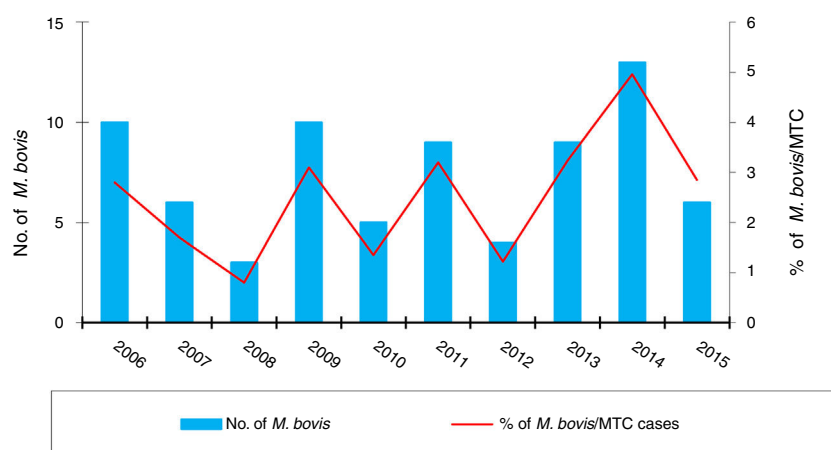


Fig. 2. Number of cases of tuberculosis due to *Mycobacterium bovis* and percentage in relation to *Mycobacterium tuberculosis* complex (MTC).

immunosuppressant conditions indicated in their medical history, excluding patients co-infected with HIV or diabetes mellitus; (e) HIV status; (f) diabetes mellitus; (g) smoker; and (h) no risk factors, patients who had none of the aforementioned risk factors.

Treatments were divided into two categories: (a) the treatment recommended by the World Health Organisation (WHO) for PZA-resistant MTC,¹⁴ consisting of two months of isoniazid (INH), rifampin (RIF) and ethambutol (ETB), followed by seven months of INH and two months of RIF and (b) others, which include combinations of INH, RIF, ETB and PZA, other combinations containing PZA or treatments which lasted less than or more than nine months.

Patient outcomes were initially recorded as resolution, death, relapse or unknown. These were then reclassified as resolution and death in order to study the influence of various risk factors on outcome.

The results were analysed using the SPSS programme (Statistical Package for Social Sciences, version 20.0, SPSS Inc., Chicago, IL, USA). Qualitative variables were analysed using the χ^2 test or Fisher's exact test, and quantitative variables using the Student's *t*-test. Differences were considered statistically significant if *p* was <0.05.

Results

Over the 10 years included in the study, 3080 cases of TB were confirmed in CyL, of which 75 (2.43%) were due to *M. bovis* and 3 (0.10%) to *M. caprae*. The overall incidence of TB decreased significantly between the first and second five-year periods (13.58 vs. 10.71 cases per 100,000 inhabitants, *p* < 0.0001). However, the

incidence attributed to *M. bovis* remained stable (0.27 vs. 0.33 cases per 100,000 inhabitants, *p* = 0.656) (Fig. 1). As a result, the proportion of TB due to *M. bovis* in relation to the total number of MTC cases showed a growing trend across both five-year periods (1.8 vs. 2.85%, *p* = 0.001). This was more due to a decrease in MTC TB than to an increase in TB attributed to *M. bovis* (Fig. 2).

Fig. 3 shows the number of cases of TB due to *M. bovis* and the cumulative incidence per province, which was significantly higher in westerly provinces (León, Zamora and Salamanca; *p* = 0.031).

PTB was predominant in this population, accounting for 44 cases (59%). Of the 31 patients who presented with EPTB, 5 also had PTB. Disseminated (26%) and lymphatic TB (26%) were the most common forms of EPTB. Others included bone and/or joint TB (5 cases), cutaneous TB (4 cases), genitourinary TB (2 cases), pleural TB (one case) and digestive TB (one case).

The microbiological, clinical and epidemiological characteristics of TB due to *M. bovis* are shown in Table 1, based on the location of the disease. PTB was significantly more common in men and in patients with a higher mean age. There were only two cases of childhood TB, where both children had been born in Spain and travelled to their parents' country of origin (Morocco). Most patients had been born in Spain (92%) and the rest came from Morocco (3 cases), Romania (one case), Thailand (one case) and "unspecified" (one case). No association was found between disease location and the patient's country of origin (*p* = 0.687). Patients with more bacilliferous forms presented with PTB (*p* = 0.006).

The most common risk factors in these patients were living in a rural environment (72%) and having direct contact with cattle

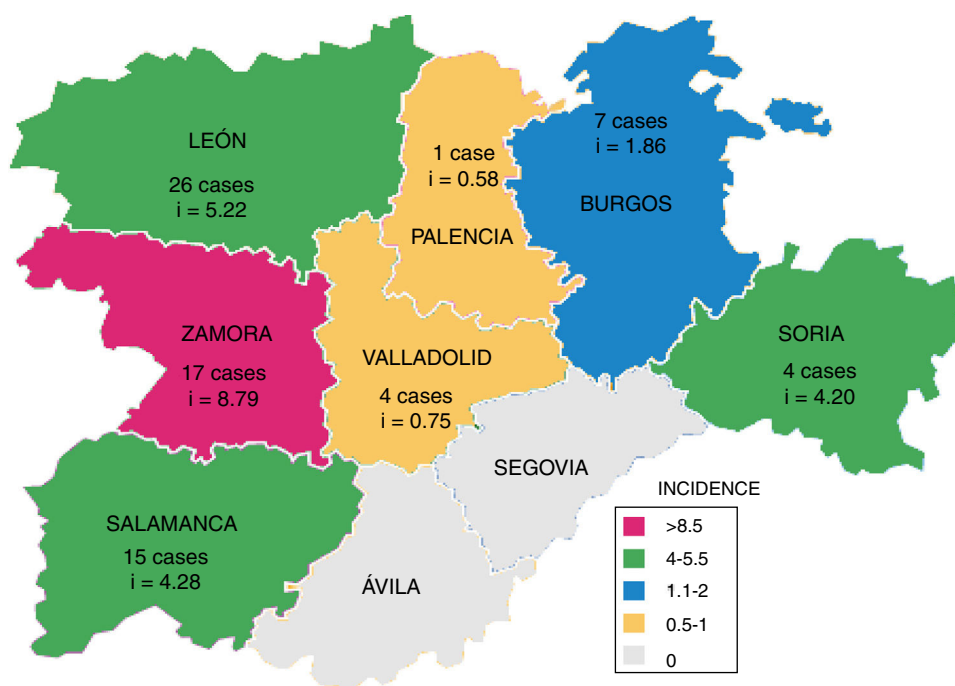


Fig. 3. Number of cases and cumulative incidence of human *Mycobacterium bovis* infections per province in Castile and León, 2006–2015.

(28%). The predisposition to PTB was greater in patients who smoked and in those who had immunosuppressive conditions.

Seventy-four of the *M. bovis* strains were mono-resistant to PZA and one strain presented INH and PZA resistance. Only 19 patients (27%) received the treatment recommended by the WHO, which was more common among patients with EPTB ($p=0.011$) and in those who recovered ($p<0.001$). Forty-one patients (63%) presented with satisfactory outcomes with a treatment that either included PZA in the first two months and/or did not have a duration of nine months. We were unable to determine the outcome of seven patients (10%) due to either a change in Health Area or failure to attend the review.

The overall mortality of these patients was 9% (Table 2): four patients died before starting the treatment and three during treatment. Mortality was higher in males, although no significant differences were found. The deceased patients had a mean age that was higher than those who recovered and immunosuppressive conditions were also significantly more prevalent. The underlying diseases of these five deceased patients were: diabetes (one case), end-stage kidney disease (one case), rectal cancer (one case), haematological malignancy (one case) and severe COPD treated with corticosteroids (one case).

Discussion

In the 10 years between 2006 and 2015, the mean annual incidence of confirmed *M. bovis* TB cases in CyL was higher than that of Spain and other European countries.¹⁵ Likewise, the proportion of *M. bovis* infections in relation to the total number of TB cases (2.43%) was also higher than in Asturias (1.4% in 2006–2014),¹⁶ Spain overall (1.9% in 2004–2007),⁷ the Netherlands (1.4% in 1993–2007)⁶ and the United Kingdom (1.5% in 2003).¹⁰ The risk of zoonotic TB in CyL may be greater than in other regions, since it is an important livestock-producing region and 26.4% of the population live in rural areas.

In our population, as in other regions with low TB rates, the efforts aimed at eradicating TB have led to a fall in *M. tuberculosis* TB. However, they have not managed to reduce *M. bovis* infections,

indicating that these two species behave in different ways.^{8,17} The fact that other domestic and wild animals may be a source of infection for humans could also have an impact in this regard.^{1,10,18}

The greatest incidence of human *M. bovis* infections were found in the most westerly provinces of the community, while the greatest prevalence of herds infected with bovine TB were located in the south of CyL.^{12,13} This disparity may be due to differences in how the disease was detected in each Health Area.

The predominance of PTB is a finding that can be compared to other populations in our setting,^{7,10,11} with the exception of the Netherlands, where EPTB accounted for 59% of the cases.⁶ Historically, *M. bovis* infections were primarily located in the digestive and lymphatic systems. The reduced incidence of EPTB could be due to the fact that air is the main mode of transmission now that milk pasteurisation has become widespread,^{10,17} as well as greater difficulties in diagnosis, which ultimately reduces clinical suspicion until the advanced stages of the disease.^{19,20}

EPTB was more common in women, as in other studies on TB caused by *M. bovis*^{7,16} or *M. tuberculosis*.²⁰ The reasons for this are not well known, although some authors have linked it to socioeconomic factors, different infection pathways in each gender, hormonal factors and cellular immunity.^{6,21}

The advanced age of the patients and the presence of immunosuppressive conditions suggest the reactivation of an old infection which occurred prior to the pasteurisation era. Thus, in regions where bovine TB has been virtually eradicated, the residual cases appearing in older patients are associated with the reactivation of a dormant infection, while TB occurring in younger patients is considered a primary infection.^{9,21} However, in the study by Palacios et al.¹⁶ 47% of the genotypes in human strains were shared by cattle, suggesting a recent transmission irrespective of age. The most common spoligotypes in the bovine TB strains of CyL are SB0121, SB0134 and SB0339.²² The first two have been identified in both humans and cattle in Asturias¹⁶ and England.¹¹ Unfortunately, the lack of spoligotyping among our strains did not allow us to draw conclusions in this regard.

Diabetes mellitus has been strongly correlated with PTB,²⁰ and HIV with EPTB.²³ However, this association was not evident in our

Table 1Characteristics of pulmonary and extrapulmonary *Mycobacterium bovis* infections in Castile and León, 2006–2015.

Variable	Pulmonary TB (n = 44) n (%)	Extrapulmonary TB (n = 31) n (%)	Total (n = 75) n (%)	p
Gender				0.009
Male	33 (77)	14 (45)	47 (63)	
Age groups, years				0.011
0–19	0	2 (6)	2 (3)	
19–45	6 (14)	8 (26)	14 (17)	
46–65	4 (9)	8 (26)	16	
>66	34 (77)	13 (42)	47 (63)	
Age, mean + SD	72.7 + 17.0	56.9 + 23.6	66.16 + 21.32	0.018
Country of birth				0.653
Spain	41 (93)	28 (90)	69 (92)	
Immigrant	3 (7)	3 (10)	6 (8)	
Sample bacilloscopy				0.006
Positive	24 (55)	7 (23)	31 (41)	
Negative	20 (45)	24 (77)	44 (59)	
Sample CRP (n = 18)				0.693
Positive	8 (88%)	8 (88%)	16 (89)	
Negative	1 (12%)	1 (12%)	2 (11)	
Risk factors				
Contact with cattle	12 (27)	9 (29)	21 (28)	0.798
Rural life	33 (75)	21 (67)	54 (72)	0.634
Immunosuppression	17 (40)	3 (10)	20 (27)	0.006
Travel to an endemic country	3 (7)	5 (16)	8 (11)	0.180
Diabetes mellitus	4 (9)	1 (3)	5 (7)	0.333
HIV positive	2 (5)	0	2 (3)	0.232
Smoking habit	13 (30)	1 (3)	14 (17)	0.003
None of the above	3 (7)	3 (10)	6 (8)	
Unknown	1 (2)	0	1 (1)	
Treatment^a				0.011
WHO recommended	7 (16)	12 (43)	19 (27)	
Other	29 (67)	16 (57)	45 (63)	
Unknown	7 (16)	0	7 (10)	
Outcome				0.463
Resolution	36 (82)	26 (84)	62 (83)	
Death	3 (7)	4 (13)	7 (9)	
Relapse	1 (2)	0	1 (1)	
Unknown	4 (9)	1 (3)	5 (7)	

^a Four patients died before receiving treatment (1 PTB and 3 EPTB).**Table 2**Outcome of patients with tuberculosis due to *Mycobacterium bovis*.

	Resolution (n = 63)	Death (n = 7)	p
Gender			0.187
Male	38 (60)	6 (86)	
Age, mean + SD	64.3 + 22.2	78.2 + 11.6	0.030
Immunosuppression	12 (19)	5 (71)	0.002
Treatment^a			<0.001
WHO	16 (25)	0 (0)	
Other	41 (65)	3 (43)	

^a Four patients died before receiving treatment. The outcome of five patients is unknown.

study, probably due to the limited number of patients with these risk factors.

Although 63% of our patients were not treated as per the WHO recommendation, they had satisfactory outcomes. The low resistance of MTC to first-line anti-tuberculosis drugs (less than 6%^{24,25}) could explain the patients' improvement, despite delayed microbiological identification. Quickly and reliably identifying the mycobacterium species and the immediate notification of the clinician could improve these results.

The mortality of the disease was lower than in similar studies.^{6,16} This was probably related to multiple factors,^{6,10,16} such as advanced age and/or immunosuppression.

In comparison to TB due to *M. bovis*, the patients with MTC TB in CyL in 2015²⁵ had a lower mean age (52.7 + 24.5) and a higher percentage of immigrants (17.5%), childhood TB (6.7%) and pulmonary forms (73%). The main risk factor was contact with a TB-positive patient (13.41%). The provinces with the highest incidences were Palencia, León and Zamora.

One of the limitations of this study was the lack of knowledge on patient chest X-ray patterns, as well as the disease resolution time.

In conclusion, the incidence of *M. bovis* TB in CyL remained stable despite the reduction in MTC TB and the fact that bovine TB is undergoing eradication. The disease mostly affected the population that had been born in Spain who had a higher mean age. The pulmonary form was the most prevalent. Detecting *M. bovis* genotypes in the population would help to detect sources of infection and prevent the transmission thereof. In our opinion, close collaboration between human and animal health is necessary in order to control and eradicate this disease.

Authorship

T. Nebreda Mayoral wrote the manuscript, which was reviewed by the co-authors. T. Nebreda Mayoral, M.F. Brezmes Valdivieso, N. Gutiérrez Zufiaurre, S. García de Cruz, C. Labayru Echeverría, R. López Medrano, L. López-Urrutia Lorente, A. Tinajas Puertas and O.

Rivero Lezcano took part in designing the study and analysing the data.

Conflicts of interest

We declare that we have no conflicts of interest.

Acknowledgements

We would like to thank Dr M.S. Jiménez (Spanish National Mycobacteria Centre, Instituto de Salud Carlos III, Majadahonda, Madrid, Spain) for her help in identifying *Mycobacterium bovis*. And thank you to all other members of the GRUMICALE (*Grupo de Micobacterias de Castilla y León* [Castile and León Mycobacteria Group]): B. Nogueira González, Hospital Clínico Universitario de Valladolid, Valladolid; R. Sánchez Arroyo, Hospital de Ávila, Ávila; S. Hernando-Real, Hospital de Segovia, Segovia; O. Rivero-Lezcano, Complejo Asistencial Universitario de León, León; B. Ullivarri Francia, Hospital Santiago Apóstol, Miranda de Ebro-Burgos; R. Rodríguez-Tarazona, Hospital Santos Reyes, Aranda de Duero-Burgos; I. Antolín Ayala, Hospital de Medina del Campo, Medina del Campo-Valladolid, for their interest in searching for *Mycobacterium bovis* cases and providing details on the TB cases from their Health Areas.

References

- Grange JM. *Mycobacterium bovis* infection in human beings. *Tuberculosis* (Edinb). 2001;81:71–7.
- Evans JT, Smith EG, Banerjee A, Smith RMM, Dale J, Innes JA, et al. Cluster of human tuberculosis caused by *Mycobacterium bovis*: evidence for person-to-person transmission in the UK. *Lancet*. 2007;369:1270–6.
- Rivero A, Márquez M, Santos J, Pinedo A, Sánchez MA, Esteve A, et al. High rate of tuberculosis reinfection during a nosocomial outbreak of multidrug-resistant tuberculosis caused by *Mycobacterium bovis* strain B. *Clin Infect Dis*. 2001;32:159–61.
- Niemann S, Harmsen D, Rusch-Gerdes S, Richter E. Differentiation of clinical *Mycobacterium tuberculosis* complex isolates by *gyrB* DNA sequence polymorphism analysis. *J Clin Microbiol*. 2000;38:3231–4.
- Richter E, Weizenegger M, Fahr AM, Rüsche-Gerdes S. Usefulness of the GenoType MTBC assay for differentiating species of the *Mycobacterium tuberculosis* complex in cultures obtained from clinical specimens. *J Clin Microbiol*. 2004;42:4303–6.
- Majoor CJ, Magis-Escarra C, van Ingen J, Boeree MJ, van Soolingen D. Epidemiology of *Mycobacterium bovis* disease in humans, the Netherlands, 1993–2007. *Emerg Infect Dis*. 2011;17:457–63.
- Rodríguez E, Sánchez LP, Pérez S, Herrera L, Jiménez MS, Samper S, et al. Human tuberculosis due to *Mycobacterium bovis* and *M. caprae* in Spain, 2004–2007. *Int J Tuberc Lung Dis*. 2009;13:1536–41.
- Gallivan M, Shah N, Flood J. Epidemiology of human *Mycobacterium bovis* disease, California, USA, 2003–2011. *Emerg Infect Dis*. 2015;21:435–42.
- Hardie RM, Watson JM. *Mycobacterium bovis* in England and Wales: past, present and future. *Epidemiol Infect*. 1992;109:23–33.
- De la Rua-Domenech R. Human *Mycobacterium bovis* infection in the United Kingdom: incidence, risks, control measures and review of the zoonotic aspects of bovine tuberculosis. *Tuberculosis* (Edinb). 2006;86:77–109.
- Stone MJ, Brown TJ, Drobnieswski A. Human *Mycobacterium bovis* infections in London and Southeast England. *J Clin Microbiol*. 2012;50:164–5.
- Ministerio de Agricultura, Alimentación y Medio Ambiente. Informe final técnico-financiero. Programa Nacional de la Tuberculosis Bovina, año 2015 [accessed 1 Feb 2017]. Available in: http://www.mapama.gob.es/es/ganaderia/temas/sanidad-animal-higiene-ganadera/informe.tb.2015_tcm7-428843.pdf.
- Ministerio, de agricultura y pesca, alimentación y medio ambiente 2017. Programa Nacional de Erradicación de Tuberculosis Bovina presentado por España para el año 2017 [accessed 15 Sep 2017]. Available in http://www.mapama.gob.es/es/ganaderia/temas/sanidad-animal-higiene-ganadera/pnetb.2017.3_tcm7-443753.pdf.
- Nahid P, Dorman SE, Alipanah N, Barry PM, Brozek JL, Cattamanchi A, et al. Official American Thoracic Society/Centers for Disease Control and Prevention/Infectious Diseases Society of America Clinical Practice Guidelines: treatment of drug-susceptible tuberculosis. *Clin Infect Dis*. 2016;63:147–95.
- European Food Safety Authority, European Centre for Disease Prevention and Control. The European Union summary report on trends and sources of zoonoses, zoonotic agents and food-borne outbreaks in 2014. EFSA J. 2015;13:4329. Available in: <http://ecdc.europa.eu/en/publications/Publications/zoonoses-trends-sources-EU-summary-report-2014.pdf>.
- Palacios JJ, Navarro Y, Romero B, Penedo A, Menéndez González Á, Pérez Hernández MD, et al. Molecular and epidemiological population-based integrative analysis of human and animal *Mycobacterium bovis* infections in a low-prevalence setting. *Vet Microbiol*. 2016;195:30–6.
- Rodwell TC, Moore M, Moser KS, Brodine SK, Strathdee SA. Tuberculosis from *Mycobacterium bovis* in binational communities, United States. *Emerg Infect Dis*. 2008;14:909–16.
- Dalovisio F, Dettler JR, Mikota-Well MS. Rhinoceros' rhinorrhea: cause of an outbreak of infection due to airborne *Mycobacterium bovis* in zookeepers. *Clin Infect Dis*. 1992;5:598–600.
- Sherman LF, Fujiwara PI, Cook SV, Bazerman LB, Frieden TR. Patient and health care system delays in the diagnosis and treatment of tuberculosis. *Int J Tuberc Lung Dis*. 1999;3:1088–95.
- Lin JN, Lai CH, Chen YH, Lee SS, Tsai SS, Huang CK, et al. Risk factors for extra-pulmonary compared to pulmonary tuberculosis. *Int J Tuberc Lung Dis*. 2009;13:620–5.
- Noertjojo K, Tam CM, Chan SL, Chan-Yeung MM. Extrapulmonary and pulmonary tuberculosis in Hong Kong. *Int J Tuberc Lung Dis*. 2002;6:879–86.
- Rodríguez S, Romero B, Bezos J, de Juan L, Alvarez J, Castellanos E, et al. High spoligotype diversity within a *Mycobacterium bovis* population: clues to understanding the demography of the pathogen in Europe. *Vet Microbiol*. 2010;141:89–95.
- Slutsker L, Castro KG, Ward JW, Dooley SW. Epidemiology of extrapulmonary tuberculosis among persons with AIDS in the United States. *Clin Infect Dis*. 1993;16:513–8.
- López-Medrano R, Nebreda-Mayoral T, Brezmes-Valdivieso MF, García-de Cruz S, Nogueira-González B, Sánchez-Arroyo R, et al. Contribution of microbiology in the diagnosis of tuberculosis in Castile and León (Spain): findings of the GRUMICALE 2013 study. *Enferm Infecc Microbiol Clin*. 2017. <http://dx.doi.org/10.1016/j.eimc.2016.11.009>.
- Informe epidemiológico sobre la tuberculosis en Castilla y León. Año 2015. Servicio de Epidemiología. Dirección General de Salud Pública. Consejería de Sanidad [accessed 15 Oct 2017]. Available in: <https://www.saludcastillayleon.es/profesionales/es/informacion-epidemiologica/enfermedades-infecciosas/tuberculosis>.