Wildlife-transmitted *Taenia* and *Versteria* cysticercosis and coenurosis in humans and other primates

Peter Deplazes*, Ramon M. Eichenberger, Felix Grimm

Institute of Parasitology, University of Zürich, 8057, Zürich, Switzerland

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**ABSTRACT**

Wild mustelids and canids are definitive hosts of *Taenia* and *Versteria* spp. while rodents act as natural intermediate hosts. Rarely, larval stages of these parasites can cause serious zoonoses. In Europe, four cases of *Taenia martis* cysticercosis have been diagnosed in immunocompetent women, and two cases in zoo primates since 2013. In North America, a zoonotic genotype related but distinct from *Versteria mustelae* has been identified in 2014, which had caused a fatal infection in an orangutan and liver- and disseminated cysticercoses in two severely immune deficient human patients in 2018, respectively. Additionally, we could attribute a historic human case from the USA to this *Versteria* sp. by reanalysing a published nucleotide sequence. In the last decades, sporadic zoonotic infections by cysticerci of the canid tapeworm *Taenia crassiceps* have been described (4 in North America, 8 in Europe). Besides, 3 ocular cases from North America and one neural infection from Europe, all in immunocompetent patients, 6 cutaneous infections were described in severely immunocompromised European patients. Correspondingly, besides oral infections with taenid eggs, accidental subcutaneous oncosphere establishment after egg-contamination of open wounds was suggested, especially in cases with a history of cutaneous injuries at the infection site. *Taenia multiceps* is mainly transmitted in a domestic cycle. Only five human coenurosis cases are published since 2000. In contrast, *T. serialis* coenurosis (1 human case since 2000) is primarily transmitted by wild canids. The etiological diagnosis of exotic cysticercoses is challenging. Usually, clinical material does not allow for a morphological identification, and serological tests are not available. These limitations have partly been overcome by molecular tools. Without claiming any dramatic emergence of cysticercoses and coenuroses transmitted by wild carnivores, further sporadic cases of such ‘exotic’ infections have to be expected.

1. Introduction

Wildlife is confronted worldwide with expanding urban and agricultural areas and increasing human leisure activities. This development has strong effects on the biodiversity of ecosystems and poses a threat to many species. However, several carnivores and other species may take advantage of urban resources and new habitats, and they are increasing their populations. Furthermore, the stone marten (*Martes foina*) is known to be adapted to urbanised habitats for centuries (Wereszczuk et al., 2017). More recently, carnivore species thriving in anthropogenic habitats include the red fox (*Vulpes vulpes*) (Chautan et al., 2000), coyotes (*Canis latrans*, in Canadian cities) (Liccioli et al., 2015), raccoons (*Procyon lotor*, in North America and Central Europe) ( Mackenstedt et al., 2015) and European badgers (*Meles meles*, in Europe) (Geiger et al., 2018). Consequently, urbanization influences the transmission of parasites with complex life cycles, which often involve intermediate and definitive host species with very different ecologies (Liccioli et al., 2015; Lafferty, 1999). For example, a distinct increase of human alveolar echinococcosis has been observed in Central Europe, approximately ten years after the establishment of stable urban fox populations (Schweiger et al., 2007). And more recently, a clear increase of angiostrongylosis in dogs has been observed in parallel with a strong increase of the prevalence of this parasite in foxes living in the Swiss Plateau (Gillis-Germitsch and Schnyder, 2018). Furthermore, increasing recreational and sports activities of the urban population, together with their dogs, in agricultural and wildlife areas have increased the transmission risks of parasites between wild and domestic carnivores. This is supported by the observations of many veterinarians reporting increasing numbers of Sarcoptes infestations in dogs after Sarcoptes mange had become more prevalent in foxes in Eastern Switzerland (Deplazes P., unpublished). Trends in the transmission of parasites between wild and domestic carnivores have been reviewed for...
protozoa and tick-borne agents (Otranto et al., 2015a), as well as for helminths and arthropods (Otranto et al., 2015b).

The causative agents of human alveolar (AE) and cystic echinococcosis (CE) are *Echinococcus multilocularis* and *E. granulosus* sensu lato (s.l.), respectively (Kern et al., 2017), while *Taenia solium* is the etiological agent for cysticercosis (Flisser et al., 2011). Zoonotic transmissions of CE and *T. solium* cysticercosis mainly involve domestic animals: pigs as the intermediate hosts for *T. solium* (Flisser et al., 2011), and dogs as the definitive hosts of *E. granulosus* s.l. (Romig et al., 2017). In contrast, zoonotic transmission of AE is mainly attributed to increased contact zones of the wild definitive hosts (red foxes, coyotes and raccoon dogs) and humans (Romig et al., 2017). Carnivore transmitted taenid infections other than echinococcosis and *T. solium* cysticercosis (Table 1) have been sporadically described in immunocompetent (*T. martis* or *T. crassiceps* cysticercosis and *T. multiceps, T. serialis, T. brauni* and *T. glomerata* coenuroses) and typically in severely immunocompromised human patients (*T. crassiceps* and *Versteria* sp. cysticercoses) (Tables 2–4), as well as in other primates.

Primates are dead-end intermediate hosts for canid-transmitted taenid species but seem to be susceptible to some degree to the development of metacestodes (larval stages) after accidental infection with eggs. The epidemiology of these rare zoonoses is not well understood. Their apparently low incidences in humans might be attributed to a very low infection pressure or to a relatively high resistance of humans against these parasites. For example, most cases of subcutaneous *T. crassiceps* cysticercosis as well as all cases of fatal *Versteria* sp. infections have been described in severely immunodeficient patients (Table 2). Interestingly, humans have been considered highly resistant against AE, which would explain the relatively low incidences in areas with high environmental contamination with *E. multilocularis* eggs (Gottstein et al., 2015). However, in patients with impaired immunological status due to AIDS (Sailer et al., 1997), cancer and drug-based immunosuppression (Chauchet et al., 2014), severe progression of AE may occur (Vuitton et al., 2015).

Based on a relatively high number of cases in relation to the small population sizes, non-human primates in zoos seem to be highly susceptible for metacestode infections. Therefore, these primates might be considered as sentinels for the zoonotic potential of cestode parasite species. Indeed, mesocestoides infections have been detected in dead-end hosts such as zoo primates (Hubbard et al., 1993; Tokiwa et al., 2014; Montalbano Di Filippo et al., 2018) and in humans (Fuentes et al., 2004). However, how humans and primates get infected is not clear, and the taxonomy of *Mesocestoides* spp. from domestic and wild carnivores is under investigation (Varcasia et al., 2015a). Furthermore, *E. equinus* (G4), initially not considered to be zoonotic, was long known to be highly prevalent in horses as intermediate and in foxhounds as definitive hosts in Great Britain. In 2012, CE caused by *E. equinus* was identified in lemurs (*Varecia rubra, V. variegata*), living in a British zoo (Boufana et al., 2012; Denk et al., 2016) and in 2018, the first human CE case caused by *E. equinus* was discovered in Uzbekistan (Timbure et al., 2018). Other examples of *T. crassiceps, T. martis*, and *Versteria sp., T. multiceps* and *T. serialis* infections in humans and other primates will be documented in more detail in this review (Tables 2–4 and supplemental material).

Genetic analyses of specified gene sequences allow for a reliable species differentiation, even if only small amounts of parasite tissue are available. The detection of *T. martis* and *Versteria* sp. cysticercosis in humans and other primates can especially be attributed to the new molecular diagnostic strategies, and probably does not represent an emergence of these zoonoses.

Furthermore, the zoonotic potential of several other *Taenia* spp. (e.g. *T. ovis, T. hydatigena* or *T. taeniaeformis*) has been claimed in textbooks based on few observations. However, for *T. hydatigena*, two well documented cases of Cysticercus tenuicollis abdominal cysticercosis with large cysts have been described in non-human primates (Table 4). As an example, intestinal *T. taeniaeformis* infections are mostly found in felids, rarely also in red foxes (Table 1). Historically, a human intestinal infection with a *Taenia* sp. named *T. infantis* has been reported in a 5-year-old boy from Buenos Aires by Bacigalupo (1922). However, Jouyoux and Baer (1929) later showed that *T. infantis* was a synonym of *T. taeniaeformis*. Furthermore, the infections of a 55 years old woman and several children from Japan were attributed to *T. taeniaeformis* (cited in Sterba and Barus, 1976). The intermediate host stage of *T. taeniaeformis*, Strobilocercus fasciolaris, infects a large range of rodents and rabbits (Table 1), but there is at least one well documented record of *T. taeniaeformis* metacestode infection of the liver in a 77-year-old man diagnosed at post-mortem examination in 1974 (Sterba and Barus, 1976). Interestingly, Sterba and Barus (1976) mentioned a Strobilocercus infection in a gibbon (*Hylobates leucicus*). However, considering the worldwide distribution of *T. taeniaeformis*, we can conclude that this species has a neglectable zoonotic potential and should not be listed as a zoonotic agent.

Similarly, coenuroses caused by *T. multiceps, T. serialis, T. brauni*, and *T. glomerata* have been described as serious zoonoses for decades.
Table 2  
Cases of Taenia crassiceps cysticercosis in humans.

| Case no.; references | Patient, immune status and case history | Pathological findings | Etiological diagnosis: morphology and DNA analysis (PCR, sequencing) |
|-----------------------|----------------------------------------|-----------------------|---------------------------------------------------------------|
| Case 1                | Shea et al. (1973); Freeman et al. (1973) | 17-year-old immunocompetent Canadian woman, with decreased visual acuity of her right eye since two months. Four months earlier she suffered from a severe generalized erythematous skin condition. A family dog infected most probably with T. crassiceps (based on proglottid morphology) lived in intimate association with the patient. | A large motile cyst with a scolex and several smaller cystic lesions were observed at the posterior pole of her right eye. With a putative diagnosis of a T. crassiceps cysticercosis, cysts were removed by surgical intervention. | Morphology. Based on clinical material and from the established isolate in mice, T. crassiceps cysticerci were identified by rostellar hook morphology and numbers as well as by the budding proliferation of the cysticerci. DNA analysis. Nd. |
| Case 2                | Klinker et al. (1992) | 33-year-old German male, AIDS patient with a Pneumocystis carinii pneumonia and cerebral toxoplasmosis presented a growing paravertebral haematoma. The patient reported a fall around 4 weeks before the first symptoms appeared. | Subcutaneous, paravertebral infiltrate resembling a haematoma which spread over several weeks to cover almost the entire back. A spontaneous rupture of the infiltrate released “whitish spherical masses, 2–3 mm in diameter”. | Morphology. T. crassiceps diagnosis was confirmed by the number of hooks and the typical budding of the cysticerci. DNA analysis. Nd. |
| Case 3                | Chermette et al. (1995) | 33-year-old Frenchman, AIDS patient (stage IV). Suffered around 2 months earlier of a haematoma of the left arm after a fall. | Subcutaneous and muscular tissues invasion of the left arm with extension to the pectoral region. A surgical intervention revealed multiple larval forms. | Morphology. Ovoid (4 mm × 2 mm), transparent vesicles of different developmental stages, some containing one scolex with 4 suckers and 2 rows of rostellar hooks, some with external buds at the opposite site of the scolex. DNA analysis. Nd. |
| Case 4                | Chuck et al. (1997) | 38-year-old USA woman, (immune status not mentioned) with blurred vision in the right eye that had persisted for 4 weeks. The patient played with a 6-month-old German shepherd shortly before the dog passed a tapeworm. | Dilated fundus examination revealed a large, elevated, clear, fluid filled subretinal mass with several oval cystic structures of varying sizes. | Morphology. One of the cysts contained a mature scolex with characteristic hooks. Budding on the distal part of the cysticerci was indicative for T. crassiceps. DNA analysis. Nd. |
| Case 5                | Francois et al. (1998) | 38-year-old Frenchman with severe AIDS. He denied using intravenous drugs. Two months before admission he noticed a rapid progressive swelling in the right arm and forearm. The patient was in close contact with dogs and frequently walked in the forests of Normandy and Jura. | MRI showed a mass, suggesting a soft tissue neoplasm with numerous, heterogeneous, invasive cystlike lesions. Surgery revealed lesions “containing a yellowish viscous fluid, hundreds of granules, and cysticerci like small vesicles”. | Morphology. DNA analysis. Nd. |
| Case 6                | Maillard et al. (1998) | 34-year-old Frenchman, AIDS patient (C3) clinical disease, developed a traumatic haematoma on the left arm after a fall in the countryside. One month later, the swelling spread to the left pectoral region. | US and MRI showed a dissociation of muscular fibres. An incision on the left arm produced “a fluid that contained many transparent, spherical masses, 1–3 mm in diameter”. | Morphology. Microscopic examination revealed parasites compatible with larvae of T. crassiceps with single or multiple external buds at the opposite side of the scolex. DNA analysis. Nd. |
| Case 7                | Heldwein et al. (2006) | 82-year-old German woman, with a history of colon cancer with hemicolecotomy and a B cell non-Hodgkin’s lymphoma treated with fludarabine phosphate and cyclophosphamide two months before admission. She had also undergone radiotherapy of retrocardiac and iliac lymph nodes. Progressive pain and swelling in the left forearm and back of the left hand had started six weeks earlier after a fall on the hand with soft tissue injury. | US and MRI of the limb showed massive oedema of subcutaneous tissue and in between muscles and tendons and multiple cystic lesions were demonstrated. At fasciotomy jelly-like tissue containing multiple spherical masses with diameters of up to 2 mm, “similar to fish spawn”, was removed from the subcutis, muscles, and tendons. | Morphology. Cysticerci with an ellipsoid cystic body, a long and retractable neck, and a single scolex with four suckers and two rows of hooklets. Additionally, buds at the posterior end of the larve were observed. DNA analysis. Nd. |
| Case 8                | Goesseringer et al. (2011); Flammer Anikpeh et al. (2014) | 47-year-old Swiss woman, severely immunodeficient (HIV-1 RNA viral load of > 4 million copies/mL). Suffered from an injury to her right wrist during her work as a zoo-employee 5 months earlier. Exposed to dogs and foxes. | Swollen and painful right forearm for 2 weeks with clinical presentation and MRI suggestive for a necrotising fascitis. “Surgical exploration revealed small transparent cystic bodies resembling white caviar”. | Morphology. Small (4–5 mm) transparent cystic bodies with a retractable neck and a single scolex with four suckers and two rows of hooklets as well as budding were indicative for T. crassiceps. DNA analysis. 125 [P6] “Sequence analysis of a fragment of the small subunit rRNA gene permitted species identification” (but this was not substantiated). |
| Case 9                | Schmid et al. (2014) | 57-year-old Swiss woman, immunocompetent (serological testing for HIV negative) and other immunological parameters were inconspicuous, presented with swelling and an 8-cm haematoma localized on the right temple. She had no history of a traumatic incident. | Sonography revealed a hypechogenic lesion within the temporal muscle. A small amount of whitish material was aspirated. | Morphology. Swarms revealed a putative tapeworm larva and accumulation of eosinophil granulocytes (Papanicolaou stains). DNA analysis. 125 [P6] sequence analysis tentatively identified the parasite as T. crassiceps nd1 [P3]: Identity 421/421 bp or 100% with T. crassiceps nd1, (GenBank AF216699). |

(continued on next page)
For *T. multiceps*, the parasite lifecycle is mainly maintained in a domestic cycle including dogs and farm ruminants (Abera et al., 2016). Nevertheless, wild carnivores such as foxes and wolves have been demonstrated to be responsible for the dissemination of tapeworm eggs (Varcasia et al., 2015b; Otranto et al., 2015b). In wolves have been demonstrated to be responsible for the dissemination (Abera et al., 2016). Nevertheless, wild carnivores such as foxes and wolves have been demonstrated to be responsible for the dissemination of tapeworm eggs (Varcasia et al., 2015b; Otranto et al., 2015b). In contrast, for *T. serialis*, the cycle is mainly based on wildlife with wild canids and herbivores (primarily lagomorphs) as natural hosts. In Africa, two more *Taenia* spp. of wildlife origin have been described causing coenurosis with zoonotic potential (*T. brauni* and *T. glomeratus*) (Morel, 1959), to which we refer as ‘African-type coenurosis’ in this review. However, based on their appearance, these species might be variants/strains of *T. serialis* (no molecular data is available). Over the last century, several cases of coenurosis have been described in humans and other primates. Clinically, *T. multiceps* coenurosis in humans caused severe central nervous symptoms and ended fatally in 42% of the collected case reports (Supplemental Table 1). In contrast, coenurosis by *T. serialis* and the ‘African-type’ coenuroses have a tropism to subcutaneous and muscle tissues, and the clinical outcome is favourable. Taking account of the high prevalence of these parasites globally and the low number of cases published, the zoonotic risk of coenurosis has to be considered minimal, irrespective of potential severe consequences.

In this review we focus on non-*T. solium* cysticercoses and on coenuroses that are transmitted through wild carnivores (mustelids, canids, and exceptionally felids), exemplifying the habitual infection biology of involved species, refer to their zoonotic potential and discuss “state-of-the-art” diagnostic strategies. The herein reported information was retrieved from full texts by comprehensive database searches (PubMed and Google Scholar). Extensive screening of reference lists for historical and missing publications was accomplished.

### 2. Mustelid-transmitted cysticercoses

#### 2.1. *Taenia martis* cysticercosis

*Taenia martis* is a large tapeworm of around 20 cm in length that develops in the small intestine of wild carnivores, mainly Mustelidae. In North America, *T. martis americana* has been described in *Martes americana* (Hoberg et al., 1990). In Europe, it was described in the stone marten (*Martes foina*) and in the pine marten (*M. martes*). However, other mustelids and rarely canids and felids might also act as definitive hosts (Table 1). *Taenia martis* is transmitted in a wild-animal cycle between mustelid carnivores and their prey, including rodents and other small mammals as intermediate hosts. In the intermediate hosts the larval form (cysticercus) reaches a size of 6–32 mm (more details see diagnostic section). It mainly develops in the pleural and peritoneal cavity as a pseudo segmented larva, not surrounded by a fibrous capsule, and without asealexpression (Schuster and Benitz, 1992).

In Europe, *T. martis* has been observed in martens in Italy (Millan et al., 2001), Belgium (Mathy et al., 2009) Poland (Korná et al., 2013) and Belarus where it has also been detected in the European polecats (Shimalov, 2010). In Southwestern Germany, *T. martis* was found in 1 (0.03%) of 3573 red foxes, in 2 (2.2%) of 84 badgers (*Meles meles*) and in 17 (37%) of 47 stone martens (Loos-Frank and Zeyhle, 1982). In southwestern Yakutia (Siberia) between 1981 and 1987, *T. martis* was found in 28% of 272 and 30% of 1548 *Martes zibellina*, respectively (Sedalishev and Odnokurtsev, 2011; Odnokurtsev and Sedalishev, 2011).

*Taenia martis* cysticercosis in rodents seems to occur all over Europe but focally in variable prevalence. As an example in Eastern Switzerland, *T. martis* larval stages (cysticerci) were detected in 43 (10.5%) of 411 *Apodemus flavicolis*, in 100 (7.8%) of 1276 *A. sylvaticus* and in 294 (24.3%) of 1211 *Myodes* (syn. *Clethrionomys*) *glareolus*, but not in 894 *Arvicola terrestris*, 347 *Microtus arvalis* and 250 *M. agrestis* of the same area (Schaerer, 1987). In contrast, in Western Switzerland (Geneva) *T. martis* was found in 2 of 99 *A. flavicolis*, but not in 466 *A. terestris*, 58 *M. glareolus* and 35 *M. arvalis* (Reperant et al., 2009). Comparable low prevalence of *T. martis* were found in Berlin, Germany (1.4% of 59 *M. glareolus*, 78 *A. agrarius* and 82 *A. flavicolis*) (Krücken et al., 2017), and in Western France in nutria (*Myocastor coypus*) (Umhang et al., 2013). In Germany, muskrats (*Ondatra zibethicus*) were highly infected with *T. martis* cysticerci (*Loos-Frank and Zeyhle, 1981*), similar to the Limburg region in the Netherlands (18.6% of 526 muskrats), but in contrast to animals from Groningen (n = 1200) with a prevalence of merely 0.7% (Borgstede et al., 2003). From Ireland, Loxton et al. (2017) reported a prevalence of *T. martis* of 0.77% (CI: 0.16–2.24) in wood mice (*A. sylvaticus*). Reports from northern Spain document the occurrence of *T. martis* in 0.7–1.0% of 376 *M. glareolus* (Ribas et al., 2009). In Serbia 4.1% of 588 *M. glareolus* were infected with *T. martis* (Bjellic-Čabriolo et al., 2011). In Belarus, the red-backed vole, the striped field mouse and the yellow-necked mouse are hosts of this helminth (Shimalov, 2010), and recently it was documented there as a parasite of red squirrels (*Sciurus vulgaris Linnaeus, 1758*) (Shimalov, 2016).

To our knowledge, four cases of human *T. martis* cysticercosis have been reported in immunocompetent women (Table 3). Interestingly, three of these patients originated from Alsace (France) and Western Germany, while the remaining fourth case was described in northern Germany. In these cases, cysticercosis manifested itself as a solitary lesion in the brain, in the peritoneum and in two cases in the eye. Furthermore, two cases in non-human primates, one originating from...
Cases of *Taenia martis*, *Versteria* sp. (zoonotic genotype) and non-specific cysticercoses in humans.

| Taeniid species, cases, citations | Patient, immune status and case history | Pathological findings | Etiological diagnosis: morphology and or DNA analysis (PCR, sequencing) |
|-----------------------------------|----------------------------------------|----------------------|-------------------------------------------------|
| *Taenia martis*                   | 43-year-old immunocompetent German woman (Freiburg in Breisgau), with mobile subretinal tumor with adjacent intraartenal and subhyaloid bleeding. | The removed cyst (after 8 days of albendazole/dexamethason therapy) showed the characteristic macroscopic and histologic features of a cysticercus bladder wall. | DNA Analysis. coxl [P1], ndl [P4], 12S [P8]; all sequences showed highest identity with *T. martis* (99%-100%). |
| Case 1, Eberwein et al. (2013)    |                                        |                      | DNA analysis. coxl [P1], 1396/1396 bp or 100% with *T. martis* EU544557, ndl [P3] (488/188 bp or 100% with EU544607) and 12S [P6] (203/265 bp or 99.2% with JX415820). |
| *T. martis*                       | 44-year-old immunocompetent French woman (Alsace), with suspected meningoencephalitis. | Thick-walled parasitic cyst with dense fibrosis and intense mononuclear inflammation contained dense fluid consisting of thick bright eosinophilic ribbons of membranous tissue and calcareous corpuscles. | Morphology. Macrocystically, the 3 mm long semi-transparent parasite had the appearance of a cestode larva. Tissue sections showed a characteristic tapeworm tegument. DNA analysis. Identities for coxl [P1], ndl [P4], 12S [P8] were 100%, 99%, and 99%, respectively with *T. martis* sequences. |
| Case 2, Brunet et al. (2015)      |                                        |                      | DNA analysis. 12S [P5]; Identity 99% with *T. martis* (DNA JF837555). |
| *T. martis*                       | 70-year-old immunocompetent woman from northern Germany; visual acuity of her left eye dropped significantly. | Intraocular inflammation and vitreous haemorrhage without fundus view in the affected eye. On US, a retinal detachment was disclosed and at surgery a moving larval parasite was extracted from the eye. | DNA analysis. Our recent re-evaluation of the 183 bp sequence of the 18S rRNA gene including the V2 region published by Olson et al. (2003) (GenBank AY199376), showed an identity of 95% (172/180 bp, 3 gaps) with the corresponding sequence of *V. mustelae* (GenBank AB731163), and the sequence had a 99.7% identity with the amplified sequence of the same gene of patient 2 (GenBank MG001670). |
| Case 3, Koch et al. (2016)        |                                        |                      | Morphology. A larva could be detected with suckers, a two-layered tegument and calcified corpuscles. DNA analysis. 12S [P5]; Identity 99% with *T. martis* (DNA JF837555). |
| *Versteria* sp. (zoonotic genotype) | 36-year-old immunocompetent woman (Germany); recurring, asymptomatic ascites, which progressed slightly over a period of 8 weeks | Histopathology assessment revealed a tumor mass with central necrosis and aggregates of epithelial cells with intermingled multinucleated giant cells of Langhans type in the periphery. | DNA analysis. Our recent re-evaluation of the 183 bp sequence of the 18S rRNA gene including the V2 region published by Olson et al. (2003) (GenBank AY199376), showed an identity of 95% (172/180 bp, 3 gaps) with the corresponding sequence of *V. mustelae* (GenBank AB731163), and the sequence had a 99.7% identity with the amplified sequence of the same gene of patient 2 (GenBank MG001670). |
| Case 1, Rudelius et al. (2017)    |                                        |                      | Morphology. Histopathology revealed cestode larva with parts of a rostellum and 2 refractile hooklets without species diagnosis. DNA analysis. Needle biopsies of the liver and formalin-fixed paraffin-embedded liver tissue were used for DNA analyses. 12S [P6]; Identity 98% with a sequence from a *Versteria* sp. isolate (GenBank KP033341) from a captive orangutan. |
| *Versteria* sp. (zoonotic genotype) | 58-year-old man from Pennsylvania (USA) with Hodgkin disease died after repeated courses of chemotherapy and radiotherapy. | The patient presented with severe pulmonary and systemic symptoms*. CT revealed mixed alveolar opacities and “a large heterogeneous central hepatic lesion (19.3 × 15 × 8.7 cm) abutting the middle hepatic and left portal veins, with multiple satellite nodules”. | DNA analysis. Suggestive of metacestodes DNA analysis. coxl [primer not specified]; Identity 98% (of 129 bp) with a sequence of *Versteria* sp. (GenBank KT223034). |
| Case 1, Connor et al. (1976)      |                                        |                      | DNA analysis. 12S [P6]; Not conclusive. |
| *Versteria* sp. (zoonotic genotype) | 53-year-old female from rural New Brunswick (Atlantic Canada) with a 3-day history of fever, productive cough, myalgia, malaise, and anorexia. Her past medical history included an obstructive nephropathy necessitating a kidney transplant with immunosuppression with tacrolimus, mycophenolate mofetil, and prednisone. | “The patient presented with severe pulmonary and systemic symptoms”. CT revealed mixed alveolar opacities and “a large heterogeneous central hepatic lesion (19.3 × 15 × 8.7 cm) abutting the middle hepatic and left portal veins, with multiple satellite nodules”. | DNA analysis. Our recent re-evaluation of the 183 bp sequence of the 18S rRNA gene including the V2 region published by Olson et al. (2003) (GenBank AY199376), showed an identity of 95% (172/180 bp, 3 gaps) with the corresponding sequence of *V. mustelae* (GenBank AB731163), and the sequence had a 99.7% identity with the amplified sequence of the same gene of patient 2 (GenBank MG001670). |
| Case 2, Barkati et al. (2018)     |                                        |                      | Morphology. Histopathology revealed cestode larva with parts of a rostellum and 2 refractile hooklets without species diagnosis. DNA analysis. Needle biopsies of the liver and formalin-fixed paraffin-embedded liver tissue were used for DNA analyses. 12S [P6]; Identity 98% with a sequence from a *Versteria* sp. isolate (GenBank KP033341) from a captive orangutan. |
| *Versteria* sp. (zoonotic genotype) | 68-year-old North American woman with hypogammaglobulinemia and previously treated lymphoma presented with fever and abdominal pain. | CT revealed numerous nodules in the lung, eye, brain, and liver. An open liver biopsy revealed “numerous nodular lesions and a mass made up of multifocal coalescing cystic lesions”*. Histopathology revealed 3-layered lesions a bladder wall and calcareous corpuscles in a matrix of granulomatous tissue inflammation with areas of necrosis. | DNA analysis. Suggestive of metacestodes DNA analysis. coxl [primer not specified]; Identity 98% (of 129 bp) with a sequence of *Versteria* sp. (GenBank KT223034). |
| Case 3, Lehman et al. (2018)      |                                        |                      | DNA analysis. 12S [P6]; Not conclusive. |
| *Taenia* sp.                      | 14-year-old (probably immunocompetent) man from Auvergne (Central France), was presented with acute ocular pain, haemorrhage of the conjunctiva and fever. The boy grew up in a rural environment with a pet dog and vegetable garden. | A “parasite” was observed in the internal chamber and surgically resected. | Morphology. A 6 mm long cysticercus was diagnosed by parasitologists, but the morphology did not allow species identification. DNA analysis. Nd. |
| Ocular cysticercosis, Mougeot et al. (1996) |                                        |                      | DNA analysis. Nd. |
| *Taenia* sp.                      | 15-year-old, Austrian, immune-competent woman presented with an iridocyclitis in the right eye. She had close contact to the young family dog. | A living parasite stage could be isolated from the anterior chamber. | Morphology. Contractable parasite stage of 2 mm length without characteristic structures such as a scolex. A putative *T. crassiceps* diagnosis was based on antibody reactions against *T. crassiceps* larval antigens in Western blots DNA analysis. Nd. |
| Ocular cysticercosis, Arocker-Mettinger et al. (1992) |                                        |                      | DNA analysis. Nd. |

Nd: Not done or no data given.

a Primer pairs used are given in squared brackets and refer to Table 6.
b Analysis could not consider *T. martis* and other rare cysticercoses.

Alsace, the other from a zoo in Rome (Italy), presented as abdominal *T. martis*-cysticercosis (Table 4). Finally, in a 14-year-old Frenchman and a 15-year-old Austrian woman, ocular cysticercoses caused by juvenile taeniid larvae could not be morphologically specified (Table 3). Similarly, a cysticercosis with fully developed cysticerci isolated from the abdomen of a lemur from a zoo in the USA in 1989 was diagnosed as *Taenia* sp. cysticercosis (Table 4).

Routes of human infection can only be speculated about. Two patients were recreational gardeners and could have been exposed to marten faeces during such activities. Extensive recreational hiking in
Table 4

| Taenia spp., case no., citation | Primate species, patient, origin, and case history | Pathological findings | Etiological diagnosis: morphology and DNA analysis (PCR, sequencing)* |
|--------------------------------|----------------------------------------------------|-----------------------|--------------------------------------------------|
| *Taenia crassiceps*            | Case 1, Baer and Schneider (1946)                  | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. crassiceps*                | Case 2, Dyer and Greve (1998)                      | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. crassiceps*                | Case 3, Young et al. (2000)                        | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. crassiceps*                | Case 4, Luzón et al. (2010)                        | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. crassiceps*                | Case 5, Ali et al. (2017)                          | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. crassiceps*                | Case 6, Bleyer et al. (2018)                       | Necropsy. Larval stages subcutaneously and in the smooth muscles, penetrating to the retroperitoneal cavity, with parasite infiltration into the spinal cord, | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *Taenia serialis*              | Case 1, Schwartz (1927)                            | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. serialis*                  | Case 2, Sandground (1937)                          | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. serialis*                  | Case 3, Elek and Finkelstein (1939)                | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. serialis*                  | Case 4, Clark (1969)                               | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. serialis*                  | Case 5, Schneider-Crease et al. (2013)             | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. multiceps* (probably misdiagnosed) | Leith and Satterfield (1974) | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |
| *T. brauni*                    | Case 5, Railliet and Marullaz (1919)               | Necropsy. Large subcutaneous tumor (1.5 kg). The majority of the mature cysts contained several hundred scolecites. | Morphology. Diagnosis based on number (n = 30–35) and size (large 182–200, small 138 < 150 μm) of rostellar hooks and budding of the cysticerci. DNA analysis. Nd. |

*nd*: not determined.

(continued on next page)
Table 4 (continued)

| Taenia sp., case no., citation | Primate species, patient, origin, and case history | Pathological findings | Etiological diagnosis: morphology and DNA analysis (PCR, sequencing)* |
|-----------------------------|-------------------------------------------------|----------------------|---------------------------------------------------------------|
| Taenia brauni Fain (1956)   | T. pisiformis frequently infected with than males (Morales-Montoret al., 2002), and females (Morales-Montoret al., 2002) | Cysts in brain (parietal lobe; 2 cm), heart (apex; 1 cm), and parotid gland | Morphology. Performed but no details provided by the author. DNA analysis. Nd. |
| Taenia sp. coeunurosis Lau et al. (1973) | White-handed Gibbon (Hylobates lar) at zoological park in the USA. Left eye progressing gradually for 3–4 weeks. Retinal mass at radiography. Euthanasia. | Cyst behind the left eyeball. | DNA analysis. Nd. |
| Taenia martis De Librero et al. (2014) | 18-year-old male ring-tailed Lemur (L. catta), immunocompetent, lives in a zoo in Rome (Italy). Apathy, loss of appetite, abdominal distension and diarrhea were observed 10 days before death. | Severe exudative fibrous-purulent peritonitis with numerous adhesions between the abdominal wall and the bowel loops. After intestine removal, two free and viable, 4 cm long, whitish, leaf-like parasitic forms were pinpointed. | DNA analysis. Nd. |
| T. martis Brunet et al. (2014) | 3-years-old subadult mail tonkana macaques (Macaca tonkana), immunocompetent, born and raised at Strasbourg University Centre of Primatology, France. | Abdominal mass (± 10 cm × 5 cm) was detected at palpation without clinical signs. | DNA analysis. Nd. |
| Versteria sp. Goldberg et al. (2014) | Juvenile (sex?) Bornean orangutan (Pongo pygmaeus), captive born in Colorado, USA, (no indications of immunosuppression). Loss of appetite and intermittent, moist cough; became increasingly lethargic and was found dead after 2 days. | Necropsy. Diffuse hemorrhages in the lungs, splenomegaly, a pale mottled liver, and thoracic and pericardial effusions. Histopathology of the liver revealed cystic structures containing eukaryotic parasite cells. Cause of death: "acute respiratory distress due to disseminated infection with an unknown parasite". | DNA analysis. Deep sequencing identified a Taenia spp. 125 [P7] and cox1 [nd1] [P1/P3] phylogenetic trees placed the organism within Versteria sp. (Cestoda: Taeniidae). |
| Taenia hydatigena Hobbs et al. (2003) | 5-year-old male rhesus macaque (M. mulatta), born and raised in a primate colony in China and imported 15 months earlier to the Oregon National Primate Research Centre. An abdominal mass was discovered during routine physical examination. | Necropsy. A pale yellow cyst attached to the greater omentum containing 500 ml of flocculent yellow fluid. | DNA analysis. Morphology. Rostellar hooks, the large size of the cysticercus, and its location within the peritoneal cavity were consistent with the diagnosis of a Taenia hydatigena cestocercosis. DNA analysis. Nd. |
| T. hydatigena Tsubota et al. (2009) | 5-years-old male long-tailed macaque (Macaca fascicularis), born and raised in a primate research colony in China in a toxicity study in Osaka (Japan). The monkey showed no clinical signs and was sacrificed at the end of a dosing experiment. | Necropsy. A yellow cyst filled with more than 100 ml of pale yellow fluid was found in the abdominal cavity. | DNA analysis. Morphology. A cysticercus with a well-developed scolex was found in the cyst. DNA analysis. Nd. [P3]: Identities of 96.7%–98.5% with T. hydatigena sequences. |
| Taenia sp. Wolff et al. (1989) | Captive-born 1-year-old male ruffed lemur (V. variegata rubra) with unremarkable routine quarantine examination with radiography (Zoo, USA). | Necropsy. Extrapleural calcified larval cestode in the left ventro-caudal thorax and pulmonary nodule with a cysticercus in the left dorso-caudal lung. | DNA analysis. Nd. |

Nd: Not done or no data given.
* Primer pairs used are given in squared brackets and refer to Table 6.
* Diagnosed as Multiceps ramosus or M. lemarit by the author, which are considered to be a synonym for T. brauni (Loos-Frank, 2000).

The Alsop was reported from another patient but without obvious contact to wild animals. All four patients could successfully be treated by surgery and anthelmintics.

Interestingly, all T. martis-cysticercosis patients described so far originated from Europe, and no data are available on the zoonotic potential of T. martis americana. Furthermore, all four human T. martis cases were documented in 36, 43, 44 and 70 years old women, though the number of cases is too small to speculate on a gender association. However, the host's endocrine system can have an impact on the susceptibility to cestode infections. For example, castration of male and pregnancy in female pigs significantly increased the prevalence of naturally acquired T. solium cysticercosis (Morales et al., 2002), female mice are more susceptible to experimental infections with T. crassiceps than males (Morales-Mentor et al., 2002), and female rabbits are more frequently infected with T. pisiformis metacestodes than males (Dominguez-Roldan et al., 2018).

2.2. Versteria mustelae and Versteria sp. cysticercosis

Versteria mustelae (new genus created by Nakao et al., 2013; syn. Taenia mustelae) is an up to 10 cm long tapeworm of mustelids with a wide range of rodents as intermediate hosts, bearing in the liver the rather small, ovoid, 0.4–2.0 mm large cysticerci containing many calcified bodies (Freeman, 1956; Slais, 1973). In Europe, weasel (Mustela sp.), stone marten (Martes foina) and pine marten (M. martes), and in the USA American pine marten (Martes americana), ermine (Mustela ermine) and mink (Neovison vison) have been described as definitive hosts (Lee et al., 2016, Hoberg et al., 1990). Between 1981 and 1987, V. mustelae was identified in 2.9% of 272 Martes zibellina in Southwestern Yakutia (Sibiria) (Sedalischev and Odnokurtsev, 2011).

A variety of rodents have been identified as intermediate hosts of V. mustelae in Europe and Asia. In eastern Switzerland, V. mustelae was found in very low prevalences in 0.1% of 1276 A. sylvaticus, in 0.1% of 1211 M. glareolus, in 0.3% of 347 M. arvalis and in 0.4% of 250 M. agrestis, but neither in 411 A. flavi- collis nor in 894 A. terrestris (Schaerer, 1987). In Zealand (Denmark), 14 of 46 M. glareolus were infected with V. mustelae (Tenora et al., 1991), and this species was found in 9.3% of 172 M. glareolus trapped in rural forest habitats and in 0.4% of 231 animals from urban forests and parks but not in 41 and 129 A. flavi-collis, respectively (Al-Sabi et al., 2015). In this study, prevalences were determined by molecular analyses with considerably higher sensitivities as compared to morphological identification. Interestingly, the authors stated that "several poorly developed cysts without specific morphology were observed". In Sweden, V. mustelae was diagnosed by...
PCR in 13% of 56 A. amphibius and in 14% of 187 M. agrestis from field habitats, and in 8.4% of 655 M. glareolus, but not in 79 A. flavidus and 206 A. sylvaticus from forest habitats (Miller et al., 2017). In Finland M. glareolus, M. rutilus, M. rufocanus, M. agrestis, M. oeconomicus and A. flavidus have been identified as intermediate hosts of V. mustelae (syn. Taenia tenuicollis) (Tenora et al., 1983). In South-Central Finland, cysts of V. mustelae were detected in 9% of 34 M. agrestis and in 27% of 117 C. glareolus in the liver in a habitat with Mustela nivalis and M. erminea as main definitive hosts (Soveri et al., 2000).

In France (Jura), V. mustelae was more frequently observed in M. glareolus (24% of 349) than in Pitymys subterraneus (5% of 75), M. agrestis (9% of 47), M. arvalis (1% of 2520) and Apodemus sp. (1% of 230) (Le Pesteure et al., 1992). In Spain, V. mustelae was reported in P. duodecimcostatus, P. lusitanicus, P. pyrenaicus, M. agrestis, M. arvalis, M. cabrerae and M. glareolus (Feliu et al., 1997). Ribas et al. (2009) documented the occurrence of T. tenuicollis (former syn. of V. mustelae, not to be confused with Cysticercus tenuicollis of T. hydatigena) in 6.7–12.6% of 376 M. glareolus caught in northern Spain.

Genetic investigations revealed that a complex of species, genera and genetic lineages exist within the genus Versteria in North America (Lee et al., 2016). The authors suggested the occurrence of a “western lineage” (present in Colorado, Oregon, and Idaho, and the Northwest Territories (NWT) of Canada) and a “northern continental lineage” (present in Wisconsin and the NWT), with sympatry in the NWT. This northern continental lineage clusters with V. mustelae from Eurasia (Finland and Siberia). The western lineage responsible for fatal infections of an orangutan and a human patient (Case 2, Table 2) also infects muskrats (Ondatra zibethicus) in Idaho, USA, and the NWT.

So far, V. mustelae genetically related to the European isolates, have not been found in primates, including humans all over the northern hemisphere. On the other hand, in North America, a “zoonotic” lineage named Versteria sp. responsible for a fatal infection of an orangutan (Table 5) and in 2018 of infections in two human patients (Table 3) seems to be genetically closely related, but distinct from V. mustelae. In this review we refer to this genotype as Versteria sp. or North American zoontic V. mustelae variant as used by Goldberg et al., 2014; Lee et al., 2016. Furthermore, based on the description of a patient from Pennsylvania (USA) with Hodgkin disease and an undefined helminthic infection (Connor et al., 1976), Olson et al. (2003) published a 18S sequence (GenBank:AY193876) without significant homology to other sequences in the database at the time of publication. We reanalysed this sequence in 2019 and identified a homology of 95% with V. mustelae (GenBank:AB731633) of European origin. Moreover, we recently sequenced the corresponding 18S region obtained by molecular analysis of clinical material of a female patient from Atlantic Canada with a Versteria sp. infection (published by Barkati et al., 2018, Table 3), and detected a homology of 99.7% with the GenBank entry (AY193876) of the Pennsylvanian patient mentioned above. Therefore, we conclude that the patient originating from Pennsylvania (USA) (Connor et al., 1976) represents the first documented case of human Versteria sp. cysticercosis (patient 1, Table 3).

Interestingly, North American V. mustelae isolates from Minnesota (USA) and from Algonquin Park (Canada) are capable of asexual multiplication in the intermediate host (Freeman, 1956), however it is unclear whether V. mustelae or the zooontic Versteria sp. have been investigated. This phenomenon is distinct from the budding multiplication of T. crassiceps, and could be responsible for the systemic and invasive infections in heavily immunocompromised human patients. However, further investigations with genetically defined isolates are needed to elucidate the fascinating biology of Versteria spp. Furthermore, based on the small, not fully developed Versteria sp. cysticerci in the liver in immunocompromised patients, this infection must be considered in the differential diagnosis of alveolar echinococcosis as proposed by Barkati et al. (2018).

3. Canid-transmitted cysticercosis and coenurosis

3.1. Taenia crassiceps cysticercosis

Taenia crassiceps is a relatively large but harmless tapeworm, 10–22 cm in length, inhabiting the intestines of carnivore definitive hosts, and it is widely distributed in the northern hemisphere (Table 1). In rodents, which are the natural intermediate hosts for the parasite, the larval (metacestode) form of T. crassiceps has a particular asexual reproduction by budding both exogenously and endogenously (Baer and Scheidegger, 1946, Freeman, 1962; Slais, 1973). Exogenous budding at the abscosex pole can produce 1–6 daughter cisticerci, which can bud off or remain attached and form a scolex of their own. This continuous and uncontrolled proliferation leads to massive infections, most frequently involving the subcutis, and both pleural and peritoneal cavities, causing death of the intermediate host within several months, or serious pathological implications in a large variety of dead-end hosts, including humans and other primates. Sporadic cases of cysticercosis caused by T. crassiceps have been documented in humans (Table 2), non-human primates (Table 4), but also rarely in e.g. domestic dogs (Ballweber, 2009; Beugnet et al., 2009; Clermont et al., 1993; Hoberg et al., 1999), a cat (Wunschmann et al., 2003), red foxes (Konjević et al., 2016; Whipp et al., 2017) and a chinchilla (Chinchilla lanigera) (Basso et al., 2014).

Natural intestinal infections with T. crassiceps have been described in the northern hemisphere mainly in the red fox (V. vulpes), but also in several other canids. A study in New Brunswick and Nova Scotia in eastern Canada in red foxes reported prevalences of 50% for T. crassiceps (Smith, 1978). T. crassiceps was also reported from Greenland (Andreassen et al., 2017), from Svalbard island (Stien et al., 2010), and from China, with 2 of 27 red foxes, and 1 of 9 Tibetan sand foxes found infected (Li et al., 2013).

In European red foxes T. crassiceps is widely distributed with prevalence varying between 4.3 and 29%, and 7.6% in foxes living in the city of Zurich, Switzerland (Häfer et al., 2000). Similar prevalence ranges have been described for Germany (17.7–28.5%, Ballek et al., 1992; Loos-Frank and Zehly, 1982; Pfeiffer et al., 1997; Welzel et al., 1995), France (15.9–29%, Pétavy and Debloc, 1980; Pétavy et al., 1990), Spain (4.3–23%, Alvarez et al., 1995; Segovia et al., 2004), and Lithuania (26.4%, Bruzinskaitė-Schmidhalter et al., 2012). In Russia, 19% of 68 hunted red foxes were infected with T. crassiceps in central Yakutia (Sedalischev and Odnokurtsev, 2013), and 49% of 247 red foxes and 7% of 43 corsac foxes (V. corsac) of Omsk Oblast (Siberia) between 2000 and 2004 (Bukova, 2005).

Taenia crassiceps was also detected in wolves in Canada, Europe and Russia (Abuladze, 1976; Craig and Craig, 2005). In Latvia 9% of 34 (Bagrāde et al., 2009), and in North-West Caucasia 2.8% of 36 wolves were infected with T. crassiceps (Itin et al., 2018). Freeman (1961) detected T. crassiceps in 1.7% of 58 Canadian wolves and in one of 6 coyote-dog hybrids, but not in 68 coyotes.

Of 17 Hungarian golden jackals (Canis aureus L., 1758), T. crassiceps has been found in 40% (Takács et al., 2014) and infections were recorded in 0.6% of 179 wolves in northern Italy (Gori et al., 2015). The raccoon dog (Nyctereutes procyonoides) seems to be another suitable host of T. crassiceps: it was identified in 6 of 72 animals from the Republic of Belarus (Subbotin, 2009), and it was found in 3.5% of 85 raccoon dogs in Lithuania (Bruzinskaitė-Schmidhalter et al., 2012).

However, not only canids seem to be susceptible to T. crassiceps intestinal infections. Raccoons (Procyon lotor) were infected with T. crassiceps as reported from the North-Western Caucasus with a prevalence of 24% in 42 animals (Itin et al., 2018). Schuster et al. (1993) identified T. crassiceps tapeworms in 2 of 25 wild cats (Felis silvestris) and Loos-Frank and Zehly (1982) found infections in 6% of 47 stone martens (Martes foina) in Germany.
Finally, in several human patients with *T. crassiceps* cysticercosis, close contacts to domestic dogs were assumed as source of infection (cases 1, 4, 5, 8, 10; Table 2). However, the epidemiological importance of domestic carnivores is probably overstated. Umhang et al. (2014) reported 5 of 817 (0.6%) dog faecal samples from eastern France to be positive for *T. crassiceps* eggs. Dyachenko et al. (2008) detected *T. crassiceps* eggs in 7 of 17,894 (0.04%) dog samples in Germany and other European countries, but in none of 9064 cat faeces. However, in an older study in southwest Germany intestinal infections with *T. crassiceps* were also observed in 1% of 387 stray cats supplied for rabies examination, some had been shot by private hunters and some were road kills (Loos-Frank and Zeyhle, 1982).

As intermediate hosts in northern America, the muskrat (*Ondatra zibethicus*), the common vole (*Microtus arvalis*), the eastern chipmunk (*Tamias striatus*), the deer mouse (*Peromyscus maniculatus*), the meadow vole (*Microtus pennsylvanicus*), but also the woodchuck (*Marmota monax*), and lemmings (*Dicrostonyx groenlandicus richardsoni, Lemmus trimucronatus trimucronatus*) were identified (Leiby and Whittaker, 1966; Albert et al., 1972). In Europe *T. crassiceps* cysticercosis seems to occur focally in variable prevalence in rodents. In Switzerland, *T. crassiceps* cysticerci were detected in 2 (0.22%) of 894 *Arvicolinae* terrestres, in 3 of 347 (0.86%) *Microtus arvalis*, and in 1 (0.4%) of 250 *M. agrestis*, but not in 411 *Apodemus flavicollis*, 1276 A. sylvaticus and 1211 *Myodes (Syn. Clethrionomy) glareolus* of the same area in eastern Switzerland (Schaerer, 1987). These very low prevalence of *T. crassiceps*, as well as very low prevalence of *E. multilocularis* (0.11%) in *A. terrestris*, and the absence in the other species mentioned above, were probably associated with the significantly reduced fox population during this time, which was attributed to the rabies epidemic and the corresponding control measurements. Later studies reported higher *T. crassiceps* prevalence, e.g., in 2.0% of 889 A. terrestris, whereas none were detected in neither 83 M. glareolus nor 154 *Apodemus* sp. (Stieger et al., 2002), while another study found *T. crassiceps* in 1.9% of 856 *A. terrestris* (Burlet et al., 2011) in the Zurich Area (Switzerland). In the Geneva area (Switzerland) Reperant et al. (2009) detected *T. crassiceps* cysticerci in 2.6% of 466 *A. terrestris*, in 2.9% of 35 *M. arvalis*, and in 1.0% of 99 *A. flavicollis*, but not in 58 M. glareolus. Interestingly, no *T. crassiceps* or *T. taeniaeformis* were identified in a study in Berlin (Germany), including 77 *A. flavicollis*, 25 *A. sylvaticus, 72 A. agrarius, 56 M. glareolus*, and 10 *Microtus* sp. (Krücken et al., 2017). Finally, in Japan *T. crassiceps* was found in three of 46 *M. montebelli* but not in 187 *Apodemus speciosus* (Ihama et al., 2000).

Interestingly, no reports on the occurrence of *T. crassiceps* were found from Scandinavia, Ireland and the UK. No *T. crassiceps* infections were detected in 197 foxes from Wales (Jones and Walters, 1992a), in 843 foxes from Southern England (Richards et al., 1995), and in 366 foxes from Northern Ireland (Ross and Fairley, 1969). Furthermore, in this paper the intestinal helminths in foxes in the UK were reviewed and *T. crassiceps* was not mentioned in several older studies. Further, no *T. crassiceps* findings are mentioned in rodent investigations in Zealand, Denmark (Al-Sahi et al., 2015; Tenora et al., 1991), in 1702 rodents investigated in Sweden (Müller et al., 2016), nor in high numbers of rodents investigated in Finland (Tenora et al., 1983; Soveri et al., 2000), and Ireland (Loxton et al., 2017).

Because of the high *T. crassiceps* prevalence in red foxes and their high population densities, the fox seems to be responsible for the perpetuation of the parasite cycle, as well as for the contamination of the environment with eggs in Central Europe. A similar situation can be observed with *E. multilocularis* (Hegglin and Deplazes, 2017). However, only a few studies have documented environmental contamination with *T. crassiceps* eggs. For example, Hauser et al. (2015) identified *T. crassiceps* eggs through DNA analyses in 5 (8.6%) of 58 fox faecal samples, and in one (0.2%) of 402 dog samples collected during the course of one year in 14 different grassland areas in the canton of Zurich, Switzerland. Furthermore, *T. crassiceps* eggs have been identified in the washing water of one of 141 samples of food, which consisted each of around 40 heads of lettuce, as well as various vegetables and fruits (Federer et al., 2016).

A number of well documented cases of *T. crassiceps* cysticercoses have been published in humans and other primates (Tables 2 and 4). Most of the cases, which have all been published in the last 30 years, originated from Central Europe (Germany, Switzerland, and France). Most cases of humans involving subcysts and muscles have been associated with underlying immunosuppression (cases 2, 3, 5–8; Table 2), except case 9, where a subcutaneous infection was associated with a haematoma localized on the right temple in an immunocompetent Swiss patient, and case 12 where a subcutaneous infection on the shoulder of a patient from USA was documented (Table 2). In contrast, an intercerebellar (case 10) or intracerebral infections (cases 1, 4, and 11; Table 2) were not associated with an impaired immune system.

Surprisingly, in all cases, the infection started uni-focally and progressed by infiltration of the surrounding tissues, especially in the non-cerebral/neural cases. Due to the systemic spread of the activated oncospheres in the blood circulation after oral egg uptake we would expect simultaneous multifocal infections as described for *T. solium* (especially in severely immunocompromised patients) and for *T. serialis*, where in some cases multiple lesions were observed in individuals without any indication of immuno-suppression. Interestingly, among the 7 cases of subcutaneous *T. crassiceps* cysticercosis (Table 2), 5 had a history of preceding injuries associated with the later development of cysticercosis. In cases 2, 3, 6, 7 a haematoma after a fall was reported at the site of subsequent cysticercosis development. In case 8, the patient remembered, that 5 months before the swelling on the same arm started, an injury to her right wrist occurred during her work as a zoo-employee.

Generally, humans acquire taeniid infection by oral ingestion of infective eggs. The most probable route of transmission after contact with taeniid eggs in the contaminated environment is the hand-to-mouth route. Hypothetically, transmission has also been linked to water or food-borne sources (vegetables/fruit/berries), but any source attribution is uncertain (Alvarez-Rojas et al., 2018). *Taenia* eggs can be dispersed from carnivore faeces with water or by adhering to objects (e.g. shoes and tyres). For example *E. multilocularis* eggs have been found on the hair coat of foxes (Nagy et al., 2011), suggesting a variety of potential infection routes to humans.

There is experimental evidence that taeniid eggs can hatch and develop further without the gastric passage. In sheep it was demonstrated that intra-tracheal *E. granulosus* egg inoculation was followed by cystic echinococcosis development in the lungs (Thompson, 1995). Furthermore, embryophore-free (based on sodium hypochlorite treatment) but not enzymatically activated *E. multilocularis* oncospheres caused local alveolar echinococcosis after subcutaneous injection in a mouse model designed for the documentation of egg viability (Federer et al., 2015). The same method using 1000 *T. crassiceps* eggs resulted in subcutaneous cysticercosis (confirmed by PCR/sequencing as described in Trachsel et al., 2007) in 2 of 3 inoculated BALB/c mice (Joekel D. and Deplazes P., unpublished data). Therefore, taeniid eggs accidently contaminating cutaneous injuries might locally hatch and further develop to larval stages. A human case of subcutaneous alveolar echinococcosis (Tschudi and Ammann, 1988) associated with a cutaneous injury, and a subcutaneous cystic echinococcosis in the popliteal fossa at the site of a previous wasp sting (Battyany et al., 2010), have been documented.
3.2. Coenurosis

3.2.1. Introduction

Coenurosis results from infection with metacestodes of *Taenia multiceps*, *T. serialis*, *T. brauni* and *T. glomeratus*. Canids and rarely other carnivores are the definitive hosts harbouring adult tapeworms in the intestine (Table 1). Taenoparasites mature within 15–42 days, with 3–4 proglottids shed daily containing more than 35,000 eggs per proglottid (Willis and Herbert, 1984). For *T. multiceps* it was shown that most eggs are released already from the proglottis in the intestine of the final host (Herbert et al., 1984; Scala and Varcasia, 2006). Each of the described tapeworm species usually infects a specific range of intermediate hosts (Table 1). After spreading of the oncosphere through the blood system, a vesicle-shaped cyst develops as coenurus, containing a few to several hundred protoscolices (Lescano and Zunt, 2013; Willis and Herbert, 1984).

Classically, only *T. multiceps* was considered to cause CNS, spinal cord, and eye infections, while *T. serialis*, *T. brauni* and *T. glomeratus* distribute to soft tissue/subcutaneous connective tissue, the musculoskeletal system, and visceral organs (Lescano and Zunt, 2013). However, a morphologically similar and genetically identical strain (based on mitochondrial genes) of *T. multiceps* (referred to as *T. gaigeri* or *T. gaigeri* in goats and *T. skrjabini* in sheep) was shown to cause non-cerebral coenurosis in goats and to a lesser extent in sheep (Christodouloupolou et al., 2013; Schuster et al., 2010; Verster, 1969). However, for none of these strains the zoonotic potential has been documented.

Humans may be infected as dead-end hosts, with the cystic larvae usually developing in the CNS, eye, or subcutaneous or intramuscular tissues. Between 1994 and 2018, 7 cases of coenurosis in humans have been described, mainly in individuals with close contact to canids (Supplemental Table 1). Human coenurosis is much less common than cystic echinococcosis, though both causing agents have the same definitive hosts. Cases have been reported predominantly from African countries, and only a few cases have been documented in South- and North America and Europe. Patients with coenurosis usually have a space-occupying lesion caused by a single cyst of 2–6 cm in diameter. However, cysts up to 10 cm in diameter have been reported.

3.2.2. *Taenia multiceps* coenurosis

- *Taenia multiceps* (formerly *Multiceps multiceps*) is a large tapeworm with variable size between 20 and 120 cm. *Taenia multiceps* coenurosis (sometimes called gid, staggers, or sturdy) has been documented in scattered foci all over the world, with the exception of Australia and New Zealand (Scala and Varcasia, 2006). In North America, however, the last case of *T. multiceps* gid has been reported in the 1980s and the disease is considered absent, despite one supposed human case in 1982 (Becklund, 1970; Ing et al., 1998; Schellhas and Norris, 1985).

- The parasite life cycle is predominantly maintained by domestic and stray dogs, when they have access to residuals from improper discarded slaughtered meat, or roaming of stray dogs around small ruminant farms and early access to animals that died in the field (Abera et al., 2016; Vasileiou et al., 2015). Global transport of domestic dogs (travel and trading) represents yet another risk for the dissemination of taenid eggs. Correspondingly, in Switzerland where only rare and sporadic cases are noticed, an outbreak of coenurosis in a dairy sheep flock has been reported, with approx. 10% of 140 sheep presenting clinical signs. The source of infection was a single livestock guardian dog imported from Italy (Schweizer et al., 2006).

- The importance of wild carnivores for the distribution of taenid eggs is valid where wild canids (e.g. wolves and foxes) have access to animals that have died in the field infected with *T. multiceps*. The role of wild carnivores was neglected as it was assumed that they have no access to sheep brains protected by the thick skull. However, firstly, the presence of cysts is associated with thinner skull bones, thus allowing access, even for foxes (Scala and Varcasia, 2006). Secondly, infection trials have demonstrated that foxes can develop gravid proglottids and infective eggs (Varcasia et al., 2015b). Correspondingly, in various regions where *T. multiceps* coenurosis is endemic in sheep, tapeworms have been reported from fox species, with 1–28.2% positive results from necropsied foxes in Russia (summarised in Varcasia et al., 2015b), 0–5% in Germany (Ballek et al., 1992; Loos-Frank and Zeyhle, 1982; Welzel et al., 1995), 2% in Peru (Moro et al., 1998), 3.8% in Jordan (El-Shehaby et al., 1999), 4.8–8.2% in Iran (Dalimi et al., 2006; Nabavi et al., 2014), and 6% in Tibetan sand foxes in China (Li et al., 2013). An extensive survey on fox hounds and foxes in Wales revealed 0.46%–1.7% dogs infected with *T. multiceps* but none of the foxes (Jones and Walters, 1992a, 1992b). However, the foxes had other *Taenia* species, including *T. serialis* which is responsible for coenurosis in lagomorphs in the UK (see below). In addition, earlier publications demonstrate the presence of *T. multiceps* in foxes in Bulgaria, Czech Republic, Italy, Poland and Switzerland (summarised in Loos-Frank and Zeyhle, 1982).

- The wolf is a known definitive host for *T. multiceps* and *T. serialis* (Craig and Craig, 2005; Otranto et al., 2015b). Data are available for European grey wolf populations with prevalences of 0.5% in Croatia (Hermosilla et al., 2017), 3.9% in Serbia (Girovic et al., 2015a), 0–9% in Italy (Gori et al., 2015; Guberti et al., 1993; Paoletti et al., 2017), 27% in Estonia (Moks et al., 2006), 28–29.8% in Spain (Segovia et al., 2001, 2003), and up to 47.1% in Latvia (Bagrade et al., 2009). Nabavi et al. (2014) found 1 of 4 wolves from Iran positive for *T. multiceps*. Interestingly, golden jackals (*Canis aureus*) from Italy as well as from the western part of Iran were negative for *T. multiceps* and *T. serialis* (Dalimi et al., 2006; Nabavi et al., 2014; Paoletti et al., 2017). In Serbia, however, *T. multiceps* was found in 1.6% of golden jackals, another wild carnivore of eastern Europe, the Middle East and Asia (Girovic et al., 2015b). These tapeworm species were absent in 80 brown bears in a *T. multiceps*/*T. serialis* endemic region in central Italy (Paoletti et al., 2017). Henke et al. (2002) described *T. multiceps* in coyotes in Texas but this report needs to be considered with care as it could be a morphological miss-diagnosis of *T. serialis*, and generally *T. multiceps* is assumed to be absent in North America.

- Metacestode infections (coenurosis) have also been reported from dead-end hosts (including dogs, horses, humans and other primates). *Taenia multiceps* coenurosis in humans and other primates represents a serious clinical condition. Supplemental Table 1 summarises cases of human *T. multiceps*-coenurosis, which were from Africa (*n* = 21), Europe (*n* = 17), Asia (*n* = 2), South America (*n* = 1), USA (*n* = 3) and the Middle East (Israel; *n* = 1). Where specified, the age of the patients was between 1 and 55 years (median: 33; SD: 17.0), showing that all age groups are potentially affected. Similar to infections in ruminants, the predominant infected tissue was the brain (*n* = 35/44), followed by intra-/periocular infections (*n* = 8/44) and rarely infections in the spinal cord (*n* = 1/44). Correspondingly, the infection was often fatal (42%) or ended with the loss of vision after eye removal (15%). Symptoms like partial paresis or paraplegia remained (12%), and only one third of the patients recovered partially or completely after surgery. However, in spite of the severity of the outcome of an infection in humans, coenurosis represents a rather minor risk for public health, given the relatively high infection pressure demonstrated by the wide occurrence and high prevalence of *T. multiceps* in canids but only 5 published cases in the past 25 years. Furthermore, there are no indications of immunodeficiency in these cases. Similarly, *T. multiceps* is rarely reported in other primates.
3.2.3. *Taenia serialis* coenurosis

The adult *T. serialis* tapeworm has a size of 20–72 cm and is morphologically very similar to *T. multiceps* but can be differentiated by hook number, size, and shape (Table 5). The life cycle of *T. serialis* includes canids as final hosts and hares and rabbits (rarely rodents) as intermediate host (Table 1), with Coenurus serialis in subcutaneous and muscle tissues, and occasionally in abdominal cavities (Paffenberger and Valencia, 1988; Verster, 1969). *Taenia serialis* is prevalent in North America, Europe and Africa. However, there are also recent reports of *T. serialis* in rabbits from Iran and China (Moshtari et al., 2018; Zhang et al., 2018). Furthermore, PCR confirmed *T. serialis* eggs in the faeces from 2 out of 1425 dogs from rural Australia (Jenkins et al., 2014). Hence, it is not surprising that this parasite has been occasionally found in Australian ring-tailed possums (*Pseudocheirus peregrinus*) and in kangaroos (*Macropus fuliginosus*) (Dunsmore and Howkins, 1968; Hough, 2000).

In contrast to *T. multiceps* with a predominantly domestic life cycle, wildlife plays a more prominent role in *T. serialis*. Surveys in the USA showed high prevalence throughout the nation, e.g. 12% (429 dogs) in Arizona and New Mexico and overall 3–29% in definitive hosts (Ing et al., 1998). With regard to intermediate hosts, 46% of 352 black-tailed jack rabbits (*Lepus californicus*) and 19% of a local rodent population were infected with *T. serialis* (Paffenberger and Valencia, 1988; Schantz et al., 1977). Henke et al. (2002) reported on *T. multiceps* in coyotes (*C. latrans*) from Texas but a misdiagnosis with *T. serialis* is probable in the absence of molecular species confirmation. Correspondingly, a recent study from Edmonton (Canada) found *T. serialis* in 13% out of 23 urban coyote carcasses (Luong et al., 2018). Interestingly, *T. serialis* was also found in arctic foxes (*Vulpes lagopus*) from north-east Greenland, demonstrating the widespread occurrence of this parasite in wild carnivores and the importance of a wildlife-cycle (Andreasen et al., 2017). In Europe, *T. serialis* occurs in regions where lagomorphs are present. In the UK, 0.3–0.6% of hunting dogs and 0.5% of 197 foxes from the same region have been found positive for *T. serialis*, respectively (Jones and Walters, 1992a, 1992b). Interestingly, *T. multiceps* was absent in the same fox population, despite being in a sheep raising region with observed cases of coenurosis. In Northern Ireland, up to 4% of the local fox population was infected with *T. serialis* (Ross and Fairley, 1969). On the European mainland, this cestode is rarely reported in foxes. Correspondingly, by examining large numbers of foxes from Germany, 0.15–2.3% of foxes were positive (Loos-Frank and Zeyhle, 1982; Pfeiffer et al., 1997; Welzel et al., 1995). In wolves, however, reported infection rates were 2.1% of 47 and 8% of 50 wolves in Spain (Segovia et al., 2001, 2003), 5.9% of 68 wolves in Portugal (Guerra et al., 2013), and 1% of 102 wolves in Serbia (Cirovic et al., 2013a). Hence, wolf migration might be a risk for the dissemination of *T. serialis* from the southern and eastern parts of Europe to new areas. In this sense, *T. serialis* was found in 1.1% of 447 golden jackals from Serbia, a species migrating into Central Europe (Cirovic et al., 2015b).

A survey in Germany could not find adult *T. serialis* in 84 badgers, 47 stone martens, and 387 cats (Loos-Frank and Zeyhle, 1982).

Rare cases of coenurosis in cats caused by *T. serialis*, but not by *T. multiceps* have been reported (Hayes and Creighton, 1978; Hsu et al., 1994; Orioles et al., 2014; Slocombe et al., 1989; Smith et al., 1988). Correspondingly, *T. serialis-coenurosis* represents a zoonosis with 11 reported cases found in literature (Supplemental Table 1). The predominant location of the coenuri was subcutaneous or intramuscularly on the trunk of the patient. While cerebral infections due to *T. serialis* have been reported in sheep (and cats), there are no such reported cases in humans. The lesions were presented as palpable swellings, which were usually painless. Where indicated, all patients recovered after surgical removal of the lesion.

Several cases of *T. serialis* coenurosis in non-human primates have been reported, in contrast to *T. multiceps* with only one described case with uncertain diagnosis (Table 4). Unlike in humans, cerebral, intrabdominal, intramuscular and subcutaneous infections have been observed. Interestingly, in a study between 1971 and 1972 in an Ethiopian national park, metacestodes of *T. serialis* have been identified morphologically in large visible subcutaneous swellings in 9.2% of 92 gelada baboons. In a follow-up study (1974–1975) observing the same animals, 10.6% were infected (Dunbar, 1980). Forty years later (2011), 4.8% of 291 other gelada baboons living in the same park had visible swellings, and in some of them, *T. serialis* was molecularly confirmed (Schneider-Crease et al., 2013). Unlike in other intermediate hosts, these visible asymmetric masses can grow very large in baboons (> 16 × 20 cm) and may show ulcerations, disturbing the animals obtrusively (Schneider-Crease et al., 2013; Schwartz, 1927). Potentially, these large lesions can develop over long time periods because of a long life span and a low predation rate of the infected animals. In fact, only 0.9% of juvenile baboons presented visible swellings, as compared to 9.9% of adults (Schneider-Crease et al., 2013). In spite of an endemic infection of primates, there were no human case-reports from Ethiopia, relativizing the zoonotic risk of *T. serialis*. Noteworthy, in contrast to e.g. *T. crassiceps* cysticercosis with single subcutaneous locations, many cases of *T. serialis-coenurosis* show multiple lesions, indicating a peroral infection and systemic parasite dissemination.

3.2.4. ‘African-type’ coenuroses

Reported cases of soft-tissue coenuroses in Africa are considered to be caused by *T. brauni* and *T. glomeratus* (syn. *T. glomerata*). *Taenia brauni* is a tropical tapeworm of Eastern and Northern Africa, Rwanda and the Democratic Republic of Congo, with a wild animal life cycle, including dog, fox, jackal and genet as definitive hosts and rodents such as the swamp rat, porcupine and gerbil as intermediate hosts (Collomb et al., 2007; Fain, 1956; Vanderlei et al., 1964). Coenuri in the brain only have been reported from rats (*Rattus r. rattus*) and non-human primates (*Fain, 1956*). *Taenia glomeratus*, which can be distinguished from *T. brauni* by careful morphological examination (Clapham, 1940a, 1940b), has only been reported from Nigeria and the Democratic Republic of Congo (Morel, 1959; Turner and Leiper, 1919). In humans, 27 cases of *T. brauni* and one case of *T. glomeratus* have been reported from Rwanda, Uganda, Nigeria, and Congo, based on morphological identification and geographical association (Supplemental Table 1). There is only one report in a non-human pri- mate (*Cercopithecus mitis doggetti*) from Rwanda with cysts in the brain, heart and parotid gland (Fain, 1956).

Overall, only limited information is available on these cestode species and any recent data is missing. It has to be considered that these species may be strains of *T. serialis*, based on the shared host range, tissue tropism, and clinical signs (Loos-Frank, 2000; Morel, 1959). However, there is no genetic data on *T. brauni* and *T. glomeratus*, and only a small number of published sequences for *T. multiceps* and *T. se- rialis*, which impair epidemiological studies (Collomb et al., 2007).

4. Diagnosis of cysticercosis and coenurosis

The tumor-like pathology of metacestode development in primates as dead-end hosts represents a challenge for the etiological diagnosis, which is mainly achieved only after necropsy, surgery or invasive diag- nostic interventions. Serological tests have not been developed or validated with the exception of *T. solium* cysticercosis. The morphologi- cal characteristics of metacestodes, especially size and form of the scolex hooks, allowed highly probable diagnoses on species level (Verster, 1969; Slais, 1973; Loos-Frank, 2000). However, incomplete or damaged larval stages without or with only a few scolex hooks might be present in clinical diagnostic material. Furthermore, their development in dead-end hosts might be atypical and lead to altered morphological characteristics. For example, Bamrungphol et al. (1972) described a juvenile strobilatetapeworm found in the spinal cord of a young man from Thailand, who had suffered from paraplegia for several months. The recovered worm (76 mm long with 300 segments) had a scolex with four muscular suckers and a rostellar with few remaining,
morphologically highly variable hooks. The authors concluded that “the immature state of the worm and the loss of many of the hooks from the scolex make its specific identification impossible”. However, they suggested that this strobilate form of a tapeworm found in extra-intestinal tissue likely belongs to the family Taeniidae.

Among the parasites discussed, three different metacestodes types exist, cysticercus, strobilocercus and coenurus (Table 5). These stages consist of a fluid-filled bladder of various sizes and forms, with a single (in the case of cysticerci or strobilocerci) or multiple (in the case of coenurus) invaginated scolex or scoleces. At the anterior end, scoleces carry a rostellum, a retractable muscular organ, that is equipped with four suckers and each a row of characteristic large and somewhat smaller hooks. *Taenia crassiceps* cysticerci are able to multiply locally and to produce daughter cysticerci by an exo- or endogenous budding process (see chapter 3.1). Cysticerci are whitish, translucent, oval bladders of 0.4 to more than 30 mm in diameter, depending on the species. At one pole, a clearly visible, white, non-translucent knob-like structure is present: the invaginated scolex. Daughter cysts usually develop at the opposite pole. More challenging is the morphological diagnosis of *Versteria* sp. cysticercosis, as this currently not well described species proliferates in the liver without many precise morphological characteristics (see chapter 2.2) and so far only 3 cases have been identified in humans based on molecular analyses (Table 3). The strobilocercus of *T. taeniaeformis* lives encapsulated in spherical cysts in the liver of its intermediate host. Isolated strobilocerci appear to be segmented and resemble adult tapeworms. However, the body has an unsegmented, flattened tube-like form, may reach several centimeters in length and carries a single scolex (Table 5). Coenuri are translucent, fluid-filled sacks of up to several centimeters in size that contain many scoleces or even clusters of scoleces (*Verster, 1969*; *Loos-Frank, 2000; Deplazes et al., 2016).

Characteristic structures that allow a tentative morphological identification include calcareous corpuscles, microtriches and scolex hooks. In HE stained histologic sections, calcareous corpuscles appear as small (5–15 μm) oval structures in the parenchyma of cestodes. The corpuscles are diagnostic for cestodes and are especially prominent in larval stages. They contain calcium carbonate whose role is not understood. The bladder wall may be smooth or have a warty appearance. Microvilli-like structures (microtriches) are present on the outer surface of the larval stages (*Ortibelt and Ash, 1995*). As shown for *T. solium*, they appear early in the development from the oncosphere to the cysticercus (Fig. 1) (*Chile et al., 2016*).

For a morphological species identification, number, form and size of the scolex hooks are important. However, reported size ranges of hooks are quite broad and do not allow for a definite species identification (Table 5). Moreover, many measurements refer to hooks of adults and some differences between adult and larval hooks and differences between the development in natural and in accidental and dead-end hosts may exist.

Today, identification of larval cestodes in clinical material is mainly based on the analyses of the nucleotide sequences of the mitochondrial 12S rRNA, ND1 (NADH dehydrogenase 1), and COI (cytochrome c oxidase subunit 1) genes or the nuclear ITS2 (ribosomal internal transcribed spacer 2) sequence (*Bowles et al., 1992; Bowles and McManus, 1993; von Nickisch-Rosenegk et al., 1999; Trachsel et al., 2007*) (Table 6). The same technique and in particular COI sequences have been broadly used for phylogenetic analyses (*Sharma et al., 2016*). Comparison of sequence data to published sequences in databases allow for a most reliable identification of cestodes, even if only small amounts of larval material is available, and this approach should be accepted as gold standard. However, sequence data especially of *T. brauni* and *T. glomeratus* are completely lacking, hindering the molecular identification in these cases. Therefore, efforts to produce and provide such data preferably based on morphologically characterized specimens should be intensified.

### Table 5

| Species         | Larval type and name | Size (mm) | Asexual replication | Number of scoleces (No) | Number of hooks (N) | Length of large hooks (μm) |
|-----------------|----------------------|-----------|---------------------|------------------------|---------------------|-------------------------|
| *T. crassiceps* | Cysticercus (Cysticercus longicollis) | 2–8 | Yes | One | 30–34 | 179–195 |
| *T. martis*     | Cysticercus          | 6–32      | No | No | 28–30 | 24–32 |
| *T. taeniaeformis* | Strobilocercus (Strobilocercus fasciolaris) | up to several cm | No | No | 4–5 | 24–42 |
| *T. multiceps*  | Coenurus (Coenurus cerebralis) | up to several cm | No | Yes | 18–24 | 14–16 |
| *T. serialis*   | Coenurus (Coenurus serialis) | 12–34 | No | Yes | 30–34 | 14–38 |
| *T. brauni*     | Coenurus (Coenurus brauni) | >1 cm | No | Many | 22–30 | 22–30 |
| *T. glomeratus* | Coenurus (Coenurus glomeratus) | 5.5–27.2 | No | Many | 28–34 | 30–34 |
| *Versteria* (Syn. *Taenia*) mustelae | Cysticercus | 0.4–2 | Yes | One | 30–74 | 14–38 |

a Cysts from an ape in various parts of the body (*Fain, 1956*).

b According to *Turner and Leiper* (1919); *Chile et al., 1996*.

c Observed with genetically not characterized North American isolates (*Freeman, 1996*).
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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ijppaw.2019.03.013.

Table 6

| Symbol (bold) and aliases | Primer designations | Primer sequence (5'-3') and approximate amplicon size | Reference and primer code [in squared brackets] as referred to in Tables 2–5 |
|--------------------------|--------------------|------------------------------------------------------|--------------------------------------------------------------------------|
| Target gene: Mitochondrial cytochrome c oxidase subunit 1 | mt-COI | JB3 | 5'-TTT TTT GGT CCG CAT CCG AGT GGT TAT-3' | Bowles et al. (1992). [P1] |
|                          |                   | JB4.5 | 5'-TAA AGA AAG AAC ATA ATG AAA ATG-3' | 529 bp |
|                          |                   | F/COI | 5'-TTG AAT TTT CCA CTT TTG AAT GC-3' | Nakao et al. (2000). [P2] |
|                          |                   | R/COI | 5'-GAA CCT AAC GAC ATA ACA TAA TGA-3' |  |
| Target gene: Mitochondrial NADH:ubiquinone oxidoreductase core subunit 1 | mt-ND1 | JB11 | 5'-AGA TTC GTA AGG GCC CTA ATA-3' | Bowles and McManus (1993). [P3] |
|                          |                   | JB12 | 5'-ACC ACT AAC TAA TTC ACT TTC-3' | 529 bp |
|                          |                   | NAD1-FF | 5'-ATT GCG TTT CAG ATG TTG TCT GAT TTA-3' | Eberwein et al. (2013). [P4] |
|                          |                   | NAD1-RR | 5'-CTC MCC ATA ATC AAA TGG ACT AGG-3' | 394 bp |
| Target gene: Mitochondrial 12S rRNA | mt-RNR1 | 12S | 5'-TTA AGA TAT GTG TAC AGG ATT AGA TAC |  |
|                          |                   | 60.fmr | 5'-AAC CGA GGG TGA CGG GGC GTG TGG ACC-3' | von Nickisch-Rosenegk et al. (1999). [P5] |
|                          |                   | 37S.rev | 5'-AAC CGA GGG TGA CGG GGC GTG TGG ACC-3' | 357 bp |
|                          |                   | Cest3 | 5'-GAA YTC TTT TTA GGG GAA GGT GTG-3' | Trachsel et al. (2007). [P6] |
|                          |                   | Cest5 | 5'-GCG GTG TGT ACM TGA GCT AAA C-3' | 267 bp |
|                          |                   | CES12SF | 5'-AGG GGA TAG GAC ACA GTG CCA GC-3' | Goldberg et al. (2014). [P7] |
|                          |                   | CES12SR | 5'-GCG TGT GTA CMT GAG YTA AAC-3' | 480 bp |
|                          |                   | 12S Cest FF | 5'-CAC AGT GCC AGC ATC YGC GGT-3' | Eberwein et al. (2013). [P8] |
|                          |                   | 12S Cest RR | 5'-GAG GGT GAC GGC CGG GTG GTA C-3' | 426 bp |
| Target gene: Ribosomal Internal Transcribed Spacer | ITS2 m2 | NC-2 | 5'-TTA GTT TCT TTT CCG CTG-3' | Gasser and Chilton (1995). [P9] |
|                          |                   | NC-6 | 5'-ATC GAC ATC TTG AAC GCA CAT TGC-3' | 600-700 bp |

Fig. 1. *Taenia crassiceps* metacestodes (case 8, Table 2). A. Cysticeri isolated at surgery (wet preparation, scale bar 5 mm). B. HE stained section of an invaginated scolex with suckers (S) and hooks (H) (scale bar 500 μm). C. Large and small hooks, calcareous corpuscles (wet preparation, scale bar 200 μm). D. Calcareous corpuscles in the parenchyma and microvilli like structures (microtriches) on the outer surface of a metacestode wall (HE stain, scale bar 20 μm).
Helmintological survey of the wolf (Canis lupus) in Estonia, with an emphasis on Echinococcus granulosus. J. Wildl. Dis. 42, 359–365.

Montibello Di Filippo, M., Meoli, R., Cavallerio, S., Eleni, C. De Libera, C., Berrilli, F., 2018. Molecular identification of Mesocestodes sp. metacestodes in a captive golden hamster (Mesocricetus auratus). Infect. Genet. Evol. 65, 399–405.

Morales, J., Velasco, T., Tovar, V., Fraguos, G., Feury, A., Beltrán, C., Villabonos, N., Aluja, A., Rodarte, L.F., Scuccio, E., Larralde, C., 2002. Castration and pregnancy of rural pigs significantly increase the prevalence of naturally acquired Taenia solium cysticercosis. Vet. Parasitol. 108, 41–48.

Morales-Montor, J., Baig, S., Hallal-Callejos, D., Camam, R.T., 2002. Taenia cricetis: andenongenesis of the host leads to protection during cestocercosis. Exp. Parasitol. 99, 209–216.

Mored, P., 1959. Les helminthes des animaux domestiques de l’Afrique occidentale. Rev. Elev. Med. Vet. Pay. 12, 153–174.

Morlo, P.L., Ballarta, J., Gilman, R.H., Legua, G., Rojas, M., Montes, G., 1998. Intestinal parasites of the Angora goat (Ovis aries capra) in the central Peruivan Andes. J. Helminthol. 72, 87–89.

Moshiri, A., Shamsian, S., Berenji, F., Jadidoleslami, A., Moghaddas, E., 2014. Coenurosis serails in northeastern Iran: a probable danger to human. J. Infect. 5, e1218.

Mouquet, G., Cambon, D., Diniego, V., Menerath, J.M., 1996. Intracerebral cestode larva in a 14-year-old boy in Auvergne. Presse Med. 25, 1168.

Nabavi, R., Manouchehri Naeini, K., Zebardast, N., Hashemi, H., 2014. Epidemiological studies of gastrointestinal helminthes of canids in chahmarahal and bakhtari province of Iran. Iran. J. Parasitol. 9, 276–281.

Nagy, A., Zadiolin, I., Schweiger, A., Schindler, M., Deplazes, P., 2011. Hair coat contamination with zoonotic helmunt eggs of farm pets and domestic foxes. Berl. Münchener Tierärztliche Wochenschr. 124, 503–511.

Nakao, M., Salo, Y., Yoku, F., Fuksanov, A., 2000. Mitochondrial genetic code in cestodes. Mol. Biochem. Parasitol. 111, 415–424.

Nakao, M., Lavikainen, I., Iwaki, T., Haukisalmi, V., Konyas, S., Oku, Y., 2013. Molecular phylogeography of the genus Taenia (Cestoda: Taeniidae): proposal for the recognition of a new species of the genus Versteria. Int. J. Parasitol. 43, 427–437.

Ntoukas, V., Tappe, D., Pfütze, D., Simon, M., Holzmann, T., 2013. Cerebral cysticercosis caused by larval Taenia cricetis tapeworm in immunocompromant woman. Germany. Emerg. Infect. Dis. 19, 2008–2011.

Nokutick, A.F., Deblock, S., Prost, C., 1990. Epidemiology of alveolar echinococcosis in central France. 1. Intestinal helminthes of the wolf, Canis lupus (Linnaeus, 1758) in the central Peruvian Andes. J. Helminthol. 72, 87–89.

Otranto, D., Cantacessi, C., Pfeffer, M., Dantas-Torres, F., Brianti, E., Deplazes, P., Genchi, C., Orihelo, T.C., Ash, L.R., 1995. Parasites in Human Tissues. ASCPPress, Chicago.

Orihelo, T.C., Orihelo, T.C., Ash, L.R., 1995. Parasites in Human Tissues. ASCCPPress, Chicago.

Orihelo, T.C., Orihelo, T.C., Ash, L.R., 1995. Parasites in Human Tissues. ASCCPPress, Chicago.

Orihelo, T.C., Orihelo, T.C., Ash, L.R., 1995. Parasites in Human Tissues. ASCCPPress, Chicago.

Pfeiffer, F., Kuschfeldt, S., Stoye, M., 1997. Helminth fauna of the red fox (Vulpes vulpes) in Germany. Emerg. Infect. Dis. 13, 223–228.

Pétabis, A.F., Deblock, D., Stock, S., 1980. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, S., Stock, G., 1990. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, S., Stock, G., 1990. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, S., Stock, G., 1990. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, D., Stock, S., 1980. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, S., Stock, G., 1990. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.

Pétabis, A.F., Deblock, S., Stock, G., 1990. Helminthes of the common fox (Vulpes vulpes) from the Netherlands. J. Helminthol. 55, 379–391.
Denmark. Helminthologia 28, 151–154.

Thompson, R.C.A., 1995. Biology and systematics of Echinococcus multilocularis. In: Thompson, R.C.A., Lymbery, A.J. (Eds.), Echinococcus and Hydatid Disease. CAB International, pp. 1–50.

Timbur, M., Sovonkubov, U., Zafar, S., Olesya, A., Kim, H.J., Yong, T., Lee, G.-J., Park, G.-M., Shin, M.-H., Yu, H.S., 2018. Genotypic and phylogenetic characteristics of Echinococcus granulosus sensu lato in Uzbekistan. In: Proceedings of the 14th international Congress of Parasitology, Daegu, Korea.

Tokw, T., Taira, K., Yamazaki, M., Kashimura, A., Une, Y., 2014. The first report of peritoneal tetrathyridiosis in squirrel monkey (Saimiri sciureus). Parasitol. Int. 63, 705–707.

Trachsel, D., Deplazes, P., Mathis, A., 2007. Identification of taeniid eggs in the faeces from carnivores based on multiplex PCR using targets in mitochondrial DNA. Parasitology 134, 911–920.

Tschni-Rosenegk, M., Lucius, R., Loos-Frank, B., 1999. Contributions to the phylogeny of the Cyclophyllidea (Cestoda) inferred from mitochondrial 12SrDNA. J. Mol. Evol. 48, 586–596.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Welzel, A., Steinbach, G., von Keyserlingk, M., Stoye, M., 1995. Zur Helminthenfauna des Rotfuchses (Vulpes vulpes L.) in Südniedersachsen. Z. Jagdwiss. 41, 100–109.

Verster, A., 1969. A taxonomic revision of the genus Taenia Linnaeus, 1758 s. str. Onderstepport J. Vet. Res. 36, 3–58.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Vuitton, L., Bresson-Hadni, S., Milton, L., 2015. Clinical epidemiology of human AE in Europe. Vet. Parasitol. 213, 110–120.

Wolff, M., Bush, M., Montali, R., Gardiner, C.H., 1989. Cysticercus pneumoniae and pleuritis in a red-ruffed lemur. J. Zoo Wildl. Med. 20, 383–385.

Wunschmann, A., Garilé, V., Averbeck, G., Kurtz, H., Hoberg, E.P., 2003. Cerebral cysticercosis by Taenia crassiceps in a domestic cat. J. Vet. Diagn. Invest. 15, 484–488.

Young, L.A., Morris, P.J., Keener, L., Gardiner, C.H., Stalis, I.H., Bicknell, E., Sutherland-Smith, M., Janssen, D.L., 2000. Subcutaneous Taenia crassiceps cysticercosis in a red ruffed lemur (Varecia variegata). B. J. Wildl. Dis. 36, 220–225.

Zhang, X.Y., Jian, Y.N., Ma, L.Q., Li, X.P., Karanis, P., 2018. A case of coenurosis in a wild rabbit (Lepus surmicus) caused by Taenia serrulata metacestode in Qinghai Tibetan plateau Aarea, China. Kor. J. Parasitol. 56, 195–198.