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Troglostrongylus brevior: a new parasite for Romania

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Abstract

Background: The genus *Troglostrongylus* includes nematodes infecting domestic and wild felids. *Troglostrongylus brevior* was described six decades ago in Palestine and subsequently reported in some European countries (Italy, Spain, Greece, Bulgaria, and Bosnia and Herzegovina). As the diagnosis by the first-stage larvae (L1) may be challenging, there is a possibility of confusion with *Aelurostrongylus abstrusus*. Hence, the knowledge on the distribution of this neglected feline parasite is still scarce. The present paper reports the first case of *T. brevior* infection in Romania. In July 2017, a road-killed juvenile male *Felis silvestris*, was found in in Covasna County, Romania. A full necropsy was performed and the nematodes were collected from the trachea and bronchioles. Parasites were sexed and identified to species level, based on morphometrical features. A classical Baermann method was performed on the lungs and the faeces to collect the metastrongyloid larvae. Genomic DNA was extracted from an adult female nematode. Molecular identification was accomplished with a PCR assay targeting the ITS2 of the rRNA gene.

Results: Two males and one female nematodes were found in the trachea and bronchioles. They were morphologically and molecularly identified as *T. brevior*. The first-stage larvae (L1) recovered from the lung tissue and faeces were morphologically consistent with those of *T. brevior*. No other pulmonary nematodes were identified and no gross pulmonary lesions were observed.

Conclusions: This paper represents the first report of *Troglostrongylus brevior* infection in Romania, so far representing the second northernmost location for this genus in Europe. The diversity of species infecting wild and domestic felids and the differences regarding the clinical significance of these nematodes highlight the need for a more intense surveillance and proper diagnosis of feline lungworm infections, especially in countries where more species were demonstrated to be present. Furthermore, an increased awareness between clinicians is needed for a correct diagnostic approach to feline lungworm diagnosis.

Keywords: Troglostrongylus brevior, Felis silvestris, Romania

Background

A wide variety of nematode species belonging to the family Metastrongyloidea are known to infect domestic and wild felids. Among them, *Aelurostrongylus abstrusus* (Strongylida: Angiostrongylidae) has been considered for a long time to be the only metastrongyloid species parasitic in bronchi, bronchioles and alveoli of felids [1]. However, other species, such as *Troglostrongylus brevior*, *T. subcrenatus* (Strongylida: Crenosomatidae) and *Oslerus rostratus* (Strongylida: Filaroididae) have been

reported more recently to cause respiratory diseases in cats [2–4]. The genus *Troglostrongylus* includes nematodes infecting domestic cats (*Felis silvestris catus*), wildcats (*Felis silvestris silvestris, Felis silvestris lybica*), leopards (*Panthera pardus*), tigers (*Panthera tigris*) and bobcats (*Lynx rufus*) [3]. *Troglostrongylus brevior* was described six decades ago in the bronchi of *F. s. lybica* and *F. chaus* in Palestine and reported more recently in domestic cats from Spain and Italy [3, 5, 6]. Since then, *T. brevior* has been found in recent years in domestic cats in Italy [7], Greece [8], Bulgaria [7], Spain [9], Cyprus [10], but also in *Lynx lynx* in Bosnia and Herzegovina [11]. The two species, *T. brevior* and *Ae. abstrusus* share a similar biology (indirect life-cycle, using intermediate and paratenic hosts); the diagnosis by the first-stage

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larvae (L1) may be challenging, which seems to have caused confusion between *T. brevior* and *Ae. abstrusus* for a long time [2, 12]. In contrast, adult nematodes inhabit specific respiratory tracts and they are clearly distinct morphologically. Adults of *Ae. abstrusus* are localized in the pulmonary tissue, forming sub-pleural nodules, while adults of *T. brevior* inhabit the trachea, bronchi and bronchioles [2, 3, 6]. *Troglostrongylus brevior* infection is more common in kittens and young cats, producing severe respiratory distress or being lethal, while reports of fatal *Ae. abstrusus* infection are unusual [2]. However, except a few large scales studies in domestic [7] and wildcats [13], the knowledge on the distribution of this neglected feline parasite is still scarce.

The present paper reports the first case of *T. brevior* infection in Romania, bringing proof for its broader geographic distribution in Europe.

Methods

Sample origin and lungworms collection

In July 2017, a road-killed juvenile male Felis silvestris in good body condition was collected near Crasna village in Covasna County (45.585185°N, 26.155503°E, 669 m altitude), Romania. The animal was identified based on morphological and morphometrical characteristics as either a wildcat or a hybrid [14] and the approximate age was established by teeth examination [15]. A full necropsy was performed and the trachea and the lungs were extracted and examined for the presence of parasites. The entire respiratory tract (trachea, bronchi and bronchioles) was longitudinally opened using scissors and observed under a stereomicroscope for the presence of parasites [6, 16]. Nematodes collected from the trachea and bronchioles were washed in saline solution and temporarily mounted on microscope slides, examined, measured and photographed, using an optical microscope (Olympus BX51; Soft Imaging solution GMBH LG20, Munster, Germany). Parasites were sexed and identified to species level, based on morphometrical features [5]. Small pieces from a female nematode were fixed and stored in 70% ethanol. A classical Baermann method [17] was performed on the lungs and the faeces to collect the metastrongyloid larvae (L1).

Molecular analysis and species identification

Genomic DNA was extracted from an adult female nematode, using a commercial kit (Isolate II Genomic DNA Kit, Bioline, London, UK), according to the manufacturer's instructions. Molecular identification was accomplished with a PCR assay, targeting the internal transcribed spacer 2 (ITS2, ~500 bp) of the rRNA gene, using primers and protocols available in the literature [18, 19]. Amplicons were purified using silica membrane spin columns (QIAquick PCR Purification Kit, Qiagen,

Hilden, Germany) and then sequenced by Macrogen Europe (Amsterdam, Netherlands). The sequence was compared to those available in the GenBank by Basic Local Alignment Tool (BLAST) analysis. Species identification (adults and larvae) was based upon morphological characteristics, correlated with molecular analysis [1, 5, 6, 20].

Results

Morphology and morphometry of Troglostrongylus brevior

Nematodes were found in the trachea and bronchioles (two males and one female) and were morphologically identified as T. brevior. The males were 7.73 mm and 7.76 mm long, with 0.29 mm and 0.31 mm maximum width, respectively, presenting a folded cuticle. The length of oesophagus varied between 0.27 mm and 0.30 mm, and the excretory pore was located at 107.45-113.13 µm from the cephalic end. The posterior end of males showed a well-developed copulatory bursa (Fig. 1), without a clear delineation of the median and lateral lobes and several distinct rays: dorsal, externo-dorsal, lateral and ventral. The dorsal ray was elongated and showed four apical papillae. The externo-dorsal ray was shorter than the dorsal ray, well-separated by dorsal and lateral rays and had a single extension. The lateral ray was divided into three branches: antero-lateral, medio-lateral and postero-lateral; the second and third were partially joined from the basis along half of their length. The ventral ray was medium sized compared to the other rays and distally split into two small extensions. The spicules were equally calibrated, measuring 0.64 mm, and 0.66 mm, respectively. Due to partial destruction during the removal from the lung tissue, the female nematode could not be measured.

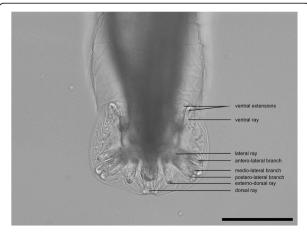


Fig. 1 Characteristic copulatory bursa in male *Troglostrongylus brevior*. *Scale-bar*: 100 µm

The first-stage larvae (L1) recovered from the lung tissue and faeces were morphologically consistent with those of *T. brevior* (average length of 0.34 mm, average width of 0.02 mm) with a sub-terminal oral opening and a pointed tail with a pronounced dorsal spine and a less deep ventral incision.

No other pulmonary nematodes or intestinal parasites were found and no gross pulmonary lesions were observed.

Molecular analysis

The BLAST analysis of the ITS2 sequence (accession number MF997544) revealed a 100% nucleotide similarity to *Troglostrongylus brevior* from Sardinia (KF241978.1).

Discussion

Wildcats are considered to be an important reservoir and also the natural hosts for Troglostrongylus spp. [16]. Troglostrongylus brevior was originally described from wild felids in Palestine [5] and since then, the nematode has been found predominantly in countries where population of wildcats are present [7, 8, 11]. Troglostrongylus infections in domestic cats were reported for the first time in Europe, after 60 years from the first description in wildcats [3, 6, 9]. This may be due to the misdiagnosed cases of Troglostrongylus in domestic cats, as the firststage larvae morphologically resemble those of Ae. abstrusus [6] and most of the diagnoses are based on coproscopy [7, 21–24]. However, T. brevior might be emerging in domestic cats, as a result of an increased interaction between wild and domestic carnivores, due to urbanization by land-clearing that forces wildlife to move into new areas, such as suburbs. The few cases of troglostrongylosis in domestic cats are characterized by severe and fatal clinical outcome. Clinical signs reported in young domestic cats include general respiratory signs, coughing, tachypnea, dyspnoea, polypnoea and depression, while in adult animals the infection is subclinical [2].

In the cat from the present study no gross lesions were noted, possibly highlighting the co-evolution of this parasite-host system. However, the absence of the pathological changes may be related also to the small number of adult parasites identified. In domestic cats, T. brevior produces lung oedema, haemorrhages, emphysema and catarrhal exudate in the airways [6, 25, 26]. Troglostrongylus brevior, Ae. abstrusus and Angiostrongylus chabaudi share a similar biology and all are using molluscs as intermediate hosts in their life-cycle [1, 5, 27], sometimes even occupying the same ecological niche and evolving as mixed infections in the same host [28]. Troglostrongylus brevior was found in association with Ae. abstrusus, rather than in single infections [9, 20, 29, 30] or also in association with An. chabaudi, a feline vascular parasite [28]. In the present study the infection was monospecific, as neither Ae. abstrusus nor An. chabaudi were present. In the last 10 years, carnivore lungworm infections were reported as spreading into new areas and in some countries the infection became emerging [31–38]. However, this might be also a result of more intense surveillance rather than a true emergence.

The diversity of species infecting wild and domestic felids and the differences regarding the clinical significance of these nematodes highlight the need for a more intense surveillance and proper diagnosis of feline lungworm infections, especially in countries where more species were demonstrated to be present [8]. Furthermore, an increased awareness between clinicians is needed for a correct diagnostic approach to feline lungworm diagnosis.

Conclusions

This paper represents the first report of *Troglostrongylus* brevior infection in Romania, so far representing the second northernmost location for this genus in Europe.

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Availability of data and materials

The nematode specimens are deposited (registration number CJ006960) in the collection of the Department of Parasitology and Parasitic Diseases, University of Agricultural Sciences and Veterinary Medicine, Cluj-Napoca. The sequence was deposited in the GenBank database (accession number MF997544).

Authors' contributions

GD performed the necropsies, identified and counted the nematodes and wrote the manuscript. AMI performed the molecular work. ADM collected the material and critically reviewed the manuscript. CMG morphologically identified and described the parasites and coordinated the study. All authors read and approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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