Evaluation of susceptibility to antimycobacterial drugs in *Mycobacterium tuberculosis* complex strains isolated from cattle in Poland

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**Abstract**

**Introduction:** Tuberculosis is a highly infectious disease affecting both humans and animals. It is caused by the *Mycobacterium tuberculosis* complex (MTBC) – *Mycobacterium bovis* and *Mycobacterium caprae*, which are aetiological factors of bovine tuberculosis (bTB). In Poland, the bTB eradication programme exists. Animals diagnosed with tuberculosis are in the majority of cases not treated, but removed from their herd and then sanitary slaughtered. **Material and Methods:** In total, 134 MTBC strains isolated from cattle in Poland were subjected to microbiological analysis. The resistance phenotype was tested for first-line antimycobacterial drugs used in tuberculosis treatment in humans: streptomycin, isoniazid, rifampicin, ethambutol, and pyrazinamide. The strains were isolated from tissues collected post mortem, so the test for drug resistance fulfilled only epidemiological criterion. **Results:** The analysis of drug-resistance of MTBC strains revealed that strains classified as *M. bovis* were susceptible to 4 antimycobacterial drugs: isoniazid, rifampicin, streptomycin, and ethambutol, and resistant to pyrazinamide. The strains classified as *M. caprae* were sensitive to all tested drugs. **Conclusion:** The results indicate that despite enormously dynamic changes in mycobacterial phenotype, Polish strains of MTBC isolated from cattle have not acquired environmental resistance. The strains classified as *M. bovis* are characterised by natural resistance to pyrazinamide, which is typical for this species.

**Keywords:** cattle, bovine tuberculosis, *Mycobacterium tuberculosis* complex, antimycobacterial drugs, Poland.

Introduction

Tuberculosis is a highly infectious disease, affecting both humans and animals, caused by the *Mycobacterium tuberculosis* complex (MTBC) – *Mycobacterium bovis* and *Mycobacterium caprae*, which are aetiological factors of bovine tuberculosis (bTB). In Poland, the bTB eradication programme exists. Animals diagnosed with tuberculosis are in the majority of cases not treated, but removed from their herd and then sanitary slaughtered. According to the classification of the World Health Organisation (WHO), bTB is rated as direct zoonosis, involving direct transmission of an infectious agent from an infected animal to a sensitive animal (human), without participation of an indirect host. Due to the direction of infectious agent transmission, bovine TB can be also rated as reverse zoonosis.
not treated, but removed from their herd and then sanitary slaughtered. The only described case of tuberculosis treatment in animals in Poland regarded a 10-year-old male giraffe kept in the Silesian Zoological Garden in Chorzów (20).

First results of the research on resistance to streptomycin (SM), isoniazid (INH), rifampicin (RMP), and ethambutol (EMB) – S.I.R.E. and pyrazinamide (PZA) of MTBC strains isolated from tissues of different animal species were published in Poland in 2011. This study is a continuation of research on MTBC strains isolated from cattle.

Material and Methods

MTBC isolates. Research material, consisting of 134 MTBC strains isolated from domestic cattle (Bos taurus), was collected from 10 provinces: West Pomeranian, Masovian, Podlasie, Lesser Poland, Pomeranian, Lower Silesian, Silesian, Łódź, Holy Cross, and Greater Poland. The strains were isolated from animal tissues retrieved post mortem in such way that estimation of drug resistance in this task met only the epidemiological criterion. The isolates were cultured on the Stonebrink medium. Affiliation to MTBC was confirmed by immunochromatographic test for detection of MPT64 protein fraction produced by MTBC cells within the culture (Identification Test MGIT TBC®, Becton and Dickinson, USA). Species identification was conducted with GenoType MTBC test (Hain Lifescience, Germany), which allowed to classify 103 strains as Mycobacterium bovis and 31 strains as Mycobacterium caprae.

The Bactec method for determination of S.I.R.E. Test for resistance to S.I.R.E. was done with the use of Bactec MGIT 960 and Middlebrook 7H10 Agar (Bactec MGIT 960 SIRE Kits, Becton, Dickinson, USA) at the concentration limit of 1.0 μg/mL for SM, 0.1 μg/mL for INH, 1.0 μg/mL for RMP, and 5.0 μg/mL for EMB. Analysis of fluorescence in test tubes containing drugs compared with fluorescence in control test tubes allowed to define sensitivity.

The Bactec method for determination of PZA. The estimation of drug resistance to PZA was carried out on the isotopic Bactec 960-Tb system (27), Middlebrook’s liquid culture medium 7H12 containing palmitic acid 14C, at pH 5.9–6.0, with PZA concentration limit of 100 μg/mL (Bactec MGIT 960 PZA Kit, Becton, Dickinson, USA). Growing bacteria metabolise the substrate present in the culture medium up to 14CO2. The amount of produced 14CO2 is directly proportional to the intensity of population growth in culture medium. The addition of PZA to culture media results in cessation of M. caprae growth. This is reflected by a decrease in amount of 14CO2 in relation to the control, in which the amount of 14CO2 and growth index increase systematically. In the case of resistant strains there is little or no difference in the growth index between the control and an analysed strain (26).

Results

In the group of 103 bovine bacillus strains identified as M. bovis, all strains demonstrated susceptibility to four basic antimycobacterial drugs: INH, RMP, SM, and EMB, and were resistant to PZA (Fig. 1). In the group of 31 strains identified as M. caprae, all strains demonstrated susceptibility to PZA and to INH, RMP, SM, and EMB. These results indicating no drug resistance except for PZA were identical to our findings of 2013 when 58 cattle isolates were examined (19).

Discussion

All confirmed M. caprae strains were susceptible to all examined drugs. Data regarding drug susceptibility of M. caprae TB in humans and animals is limited. Strains classified as M. bovis are marked by a feature which is unique for this type, namely their natural resistance to PZA, which was proved in the study. These results indicating no drug resistance, except to PZA, were identical with research by Daly et al. (12) and Fitzgerald et al. (13). The material for their study consisted of 100 strains of M. bovis: 91 from deer, 4 from coyotes, 3 from cattle, and 2 from raccoons. The obtained results are similar due to the lack of antimycobacterial therapy which, if applied to either cattle or wildlife species, would cause them to develop environmental resistance. Resistance of cattle M. bovis strains to INH and RMP was reported in Northern Italy (7). Fifteen out of 61 tested strains were resistant to either INH or RMP, or both drugs, showing multidrug resistance (MDR). Detection of resistant cattle M. bovis strains appears of great interest.

The literature describes also MDR M. bovis strains from human clinical samples (33). This type of resistance is mainly observed in countries with a high prevalence of HIV infection (4). The emergence of
drug-resistant *M. bovis* is an important public health problem that jeopardises the success of human TB control programmes.

The incidence rate of bovine TB among people in Europe is higher than in last few years. That is why establishing the phenotype of resistant MTBC strains retrieved from animals on anti-tubercle bacilli and TB treatment in humans seems to be justified in the public health aspect. Data pertaining to incidence of bovine tuberculosis among humans in 2014, which were presented in the EFSA report, were gathered from 27 EU members (excluding France) and Iceland, Norway, and Switzerland. As many as 134 incidents of *M. bovis* were confirmed, among which the highest number occurred in Germany – 45 people, next in UK – 29, and in Spain – 25 (28).

The WHO estimates that in the countries where programmes against TB were insufficiently monitored, 1% of TB incidence is caused by tubercle bacilli. So far, there have been two incidents of bovine TB among people in Poland, specifically in the south of the country (16). According to the EFSA report, the main source of *M. bovis* in humans is the milk originating from infected cows.

It should be emphasised that wildlife, especially infected wild animals living in the vicinity of households (9), is considered one of the main reservoirs of these microorganisms which may be transmitted directly and indirectly to livestock (common pasture) and even humans. Although the biggest natural reservoir of *M. bovis* in Poland is sick cattle (17), the main source of tuberculosis in Spain is the wild boar (3), and the reservoir of *M. bovis* in England is the population of badgers (14). There was one incident noted in the area of Cornwall (England) where a veterinary nurse had a pulmonary *M. bovis* TB infection. Prior to that, the patient assisted in tuberculin tests of cattle herds and had contact with badgers in the past (reservoir of *M. bovis*). Four months later, the dog in the patient’s household displayed pulmonary symptoms of bTB. The genotype of isolated *M. bovis* strains was identical, which allowed to establish the common source of infection (31).

*M. bovis* can cause tuberculosis in other livestock species as well. It is of high virulence under natural conditions among goats (11), pigs (2, 5), sheep (25), and cats (23), whereas it is lower in horses (24) and dogs (32).

Cattle demonstrates low susceptibility to tubercle bacillus, but some incidents of *M. tuberculosis* among cattle were reported as well (18). The most essential components of modern programmes that help to reduce tuberculosis incidence are primarily its early diagnosis (21, 30) and appropriately selected anti-tuberculosis treatment (22). The biggest obstacle in carrying out programmes against tuberculosis all over the world is the ever-increasing occurrence of the bacillus drug resistance phenomenon as well as the absence of new drugs (7) since the invention of RMP method in 1965. Drug-resistant tuberculosis, especially its specific MDR strains and extensively drug resistant (XDR) strains, is a disease of great mortality, with the risk especially increased in people diagnosed with HIV (21, 34).

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