(Romania)

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Abstract

Study Design: Case report.

Objective: To characterise the paleopathology presented in the skeleton of a 45-50-year-old male indicative of tuberculous spondylitis and to confirm by the detection of ancient DNA. Summary of Background Data: Tuberculosis (TB) is an infectious disease prevalent in both present and ancient human populations. The disease is primarily located within the lungs, so although characacteristic bone lesions can lead to a clear diagnosis, skeletal TB occurs in only 5-6% of TB infections, even in historical cases. In addition, the visual appearance of human skeletal remains may be influenced by the environmental conditions at the burial site. However, it is important to recognise ancient skeletal TB, because this can provide important data on the history of Mycobacterium tuberculosis and gives an unique opportunity for physicians to observe the natural outcome of the infection from the pre-antibiotic era.

Methods: Paleopathological analysis was carried out using careful visual observation supported by ancient DNA analysis. Approximately 60 mg of bone powder from rib fragments was examined and DNA from the *M. tuberculosis* complex was detected by PCR targeting specific genetic loci of the IS6110 and IS1081 regions.

Results: The skeleton is part of a human osteoarchaeological collection (n=274) from the 12th-13th century Transylvanian archaeological site of Peteni, in modern-day Romania. The individual, a 45-50-year-old male, showed gross pathology typical of tuberculous spondylitis. The paleopathological diagnosis was supported by analysis for *M. tuberculosis* complex ancient DNA.

Conclusions: This case demonstrates that TB was present in Transylvania (Romania) during the 12-13th century and adds to the growing body of knowledge on the history of this disease. **Key Words:** ancient DNA;Mycobacterium tuberculosis complex, paleopathology, PCR, skeletal tuberculosis,Transylvania (Romania)

Mini Abstract/Précis.

The skeleton of a 45-50-year-old male from the 12-13th century in Transylvania, present-day Romania, demonstrated paleopathology typical of spinal tuberculosis. The morphological diagnosis was supported by the detection of ancient DNA specific for the *Mycobacterium tuberculosis* complex.

Key Points:

Spinal tuberculosis was present during the 12-13th century in Transylvania.

The presented case was examined morphologically and supported by the detection of ancient DNA from the *Mycobacterium tuberculosis* complex.

Introduction

Tuberculosis (TB) is an infectious disease caused by members of the *Mycobacterium tuberculosis* complex. The infection is normally located within the lungs. However, the infection may become generalised and affect the skeleton, mainly the thoracic and the lumbar sections of the spine, the great joints, the costae and the endocranial surface. Skeletal TB is, with rare exceptions, the result of limited hematogenous spread.¹

The appearance of ancient TB in Europe, including Hungary and the whole Carpathian Basin, has been the subject of several previous investigations.²⁻⁶ From Romania only four cases have been published which describe bone alterations suggestive of TB, from the Medieval cemetery of Sibiu-Piaţa Huet.⁷

The purpose of the current study is to present a new skeletal TB case from Transylvania, in present-day Romania.

Material and methods

A human osteoarchaeological collection (n=274) dated to the 12-13th century, from the Transylvanian archaeological site of Peteni,⁸ served as a source of material. The age at death and the morphological sex was estimated based on the method of Acsádi and Nemeskéri.⁹ Grave 107 contained the skeleton of a 45-50-year-old male which showed possible

pathological changes. A paleopathological visual analysis was followed up with ancient DNA (aDNA) analysis for the *Mycobacterium tuberculosis* complex.

Molecular analysis of DNA for the Mycobacterium tuberculosis complex

Rib fragments were used for DNA extraction. The recommended protocols for aDNA were followed and approximately 60 mg of bone powder was examined and DNA extracted.^{5,10} PCR was used to amplify any DNA from specific regions of the multicopy IS6110 and IS1081 regions of the *M. tuberculosis* complex. Amplified DNA was examined by agarose gel electrophoresis and confirmed by sequencing.^{10,11}

Results

Paleopathological assessment

Cervical (C) spine: From C1 to C6 all vertebrae are absent post-mortem. A 8-9 mm wide probable lytic hole can be observed on the superior surface of the corpus of C7.

Thoracic (T) spine: T1, T2 and T6 are absent post-mortem. The superior surface of T3 is porotic and pitted. There are two small pits on the superior surface of T4. The corpus is porotic and the inferior surface is eroded (Fig. 1). The inferior surface of T5 is pitted. One-quarter of the inferior surface of T7 is missing due to lytic destruction. This reduced the height of the corpus so the corpus has started to collapse (Fig. 2). The superior surface of T8 and the inferior surface of T11 are pitted. There is one major lytic hole on the inferior surface of T9.

Lumbar (L) spine: Slight spondylosis deformans can be seen on L1 – the inferior surface of the corpus is pitted (Fig. 3). Osteophyte formation can be seen on the right side of L2 and the height of the corpus is reduced. The inferior surfaces of the corpus of L2 and L3 are pitted. Besides possible post-mortem changes, large pre-mortem perforations can be seen on the corpus of L3. The height reduction of the corpus is major. There are more cavities in the corpus of L4 (Fig. 4). The cavities are smaller on the inferior surface than the superior surface

and also the height of the corpus is reduced. The superior surface of the corpus of L5 is eroded by smaller pits and a major cavity has formed. Osteophyte formation appears between the left side of the corpus of L5 and the sacrum.

Ribs: No active periosteal reactions were observed on the ribs. However, on the costal groove and on the inner surfaces of the ribs, disseminated small new bone formations (probably healed pleuritis/periosteal appositions) can be seen (Fig. 5).

Endocranial surfaces, pelvic girdle and the bones of the lower limbs: No pathological changes consistent with any possible infectious diseases were observed on these bones.

Diagnosis: Possible spinal TB. However, other infectious processes cannot be excluded.

Molecular examination

Positive PCR results were obtained using primers for IS6110 (123-bp) and IS1081 (113-bp) (Fig. 6). The IS1081 amplified DNA was sequenced (Figs. 7A-B) which confirmed that it was from the *M. tuberculosis* complex.

Conclusions

Ancient skeletal TB has been described previously in the Carpathian Basin.^{5,6} Based on morphological and molecular results, this Transylvanian case from the Medieval Period, in present-day Romania, contributes to our knowledge on the paleopathology of spinal TB and its occurrence in the Carpathian Basin in the past.

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Figure legends

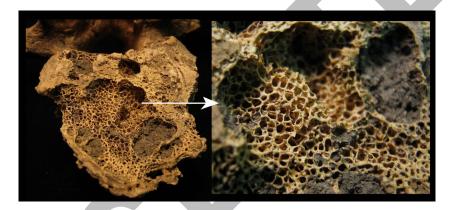


Figure 1. The inferior surface of the corpus of the 4th thoracic vertebra is eroded by lytic holes.



Figure 2. The height of the corpus of the 7th thoracic vertebra is reduced and a collapse started to form.



Figure 3. The inferior surface of the corpus of the first lumbar vertebra is pitted.



Figure 4. More cavities can be seen on the inferior surface of the 4th lumbar vertebra (the photo was taken by Elek Benkő).



Figure 5. Disseminated small new bone formations can be observed on the costal groove and on the inner surface of the ribs.

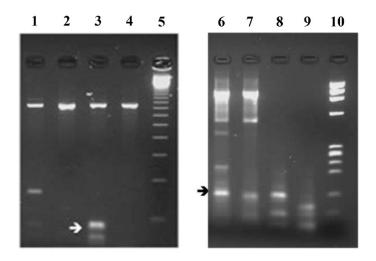


Figure 6. Agarose gel electrophoresis of PCR amplicons from the *M. tuberculosis* complex-specific loci IS6110 (on the left side) and IS1081 (on the right side). The gels were loaded as follows (left to right): lanes 1–3, different fractions of DNA extract from Peteni 107 (see white arrow); lane 4, negative extraction control; lane 5, molecular markers, lanes 6-9, different fractions of DNA extract from Peteni 107 (see black arrow), lane 10, molecular markers.

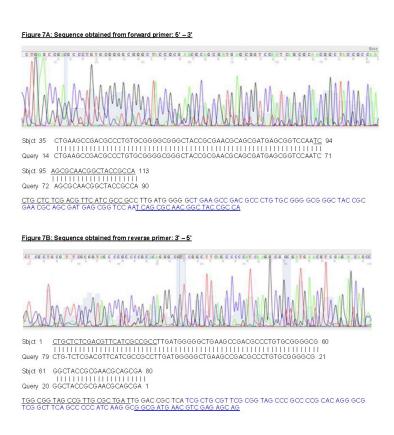


Figure 7. DNA sequences obtained from the *M. tuberculosis* complex specific region of IS*1081*. For the forward (Fig. 7A) and reverse (Fig. 7B) primers the following data are shown: Top: the actual sequence (electropherogram); Centre: the experimental sequence obtained by PCR aligned with the corresponding sequence in the NCBI nucleotide database (http://www.ncbi.nlm.nih.gov/nucleotide/); Bottom: the forward and reverse sequence for the target locus showing the primers (underlined) and the bases sequenced from amplified DNA (blue text).