Current Status of Epidemiology and Diagnosis of Human Sarcocystosis

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Species of Sarcocystis are Apicomplexan parasites requiring intermediate and definitive hosts to complete their life cycle. Humans are one of many natural host species and may serve as both intermediate and definitive hosts. However, the extent and public health significance of human Sarcocystis infection are incompletely known. In this minireview, we provide an update on the epidemiology and diagnosis of human sarcocystosis and propose some tools that could contribute to a better understanding of the clinical significance and epidemiology of Sarcocystis infections.

Humans are natural hosts of multiple Apicomplexan genera and usually serve as either the intermediate (e.g., Toxoplasma, Plasmodium) or definitive (e.g., Cryptosporidium [monoxenous life cycle]) host. Meanwhile, Sarcocystis comprises Apicomplexan species with a two-host life cycle, including an intermediate host (prey) and a definitive host (predator), and humans may be both intermediate and definitive hosts for some of these species (1–3) (Fig. 1). Sarcocystis undergoes sexual reproduction in the intestinal epithelium, and oocysts are subsequently shed in host stool in definitive hosts, while the parasite forms the so-called “sarcocysts” in muscle tissue of distinct types, including striated, cardiac, and smooth muscle, in intermediate hosts. At least two species, namely, Sarcocystis hominis and Sarcocystis suihominis, exist with humans as definitive hosts and hence as potential causes of human intestinal sarcocystosis, with cattle and pigs as intermediate hosts, respectively. Humans can also be accidental intermediate hosts and develop muscular sarcocystosis.

Muscular sarcocystosis in the farming industry is a well-documented problem. A high prevalence of Sarcocystis infection is seen in cattle, pigs, and sheep in both developing and industrialized countries. Infection with Sarcocystis in the intermediate host is known to cause mortality, morbidity, abortions, lower meat yield, and meat condemnation due to visible sarcocysts (1, 2). Meanwhile, the extent and importance of human sarcocystosis remain less well described despite the potential for muscular or intestinal sarcocystosis to afflict a large number of people (3–6).

In this minireview, a short introduction to Sarcocystis taxonomy has been included. A systematic literature search has been performed on the prevalence of both intestinal and muscular sarcocystosis in humans. Lastly, diagnostic methods and their limitations are summarized with suggestions for development and application of molecular diagnostic tools to facilitate detection and a better understanding of the epidemiology and clinical significance of Sarcocystis infections of humans.

TAXONOMY

The taxonomy of Sarcocystis was updated in 1998 by Odening (7), who included 189 species. Since then, the number of species has increased, owing in part to an expanding interest in Sarcocystis in wildlife. The taxonomy of Sarcocystis is currently under revision; a common source of confusion has been the classification of identical Sarcocystis species from intermediate and definitive hosts as separate species. Traditionally, a new species of Sarcocystis is accepted when described in a new intermediate or definitive host (8). The sarcocyst wall structure is utilized for species identification due to the variability between Sarcocystis species (2); however, the wall structure is subject to change depending on the age of the sarcocyst (3) and so may have limited applicability as a morphological criterion. Molecular characterization combined with phenotypic description is critical to the ongoing revision of species in the genus Sarcocystis and is useful to study Sarcocystis transmission patterns in intermediate and definitive hosts, circumventing the need for expensive and laborious experimental infections (9, 10). This approach has been employed to some extent and recently led to the division of Sarcocystis in cervids into multiple species (10).

With the utilization of molecular methods for species identification, there is a need for guidelines as to what genes are useful for this purpose. Currently, the small-subunit (SSU) rRNA, cytochrome oxidase 1 (cox1), and internal transcribed spacer (ITS) genes are used for species identification (9, 10). There appear to be no guidelines, however, with respect to the amount of genetic diversity necessary to separate species and how to best analyze genetic diversity in terms of genetic markers, amount of DNA sequence data needed (e.g., sequence read length), and phylogenetic models. Currently, however, maximum-parsimony analysis of cox1 data appears to be the best possible method for delineation of closely related species, which may in part be due to the fact that cox1 DNA sequences are less difficult to align than SSU ribosomal DNA (rDNA) sequences (10).

With molecular confirmation pending, it is unknown whether Sarcocystis species other than S. hominis and S. suihominis can cause intestinal sarcocystosis (3). Until recently, the Sarcocystis species causing muscular sarcocystosis in humans were unknown (3, 5). Recently, 18S ribosomal DNA sequencing performed on a human muscle biopsy specimen containing Sarcocystis identified the species as Sarcocystis nesbitti (11). S. nesbitti has been described in muscle tissue of nonhuman primates (NHP), implying that...
NHP may indeed serve as reservoir hosts for *Sarcocystis* species causing muscular sarcocystosis in humans.

**EPIDEMIOLOGY AND CLINICAL SIGNIFICANCE**

Natural enteric infections with *S. hominis* and *S. suihominis* are described mostly as asymptomatic (4, 12, 13), although Yu (14) reported that 10% of infected individuals had symptoms associated with *Sarcocystis* infection. Meanwhile, experimental infections have been associated with gastroenteritis and eosinophilia (2, 3, 15, 16).

The prevalence of intestinal *Sarcocystis* infection in humans has been estimated to be between 1.1% and 10.4% in Europe, between 0.4% and 23.2% in Asia, 0.5% in Australia, and 0% in Argentina (Table 1). In general, caution must be taken when comparing prevalence figures reported in these studies due to the differences in diagnostic approaches. In addition, population bias may have been a factor, since the individuals in the study groups were not selected randomly from the population. A relatively high prevalence was observed in most of the studies, indicating that a large population was infected, but there is a need for further studies to confirm this. There is a tendency of recent studies to report prevalence figures lower than those reported in older studies (Table 1).

Potential geographical variation in the prevalence of intestinal sarcocystosis is difficult to assess due to many of the prevalence studies not being available in English, and so the methodology cannot be evaluated systematically.

Recent prevalence studies based on morphology and molecular data on *S. hominis* in cattle revealed that the occurrence of *S. hominis* infections is rather common, with prevalences of 6.2% to 97.4% (32–34), 53.5% (35), and 94% (16) in Europe, Asia, and South America, respectively. However, no cases of *S. hominis* infection were observed in a study performed in North America (36). Data on the prevalence of *S. suihominis* infection is scarce compared to those on the prevalence of *S. hominis* infection, and so recent prevalence studies estimating the prevalence of *S. suihominis* infection in pigs appear not to be available. Again, it should be emphasized that studies vary with regard to the type of sample (for instance, processed or unprocessed meat), type of muscle tissue examined, animal age, and number of samples analyzed. The fact that *S. hominis* or *S. suihominis* is regularly detected in intermediate hosts means that humans must be infected with these parasites.

Possible risk factors associated with acquiring intestinal *Sarcocystis* infection include eating raw or undercooked beef and pork and unsanitary disposal of human feces that can infect the inter-
mediate host and maintain the life cycle of Sarcocystis species causing intestinal sarcocystosis in humans (4).

Symptoms of muscular sarcocystosis include acute fever, myalgias, myositis, vasculitis, bronchospasm, and pruritic rashes (3, 5, 6, 37, 38). Muscular sarcocystosis has also been associated with nonspecific rheumatic diseases associated with myositis (39). No experimental-infection experiments have been performed with humans as intermediate hosts to our knowledge.

Treatment with albendazole alone or in combination with prednisone may lead to improvement in some cases (6, 37). However, as with intestinal sarcocystosis, there are currently no recommendations regarding treatment of muscular sarcocystosis, despite the severity in some cases, mainly due to lack of experience (3, 5).

The prevalence of muscular sarcocystosis in humans has been estimated to range between 0% and 3.6% in western countries and 21% in Malaysia from postmortem examination (Table 1). The high prevalence in Malaysia compared with western countries is likely due to NHPs being common in Malaysia, and humans can serve as intermediate hosts for the Sarcocystis species found in NHPs (40). Serological methods such as enzyme-linked immunosorbent assays (ELISA) and indirect fluorescent antibody tests (IFAT) have also been used to estimate the seroprevalence in Australia, the former Social Federal Republic of Yugoslavia, and Malaysia, producing seroprevalence figures of 34%, 22.5% to 44.4%, and 19.7%, respectively (Table 1) (25, 26, 28). The discrepancy between the higher prevalences observed in serological studies and those observed by direct detection in postmortem studies might be explained by low diagnostic sensitivity of direct detection and/or by low diagnostic specificity of serology. Another explanation could be persistence of a serological response after clearance of infection. There is a need for further studies to clarify this.

In 2004, Fayer (3) calculated the total number of muscular sarcocystosis cases described to be 92, but that many cases were probably unreported. Recently, the largest outbreak of human muscular sarcocystosis reported to date, with 100 people infected,

<table>
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<th>Reference and sarcocystosis category</th>
<th>Country location and study group</th>
<th>% prevalence value(s) (species; no. of samples examined)</th>
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<tr>
<td>Intestinal</td>
<td>Poland</td>
<td>10.4 (125)</td>
<td>Kato-Miur</td>
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<td></td>
<td>France</td>
<td>2.0 (S. hominis; 3,500); 0.7 (S. suihominis; 700)</td>
<td>Ethertype followed by flotation</td>
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<td></td>
<td>Germany, routine fecal samples from hospitals and clinics from patients with intestinal complaints</td>
<td>2 (403)</td>
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<td>Tibet in 3 counties</td>
<td>20.5, 22.5, and 22.9 (S. hominis; 926); 0.0, 0.6, and 7.0 (S. suihominis; 926)</td>
<td>Zinc sulfate flotation</td>
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<td></td>
<td>Laos</td>
<td>7.9 (767)</td>
<td>Kato-Katz &amp; merthiolate-iodine-formaldehyde concn</td>
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<td>Central Slovakia, workers from Vietnam</td>
<td>1.1 (1,228)</td>
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<td>Western Australia, Aboriginal communities</td>
<td>0.5 (385)</td>
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<td>Thailand, Thai laborers</td>
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<td>Thailand, rural communities</td>
<td>7.0 (1,603)</td>
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<td></td>
<td>Northeast Thailand, rural communities</td>
<td>0.4 (253)</td>
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<td>Muscular</td>
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<td>Malaysia, blood samples from blood donors</td>
<td>19.7 (243)</td>
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<td>Yugoslavia, blood samples obtained from cases suspected to be infected with Toxoplasma gondii</td>
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<td>Denmark, autopsy samples</td>
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<td>Australia, blood samples</td>
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<td>Malaysia, autopsy samples</td>
<td>21 (100)</td>
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<td>Australia, Sydney, autopsy samples</td>
<td>0 (50)</td>
<td>Two methods: (i) examination of formalin-fixed, paraffin-embedded sections from the psoas and diaphragm and (ii) digestion of (sieved and centrifuged) pepsin followed by microscopic examination</td>
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<td></td>
<td>Northern Ireland, autopsy samples</td>
<td>0 (50)</td>
<td>Examination of formalin-fixed, paraffin-embedded sections of tongue and diaphragm tissues</td>
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was investigated on Tioman Island, Malaysia (5). The outbreak has received attention due to the severity of disease and because the infected individuals were almost exclusively European tourists (5, 6). Also supported by the prevalence studies in Malaysia (25, 29), this indicates that the indigenous population is at substantial risk of contracting disease. In addition, an outbreak of muscular sarcocystosis was observed in 89 Malayan college students who visited Pangkor Island (38).

Little is known about the risk factors associated with acquiring muscular sarcocystosis. Humans have most likely been infected by consuming water or food contaminated with oocysts/sporocysts from definitive hosts (5, 6).

DIAGNOSIS OF INTESTINAL SARCOCYSTOSIS

A suspicion of intestinal sarcocystosis may be based on a history of gastroenteritis combined with habits of eating raw or undercooked meat that might contain infective Sarcocystis. The “diagnostic window” is limited to the time of oocyst/sporocyst excretion. The prepatent periods for S. hominis and S. suihominis are 14 to 18 days and 11 to 13 days, respectively, and oocyst/sporocyst excretion can last for at least 1 month (1–3). Importantly, this may imply that Sarcocystis may not be shed in feces in the acute stage of intestinal infection.

Methods for coprological examination include different flotation techniques (2, 3), modified Kato thick smear, the formalin-ether technique, and direct smear (22). The sensitivity of these methods is to a large extent unknown due the lack of studies addressing this specifically, but it should be possible to determine sensitivity by spiking samples with known amounts of oocysts/sporocysts. In animals, intestinal Sarcocystis infection is often detected postmortem by examination of mucosal scrapings rather than stool due to light fecal shedding (8).

Oocysts of S. suihominis measure 12.3 to 14.6 μm by 18.5 to 20.0 μm; fewer detailed data are available for oocysts of S. hominis, which may measure 20 to 23 μm in diameter. Sporocysts of S. hominis average 9.3 by 14.7 μm, and those of S. suihominis average 10.5 by 13.5 μm (3, 41). Both oocysts and sporocysts may be seen in fecal samples from definitive hosts; the oocyst wall may be difficult to discern (Fig. 2), but the use of stains such as periodic acid-Schiff stain may enable its detection. Given the shape and size of the oocysts and sporocysts, there should be little risk of confusing them with oocysts of Cryptosporidium, Cyclospora, and Cystoisospora. Still, a number of predicaments may impede our ability to detect Sarcocystis in human fecal samples on the basis of traditional parasitological methods and conventional reasoning. The relatively long prepatent period has already been highlighted, and it is reasonable to believe that fecal samples are sometimes being analyzed from patients with gastroenteritis before oocyst shedding commences. Moreover, oocysts/sporocysts do not consistently take up acid-fast stain (Fig. 2), and so they may be missed by the Ziehl-Neelsen staining commonly used for traditional parasitological detection of Apicomplexan parasites in stool. Moreover, in a recent study, we showed that none of 16 of 889 fecal samples positive for Cryptosporidium by real-time PCR were post-
itive by Ziehl-Neelsen staining, which was probably due to the presence of only light infections, as evidenced by high threshold cycle (CT) values (42). Since fecal shedding of Sarcocystis is presumably usually light, the parasite may therefore easily be overlooked in Ziehl-Neelsen preparations, and given the variability in uptake of acid-fast stain, the Ziehl-Neelsen method should not be recommended for detecting Sarcocystis.

To detect Sarcocystis in stool, a DNA detection-based approach therefore appears relevant. Real-time PCR assays have been published for several intestinal protozoa, but so far, none has been published for Sarcocystis. This may be due to a variety of reasons. Generally, if laboratory methods fail to enable detection of a given parasite, the parasite will probably be thought of as rare, and so the incentive to develop DNA-based methods for its detection may remain low. More specifically, regions of the small-subunit (SSU) rRNA gene conserved across the Sarcocystis genus may be so genetically similar to those found in competing templates in fecal genomic DNA that the applicability of this marker in genus-specific PCR analyses performed on DNA extracted directly from feces is reduced due to decreased PCR specificity and sensitivity. Nevertheless, the first molecular diagnostic assay to diagnose DNA of oocysts/sporocysts of Sarcocystis in fecal samples was developed only recently. The described method included seminested amplification of the SSU rRNA gene with subsequent restriction fragment length polymorphism (RFLP) analysis, making species differentiation possible in cattle (43). The primers target conserved regions; however, the forward primer “18s 2L” given in a paper by Xiang et al. (43) and used in both first and second PCR reads 5’-GGATAAACCGTGGTAATTCT ATG-3’, while the SSU rDNA template reads 5’-GGATAACCC GTGGTAATTCTATG-3’. The two primer combinations used in the nested PCR appear relatively nonspecific, and in silico analysis suggests that they may amplify a variety of Apicomplexan parasites. A different gene target, cytochrome oxidase I (cox1), exhibits extensive genetic diversity and can be used to discriminate between even closely related species of Sarcocystis (10). Unfortunately, cox1 DNA sequences are unavailable for the two species commonly thought to be involved in human sarcocystosis, namely, S. hominis and S. suihominis. Once those sequences are available, and given the amount of sequence data in GenBank available for multiple species of Sarcocystis, it should be possible to develop diagnostic species- and group-specific PCRs based on the cox1 gene in a real-time or nested PCR format to increase sensitivity. Relevant pretreatment of fecal samples could include purification and concentration of oocysts using gradient (zinc sulfate, sodium chloride, sucrose, etc.) methods, inherently increasing the chances of detecting other single-celled parasites of similar densities as well, with subsequent microscopy and DNA extraction of the meniscus. At present, nearly full-length SSU rDNA sequences (~1.9 kbp) and partial cox1 sequences (~1.0 kbp) of about 40 and at least 20 Sarcocystis species, respectively, are available (10), potentially facilitating the design of relevant primers and probes for conventional or quantitative PCR-based detection of oocysts/sporocysts in feces.

**DIAGNOSIS OF MUSCULAR SARCOCYSTOSIS**

Muscular sarcocystosis is suspected when compatible symptoms are present and especially if traveling has occurred to places where native wildlife includes NHPs. The identification of sarcocysts in muscle is considered the sole definitive diagnostic criterion (5). It is, however, rarely possible to obtain a biopsy specimen containing sarcocysts due to both issues related to obtaining consent from the infected individual and to the low chance of retrieving a specimen that contains sarcocysts.

ELISA and IFAT methods are used in serodiagnosis of muscular sarcocystosis (39, 44). There has been no report of cross-reactivity with, e.g., *Toxoplasma*, indicating that methods in use have high specificity (44). One IFAT method showed complete agreement with morphological examination of freshly slaughtered cattle (45). Similar prevalence figures in humans were also obtained in Malaysia, where Wong and Pathmanathan (29) used microscopy to examine tongue tissue while Thomas and Dissanaike (25) used IFAT, reporting prevalences of 21% and 19.7%, respectively. However, the two studies were performed on two different groups of people. Ideally, ELISA and IFAT methods should not be able to detect intestinal sarcocystosis and classify it as muscular sarcocystosis (44); however, the high prevalence figures observed in the former Socialist Federal Republic of Yugoslavia and Australia where no species of NHP is autochthonous indicate that this may not be the case (26, 46). A Western blot analysis for the confirmation of muscular sarcocystosis is currently under development (5).

An inherent problem with the utilization of antibodies as a principle of diagnosis is the lack of differentiation between early-stage infection and late-stage infection (47). This is of particular importance in relation to the development and clearance of sarcocysts in humans. Among animals, an ELISA enabled detection of IgM in acute-phase infections in mice and sheep; however, the IgG response was detected earlier than the IgM response in infections in pigs (48). No data on IgM development in humans are available. A different approach to detecting acute muscular sarcocystosis using nested PCR on blood samples containing circulating merozoites has been developed for testing in sheep (47). Molecular studies are warranted to identify the species that can cause muscular sarcocystosis to enable making epidemiological inferences that can be used to assess the extent and importance of human muscular sarcocystosis plus the modes of transmission.

**CONCLUDING REMARKS**

Diagnostic limitations have impeded our ability to increase our understanding of the epidemiology and public health significance of *Sarcocystis* infections in humans. Due to light shedding in feces, PCR-based methods for detection of DNA from oocysts/sporocysts in feces appear relevant; multiple species of *Sarcocystis* have now been molecularly characterized, and available sequence data should greatly enhance the possibilities for development of relevant diagnostic PCR assays and assays serving to map host specificity and transmission modes. Meanwhile, direct detection of the parasite in cases of human muscular sarcocystosis is hampered by the apparent absence of a predilection site, and so indirect detection by serological methods is preferred to testing of tissue biopsy specimens.

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