



Review

Human toxocariasis – A look at a neglected disease through an epidemiological ‘prism’



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ABSTRACT

Toxocariasis, a disease caused by infection with larvae of *Toxocara canis*, *T. cati* and/or congeners, represents clinical syndromes in humans including visceral and ocular larva migrans, neurotoxocariasis and covert/common toxocariasis. It is reported to be one of the most widespread public health and economically important zoonotic parasitic infections that humans share with dogs, wild canids, including foxes, and possibly other mammals. Humans become infected by accidental ingestion of embryonated *Toxocara* eggs, or larvae from tissues from domestic or wild paratenic hosts. Most infections are asymptomatic, and human disease may go unnoticed, as clinical investigation is often not pursued and/or diagnostic testing not conducted. Sometimes toxocariasis can be associated with complications, such as allergic and/or neurological disorders, possibly including cognitive or developmental delays in children. There is no anti-toxocariasis vaccine, and chemotherapy in humans varies, depending on symptoms and location of larvae, and may include the administration of albendazole or mebendazole, together with anti-inflammatory corticosteroids. Some recent studies indicate that toxocariasis is having an increased, adverse impact on human health in some, particularly underprivileged, tropical and subtropical communities around the world. Although tens of millions of people, especially children, are expected to be exposed to, or infected with *Toxocara* species, there is limited precise epidemiological data or information on the relationship between seropositivity and disease (toxocariasis) on a global scale. To gain an improved insight into this area, the present article reviews salient clinical aspects of human toxocariasis and the epidemiology of this disease, with particular reference to seroprevalence, and discusses future research and approaches/measures to understand and prevent/control this socioeconomically important, yet neglected zoonosis.

1. Introduction

Toxocariasis results from the transmission of *Toxocara* species from carnivores, including canids and felids, to humans (Dantas-Torres and Otranto, 2014; Macpherson, 2013; Nichols, 1956; Wilder, 1950). *Toxocara* adults live in the small intestines of wild or domestic definitive hosts. The geographically most widespread and important zoonotic species, *T. canis*, infects a wide variety of canids, including dogs, foxes, wolves, jackals and coyotes, whilst *T. cati* and *T. malaysiensis* infect felids (Fisher, 2003; Gibbons et al., 2001; Schnieder et al., 2011). Infected definitive hosts excrete unembryonated *Toxocara* eggs in faeces,

which contaminate the environment and/or the hosts' hair. Under favourable temperature and humidity conditions, eggs embryonate over weeks to months and can remain infective in the environment for months to years (El-Tras et al., 2011; Overgaauw et al., 2009; Roddie et al., 2008).

Toxocara eggs containing infective third-stage larvae (L3s) are often accidentally ingested by humans from contaminated food, water and/or the environment (soil or sand). In the small intestines of definitive hosts, L3s emerge from the eggs, migrate through the intestinal wall and then via the circulatory system to organs including liver, lungs, central nervous system and/or musculature; L3s can undergo arrested

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development within tissues for at least several months (Schnieder et al., 2011). Infective eggs ingested by transport or paratenic hosts, including rodents (mice and rats), lagomorphs (rabbits), ruminants (cattle), suids (swine) or birds (chickens), undergo a similar fate, with L3 s migrating to and then arresting in tissues (cf. Strube et al., 2013). If L3-infected tissues from paratenic hosts are food sources for humans and definitive hosts, ingestion can result in infection (Schnieder et al., 2011; Strube et al., 2013). The consumption of infected raw or undercooked ruminant liver has been implicated in human toxocariasis (Salem and Schantz, 1992; Yoshikawa et al., 2008).

Thus, humans are infected via a variety of routes, and it is thought that children usually become infected by accidentally ingesting infective eggs of *Toxocara* from the environment or, occasionally, by eating invertebrates, such as earthworms (Cianferoni et al., 2006), whilst some people become infected by ingesting the tissues of infected vertebrate (paratenic) hosts. Ingested larvae penetrate the intestinal wall, invade various tissues and cause immune and inflammatory responses that can lead to symptoms including fever, headaches, coughing, and abdominal or limb pain (Pawlowski, 2001; Rubinsky-Elefant et al., 2010; Taylor et al., 1988). The clinical syndromes of human toxocariasis include: visceral larva migrans (VLM), ocular larva migrans (OLM), neurotoxocariasis (NT) (e.g., eosinophilic meningoencephalitis) and covert/common toxocariasis (CT) (Finsterer and Auer, 2013; Nicoletti, 2013; Rubinsky-Elefant et al., 2010).

Despite the public health and clinical significance of human toxocariasis, particularly in tropical and subtropical regions of the world and in underprivileged communities in temperate climatic zones (Fialho and Corrêa, 2016; Hotez and Wilkins, 2009; Lee et al., 2014; Macpherson, 2013; Overgaauw, 1997; Rubinsky-Elefant et al., 2010), considerable knowledge gaps in the epidemiology of this disease exist (Çelik et al., 2013; Cooper, 2008; Holland, 2015; Hotez and Wilkins, 2009; Le et al., 2016; Poulsen et al., 2015; Smith et al., 2009). Recent studies indicate that toxocariasis is neglected, and it is not a notifiable disease (Fu et al., 2014; Hayashi et al., 2005; Won et al., 2008). In some communities, despite the humane reduction or treatment of canids and felids to decrease environmental contamination with infective eggs, the anti-*Toxocara* serum antibody prevalence in humans can remain high (Fu et al., 2014; Hayashi et al., 2005).

This review examines the clinical features and epidemiology of human toxocariasis, and discusses future research needs and approaches/measures to understand and to prevent/control this important and widespread zoonosis.

2. Toxocariasis – clinical forms and treatment options

Of the four main clinical forms of toxocariasis (Fig. 1), although relatively uncommon, VLM is the most advanced form of human toxocariasis, mostly affecting children, being associated with signs such as coughing, wheezing, myalgia and/or cutaneous abnormalities (e.g., eczema, pruritus, rash and/or vasculitis) (Despommier, 2003; Gavignet et al., 2008). In addition, VLM cases can present with

lymphadenopathy, hepatitis, myocarditis, nephritis and/or arthritis (Despommier, 2003; Kuenzli et al., 2016; Pawlowski, 2001). Asthma and pulmonary fibrosis can be associated with VLM (Aghaei et al., 2018; Cooper, 2008). Eosinophilia and elevated IgE levels are common. Cerebritis can also occur as a result of larval invasion into the central nervous system to produce neurotoxocariasis (NT), although NT can also present as a separate syndrome (see below).

OLM refers to human toxocariasis only affecting the eye. It is relatively uncommon, but mostly reported to occur in children between three and 16 years of age (Despommier, 2003; Good et al., 2004; Pivetti-Pezzi, 2009). Uni-ocular visual impairment might occur, accompanied by chronic endophthalmitis/retinitis and/or posterior/peripheral granulomata (Despommier, 2003). The extent of this impairment is linked to migrating or dead larvae and/or resultant immune reactivity against them as well as location in the eye (Pivetti-Pezzi, 2009). Blindness can result from tractional retinal detachment, vitritis and/or cystoid macular oedema (Martínez-Pulgarin et al., 2015; Pivetti-Pezzi, 2009).

NT is reported to occur mainly in middle-aged people and less in children (Deshayes et al., 2016). Signs relate to the larval migration in the central nervous system and ensuing meningitis, encephalitis, cerebral vasculitis and/or myelitis, usually relating to relatively non-specific clinical signs, such as headaches and fever (Caldera et al., 2013; Deshayes et al., 2016; Finsterer and Auer, 2007; Vidal et al., 2003). There has been discussion in the literature about possible associations between cerebral toxocariasis and neurodegenerative disorders, including seizure, schizophrenia, cognitive deficits, idiopathic Parkinson's disease and/or dementia (cf. Çelik et al., 2013; Fan et al., 2015; Finsterer and Auer, 2007). There is particular concern about cognitive or developmental delays in children in socioeconomically disadvantaged communities (Walsh and Haseeb, 2012). Some studies suggest that NT might represent a cause of poor school performance and other psychological disturbances in this cohort (Hotez, 2014). Although associated with logistical and ethical challenges, there is a need for epidemiological studies of the relationship between *Toxocara* and cognitive deficits. Involvement of the peripheral nervous system, reflected in radiculitis, inflammation of cranial nerves and/or skeletal muscles, is rarely recorded in humans (Finsterer and Auer, 2007). In a meta-analysis of published information/data representing 2159 people with epilepsy and 2581 people without epilepsy (Luna et al., 2018), it was shown that seropositivity to *Toxocara* was significantly associated with epilepsy (odds ratio [OR], 1.69; 1.42 to 2.01).

CT, representing 'common toxocariasis' in adults and 'covert toxocariasis' in children, is challenging to diagnose clinically because of non-specific symptoms (Taylor et al., 1988). Clinical signs, such as weakness, pruritus, rash, pulmonary dysfunction, pulmonary insufficiency and/or abdominal pain can be seen in adults, whereas fever, anorexia, headache, nausea, abdominal pain, vomiting, wheezing, lethargy, sleepiness and behavioural disorders, pulmonary symptoms and/or limb pain, might be observed in children (Walsh and Haseeb, 2014). Eosinophilia and elevated IgE levels are common. There appears to be an association between CT and asthma, although some investigators might place developmental and cognitive delays in this category rather than NT (Hotez, 2014; Li et al., 2014; Mendonça et al., 2012). These clinical characteristics can associate with moderate to high anti-*Toxocara* serum antibody titres (Pawlowski, 2001; Rubinsky-Elefant et al., 2010; Taylor et al., 1988). A meta-analysis (Aghaei et al., 2018) inferred an increased risk for asthma in children who were seropositive to *Toxocara* (OR, 1.91; 1.47 to 2.47). Another such analysis (Mohammadzadeh et al., 2018) revealed an increased risk of allergic skin disorders in individuals who were seropositive to *Toxocara* (OR, 1.75; 1.16 to 2.64), with seropositivity being significantly associated with urticaria (OR, 2.97; 1.53 to 5.76) but not with atopy (OR, 1.08; 0.55 to 2.15) or eczema (OR, 1.62; 0.95 to 2.78).

The chemotherapy of human toxocariasis is challenging; some drugs do not reach and penetrate *Toxocara* larvae in tissues and their efficacy

Visceral larva migrans (VLM)	Coughing, wheezing, pruritus, rash, lymphadenopathy, hepatosplenomegaly, nodules myocarditis, nephritis, eosinophilia present
Ocular larva migrans (OLM)	Chronic endophthalmitis, retinitis, posterior/peripheral granulomata, Uni-ocular visual impairment, blindness
Neurotoxocariasis (NT)	Fever, headache, meningitis, encephalitis, cerebral vasculitis, myelitis, epilepsy, neurodegenerative disease, cognitive and developmental delays
Covert toxocariasis (CT)	Nonspecific symptoms, asthma and developmental delays accompanied with eosinophilia and elevated IgE

Fig. 1. The range of clinical syndromes associated with human toxocariasis.

cannot be rigorously assessed in patients. Nonetheless, anthelmintic treatment is recommended for acute toxocariasis, principally to prevent larvae from reaching the central nervous system (Pawlowski, 2001; Wiśniewska-Ligier et al., 2012). Despite limited efficacy, albendazole and mebendazole are commonly used for oral treatment of VLM (Caumes, 2003), albeit these benzimidazoles are poorly soluble, poorly absorbed and undergo extensive, first-pass metabolism in the liver. Albendazole is preferred due to its distribution (of a metabolite) across distinct tissues compared with mebendazole. In humans, albendazole is absorbed better than mebendazole (bioavailability of $\leq 2\%$, following a single oral dose); also, albendazole passes through the blood-brain barrier, and is better tolerated than mebendazole (Othman, 2012). In addition, nonsteroidal anti-inflammatory drugs or corticosteroids, such as dexamethasone or prednisolone, can be administered to reduce clinical signs linked to allergic responses to *Toxocara* antigens (Despommier, 2003). However, although treatment with albendazole and steroids may result in a favourable clinical outcome (Barisani-Asenbauer et al., 2001; Martínez-Pulgarin et al., 2015), the efficiency of such treatment of OLM (e.g., following ophthalmologic surgery) is unclear (Despommier, 2003). Although animal experiments (C57BL/6 mice) have indicated promising efficacy of fenbendazole formulations at killing *T. canis* larvae in musculature (89.5–100%) and/or brain (66.1%) (Hrčková et al., 2007; Leonardi et al., 2009), there is a major need for well-controlled evaluations in other animals. Liposome-encapsulated and polyethylene glycol (PEG)-conjugated compounds show promise for treatment (Barrera et al., 2010; Horiuchi et al., 2005; Hrčková and Velebný, 2001; Hrčková et al., 2007; Moreira et al., 2014), but detailed evaluations are needed. There is also an urgent need for clinical studies in people, particularly children, to evaluate whether new drug formulations can relieve persistent symptoms, such as headache (cf. Wiśniewska-Ligier et al., 2012). Although other synthetic chemicals and natural products seem to show larvicidal or larvostatic activity (e.g., linoleylpyrrolidilamide, phenazines, tribendimidine and nitazoxanide; extracts from *Chenopodium* sp., and Nutridesintox®), they require extensive future investigation to assess their potential against *T. canis* and *T. cati*.

3. Diagnosis of infection and disease, and its challenges

In humans, the diagnosis of *Toxocara* infection or toxocariasis is usually achieved by serology, occasionally combined with imaging methods to detect granulomata (with encapsulated larvae) in tissues. Methods such as enzyme-linked immunosorbent assays (ELISAs), utilising *Toxocara [canis]* excretory/secretory (TES) antigens (de Savigny et al., 1979), have been used for seroprevalence studies of *Toxocara* infection/exposure in humans (Elefant et al., 2016; Jin et al., 2013; Magnaval et al., 1991; Mohamad et al., 2009; Noordin et al., 2005; Peixoto et al., 2011; Roldán et al., 2015; Watthanakulpanich et al., 2008; Table 1). ELISA results can be verified by Western blotting to reduce false-positive results relating to background- or cross-reactivity

Table 1

Key serological/immunological methods in current use for the detection of *Toxocara* infection(s) or exposure, and/or the diagnosis of toxocariasis, and brief description of purpose and performance.

Methodology	Purpose and performance	References
IgG-TCLA-ELISA	Detects specific total IgG in human serum; 92% sensitivity; 87% specificity	Jin et al. (2013)
IgG-TES-WB	Detects specific total IgG in human serum; “high” specificity; “minor” cross-activity	Magnaval et al. (1991)
IgG-TES-ELISA	Detects specific total IgG/IgG subclass in human serum; 60–98% sensitivity; 36–81% specificity	Noordin et al. (2005) Watthanakulpanich et al. (2008) Peixoto et al. (2011)
IgG-dTES-WB	Detects specific total IgG in human serum; no cross-reactivity with 32, 55 and 70 kDa fractions of dTES	Roldán et al. (2015)
IgG-dTES-ELISA	Detects specific total IgG in human serum; 100% sensitivity and specificity	Roldán et al. (2015)
IgG4-rTES-ELISA	Detects specific IgG4 subclass in human serum; 93% sensitivity and “increased” specificity	Mohamad et al. (2009)
IgG-DiM-BSA-ELISA	Detects specific total IgG in human serum; 92% sensitivity; 95% specificity	Elefant et al. (2016)

IgG: immunoglobulin G; TCLA: crude antigens from *T. canis* larvae; ELISA: enzyme-linked immunosorbent assay; TES: *Toxocara canis* excretory/secretory antigens; WB: Western blot; dTES: deglycosylated TES antigens; rTES: recombinant TES antigens; DiM-BSA: di-O-methylated coupled bovine serum albumin.

with other parasites (Moreira et al., 2014). This combined approach is preferable (Moreira et al., 2014), but it is important to emphasise that current *Toxocara* infection cannot be unambiguously differentiated from past infection or exposure (Holland, 2015; Smith et al., 2009). Clearly, problems exist with cross-reactivity in some serodiagnostic tools, leading to false-positive results, especially in geographic areas where polyparasitism is prevalent (Fillaux and Magnaval, 2013; Moreira et al., 2014). Moreover, it is not possible to unambiguously distinguish between *T. canis* and *T. cati* (or possibly *T. malayensis*) infections using current serological or immunological techniques (cf. Poulsen et al., 2015; Schabussova et al., 2007), such that optimised, species-specific diagnostic tools are required to support future epidemiological investigations.

The sensitivity and specificity of serodiagnostic tests for human toxocariasis depend on the antigens used (e.g., somatic products from *T. canis* larvae, native or recombinant TES products, or either glycan antigens or deglycosylated TES), the isotype of antibodies (e.g., IgG, IgG subclass or IgM) being detected and the extent of optimisation of the assays (Peixoto et al., 2011; Roldán et al., 2015). Thus, numerous ELISAs have been established and evaluated including those listed in Table 1. Assays for the specific detection of IgG3 and IgG2 anti-*Toxocara* antibodies are reported to achieve increased sensitivity and specificity (Magnaval et al., 1991). By contrast, an IgG4-ELISA, employing recombinant proteins (rTES-120 and rTES-30) was reported to achieve 100% sensitivity, and an IgG-ELISA using deglycosylated TES is claimed to have 100% sensitivity and specificity (Mohamad et al., 2009; Roldán et al., 2015). Other recombinant proteins have also been assessed as diagnostic antigen candidates (Anderson et al., 2015; Zhan et al., 2015). Although IgG antibody avidity or eosinophil cationic protein levels can be measured to suggest current or past *Toxocara* infection (Boldiš et al., 2015; Dziemian et al., 2008; Hübner et al., 2001; Magnaval et al., 2001), each assay needs to be critically evaluated.

Especially for the diagnosis of OLM, specific anti-*Toxocara* antibody in blood (although false-negative results are common) and vitreous or aqueous humour should be assessed, while the detection of specific antibodies in serum as well as eosinophils in cerebrospinal fluid (CSF) need to be evaluated for the diagnosis of NT (Despommier, 2003; Macpherson, 2013). In addition to serology, imaging techniques can assist clinical diagnosis in patients with VLM or OLM; this might involve the use of computed tomography (CT), magnetic resonance-imaging (MRI) and/or ultrasound to scan liver lesions relating to VLM, and fluorescein angiography, optical coherence tomography (OCT), ocular ultrasound and/or CT can also be employed to support OLM diagnosis (Macpherson, 2013; Strube et al., 2013). However, better diagnostic algorithms and clinical definitions are needed for both CT and NT.

In paratenic or accidental hosts, the diagnosis of toxocariasis/*Toxocara* infection can be achieved by histopathological examination, morphometric assessment of larvae (if detected) and/or the specific detection of larval DNA in/from tissues or body fluid samples by molecular means (Fillaux and Magnaval, 2013; Gasser, 2013). As the

results from serological tests alone do not allow for an unequivocal diagnosis of infection (Poulsen et al., 2015; Smith and Noordin, 2006), polymerase chain reaction (PCR)-based tools, using genetic markers in the first (ITS-1) and second (ITS-2) internal transcribed spacers (ITS) of nuclear ribosomal DNA (rDNA) or select mitochondrial (mt) genes, can achieve specific diagnosis (e.g., Gasser, 2006, 2013; Jacobs et al., 1997; Jex et al., 2008; Li et al., 2008; Wickramasinghe et al., 2009), and are employed for epidemiological, population genetic and/or systematic investigations. Importantly, these tools allow *Toxocara* species to be identified molecularly and can be used to discover “new” species, such as *T. malaysiensis* (Gibbons et al., 2001; Zhu et al., 1998), whose transmission patterns and zoonotic potential are unknown. Studies demonstrating the specific detection of *Toxocara* DNA by PCR from cerebrospinal fluid from humans or bronchoalveolar lavage from mice show promise for future complementary diagnostic tools (Caldera et al., 2013; Pinelli et al., 2013).

4. Epidemiology and global health significance

Toxocara canis and *T. cati* have a worldwide distribution (Fisher, 2003; Lee et al., 2010). Some nation-wide studies of these nematode species in dogs and cats estimated respective prevalences of 6.1% and 4.7% in Germany (Barutzki and Schaper, 2011), ~4.5% in the Netherlands (Overgaauw et al., 2009), and 1.2%–3.2% in Australia (Palmer et al., 2008), with marked prevalences (51–100%) in puppies and in adult dogs (1–45%) and cats (3.2–91%) in some surveys in countries including Portugal, Nigeria, India and China (Dai et al., 2009; Fisher, 2003; Lee et al., 2010; Sowemimo, 2007; Traub et al., 2005; Waap et al., 2014). It appears that there are > 77 million dogs and 93 million cats in the USA (Gompper, 2013), some of which are expected to play a key role in the dissemination of toxocarasis via the excretion of eggs in faeces into playgrounds, sand pits, gardens, parks and beaches, which represents a substantial risk factor of human infection if eggs survive and then become infective in the environment (Fisher, 2003; Manini et al., 2012).

Humans acquire *Toxocara* infection(s) by accidentally ingesting infective eggs from contaminated soil, water, fruit or vegetables (e.g., Dubná et al., 2007; Poepl et al., 2013). Humans may also become infected through direct contact with dogs or cats (Poepl et al., 2013) – although eggs have been detected on the hairs of definitive hosts (El-Tras et al., 2011; Roddie et al., 2008; Overgaauw et al., 2009), the numbers of embryonated eggs on well cared-for dogs are negligible (Holland, 2015; Keegan and Holland, 2010). People can also become infected by ingesting larvae present in raw or undercooked meat or organs from paratenic hosts, such as rabbits, ruminants and poultry (cf. Dutra et al., 2014; Salem and Schantz, 1992; Taira et al., 2004; Yoshikawa et al., 2008). The apparent broad geographic distribution of *Toxocara* and multiple transmission routes indicate that toxocarasis is a common helminth infection in humans (Fig. 2; Appendix Table 1).

4.1. Recent estimates of the prevalence of *Toxocara* eggs in public places worldwide

Although many investigations had indicated that environmental contamination with *Toxocara* eggs is a major risk factor for human toxocarasis, there was, until recently, no detailed analysis of published information. For this reason, we carried out the first systematic review and meta-analysis to evaluate the global prevalence of *Toxocara* eggs in public places, including playgrounds, parks and beaches (Fakhri et al., 2018). We identified 109 peer-reviewed studies by searching four public databases (Embase, PubMed, Science Direct and Scopus); in these studies, a total of 42,797 soil samples from 40 countries around the world were tested for the presence of *Toxocara* eggs using conventional diagnostic methods (usually flotation). The results revealed that 21% (range: 16–27%) of public places were contaminated with *Toxocara* eggs, with the highest level being in the Western Pacific

region (mean: 35%; range: 15–58%) and the lowest in the North and Central Americas (mean: 13%; range: 8–23%). In other WHO regions, contamination (i.e. prevalence) rates varied from 3 to 49%: Africa (mean: 27%; range: 11–47%), South America (mean: 25%; range: 13–33%), South-East Asia (mean: 21%; range: 3–49%), Middle East and North Africa (mean: 18%; range: 11–24%) and Europe (mean: 18%; range: 14–22%). Moreover, our findings showed that the extent of contamination in public places was linked to climatic and geographical parameters, being higher in regions with a higher relative humidity and being located at higher longitudes and lower latitudes. Overall, this investigation (Fakhri et al., 2018) showed that public places are frequently heavily contaminated with *Toxocara* eggs. This finding calls for increased awareness of this risk factor and for measures to prevent the transmission of toxocarasis to humans in public places. There is also a need for well-designed, longitudinal and hypothesis-driven investigations using standardised experimental methods and robust statistical analyses.

4.2. Estimates of nation-wide seroprevalence rates in humans

Over the past 36 years, nation-wide surveys of humans have estimated prevalences of specific anti-*Toxocara* serum antibodies at 0.7% in New Zealand, 1.6% in Japan, 2.4% in Denmark, 6.3% in Austria, 7% in Sweden, 14% in the USA and 31% in Ireland, but prevalences are > 20% in some ethnic and socioeconomically disadvantaged groups, 22% in Iran and 81% in Nepal (Appendix Table 1). Notably, a seroprevalence of 85% has been recorded in school children in Manado, Indonesia (Hayashi et al., 2005), and 87% on the Marshall Islands (Fu et al., 2014). However, there are still limited detailed epidemiological data for most countries around the world.

Although human toxocarasis is expected to be common in underprivileged communities in which there is a close relationship between wild or domestic canids/felids and humans, it has been challenging to evaluate the global impact of this neglected disease (Hotez and Wilkins, 2009; Smith et al., 2009), because of some limitations of available diagnostic tools (see section 3; e.g., Fillaux et al., 2013; Holland, 2015; Moreira et al., 2014). Nonetheless, there is an exemplary, large national survey by Won et al. (2008), who utilised representative samples and a well-defined approach (e.g., sample dilution, immunoassay and statistical analysis) to obtain seroprevalence data. This investigation revealed an age-adjusted seroprevalence for toxocarasis of 13.9%; the prevalence in Mexican Americans (10.7%) and non-Hispanic whites (12%) was lower than in non-Hispanic blacks (21.2%). These numbers were updated in 2018, to find an overall lower prevalence of 5% but similarities, in terms of higher prevalence rates among selected at-risk groups, including non-Hispanic blacks and Hispanics, living in poverty existed (Liu et al., 2018). The authors suggested a link to ethnicity and emphasised that this seroprevalence difference might be an important health education message. Nevertheless, it is still challenging to serologically distinguish infection from exposure to *Toxocara* (Fisher, 2003; Gasser, 2013; Poulsen et al., 2015), which hinders a better understanding of the epidemiology (e.g., transmission dynamics) of human toxocarasis.

4.3. Seroprevalence estimates for human toxocarasis worldwide

There has been a need for investigations to estimate the prevalence of human toxocarasis worldwide to justify investments in prevention programs. In spite of the many epidemiological studies of toxocarasis and *Toxocara* spp., until recently, there had been no seroprevalence estimates for the general human population at the global, regional and national levels. For this reason, we performed a systematic review and meta-analysis of the literature to estimate the regional and global prevalences of anti-*Toxocara* serum antibodies in human populations around the world (Rostami et al., 2019); the authors used the term ‘T-seroprevalence’, so as to emphasise that the presence of anti-*Toxocara*

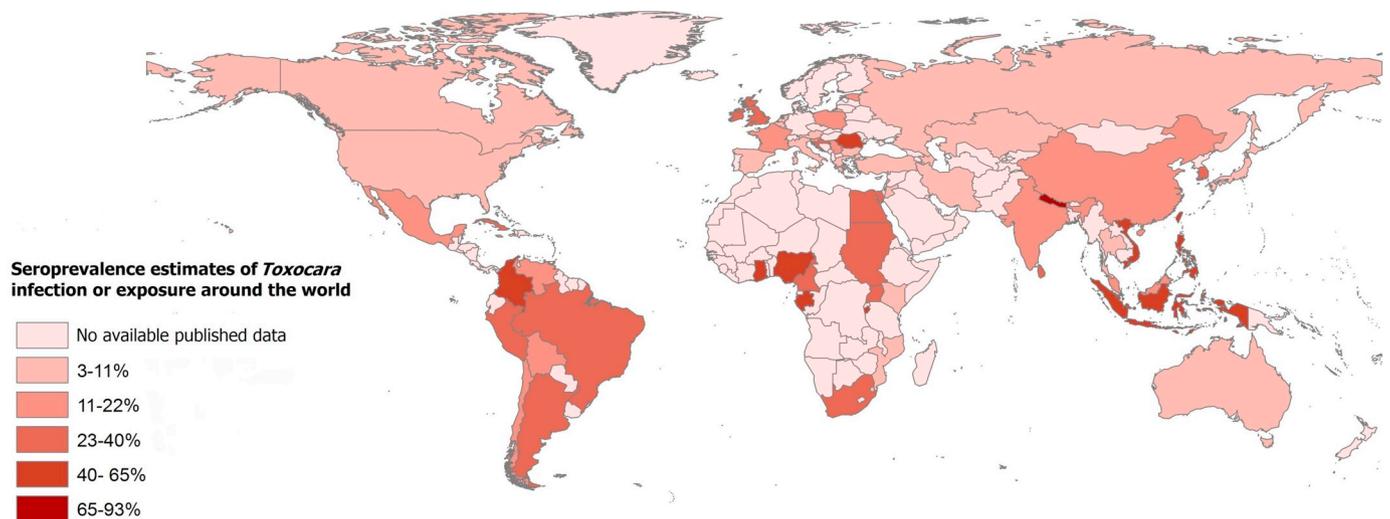


Fig. 2. Estimated seroprevalences of *Toxocara* infection or exposure around the world. Prevalence values (%) are estimated from published works (for details refer to Appendix Table 1).

serum antibodies can reflect infection or past exposure.

A systematic search was undertaken using data and information (1980 to 2019) from five public databases (Embase, PubMed, SciELO, Scopus and Web of Science). Random effect model-based meta-analysis was employed to calculate overall the prevalence of anti-*Toxocara* serum antibodies in people on a national, regional or global scale. Moreover, we evaluated the impact of socio-demographic, geographical and climatic parameters on seroprevalence, and assessed risk factors for seropositivity. By reviewing 250 studies of 265,327 people from 71 countries in WHO regions, we concluded that almost one fifth (19%; 1,411,749,590 individuals) of the world's human population is seropositive to *Toxocara*. The highest seroprevalence rates were found in Africa (mean: 37.7%; range: 25.7–50.6%) and the lowest in the Eastern Mediterranean region (mean: 8.2%; range: 5.1–12.0%). Relatively high mean seroprevalence rates were also estimated for the South-East Asia region (mean: 34.1%; range: 8.5–12.8%), the Western Pacific region (mean: 24.2%; range: 19.7–26.0%) and South American countries (mean: 27.8%; range: 23.1–32.7%), and lower rates for North American (mean: 12.8%; range: 10.0–15.8%) and European (mean: 10.5%; range: 8.5–12.8%) regions. A country or region with a low human development index (HDI), income level, latitude, and high environmental temperature and precipitation was more likely to have a high prevalence of anti-*Toxocara* serum antibodies in the human population. The findings also indicated that potential risk factors associated with *Toxocara* seropositivity were male gender; living in rural areas; young age; close contact with dogs, cats or soil; consumption of raw meat; and consumption of untreated drinking water.

The findings of a recent study (Rostami et al., 2019) indicated high levels of infection by or exposure to *Toxocara* spp. in many countries, which calls for increased attention to human toxocariasis and improved measures to prevent adverse health risks of this disease. The risk predictors identified in this investigation should be considered in the design of human toxocariasis prevention programs. Clearly, epidemiological and serological surveys are needed in countries for which prevalence data are lacking, in order to provide a basis for the implementation of future prevention and/or control programs.

4.4. Molecular epidemiology of *Toxocara* species of animals

Toxocara species have been identified and distinguished according to their morphological characteristics and also their particular host species. Recognised species include *T. canis* and *T. tanuki* (from canids), *T. cati* (felines), *T. vajrasthira* (mustelids), *T. lyncus* (caracals), *T. vitulorum* (bovids), *T. apodemi* and *T. mackerrasae* (rodents), *T. paradoxura*

and *T. sprengi* (viverrids), and *T. pteropodis* (bats) (cf. Bowman, 2014). Traditional taxonomic methods can have limitations for the identification and differentiation of some *Toxocara* species, especially at the larval and/or egg stages (Chen et al., 2012; Gasser, 2013; Gasser et al., 2016).

In the late 1990s, it was shown that PCR utilising sequences of ITS-1 and ITS-2 of rDNA were very useful for the differentiation of *Toxocara* species. Using ITS-2, Jacobs et al. (1997) successfully distinguished *T. canis*, *T. cati* and *Toxascaris leonina* from each other and from other ascaridoids that might be found in human tissues. Subsequently, PCR-based methods using ITS-1 and ITS-2 were employed to identify and differentiate of *Toxocara* species that were morphologically very similar or ambiguous (Gibbons et al., 2001; Zhu et al., 1998, 2001). For example, this approach was used to characterize a *Toxocara* species from cats from Malaysia, that, morphologically, was neither *T. canis* nor *T. cati*, and, using molecular methods was shown to represent a distinct species. This species was described and named *T. malaysiensis* (Gibbons et al., 2001; Zhu et al., 1998). In additional studies, *T. malaysiensis* was shown to occur in cats in China and Vietnam by molecular characterization using ITS-2 as the genetic marker (Le et al., 2016; Li et al., 2006). Our literature searches of international databases and also of the GenBank database revealed that *T. malaysiensis* has not yet been found in countries other than Malaysia, China and Vietnam. In addition to rDNA sequences, some studies have indicated that mt genes can be useful alternative markers for identification, and systematic and phylogenetic relationship analyses of *Toxocara* species (e.g., Hu et al., 2004; Hu and Gasser, 2006; Jex et al., 2008; Li et al., 2008; Wickramasinghe et al., 2009). Phylogenetic analyses using ITS-1, ITS-2 and mt gene data sets have shown that *T. malaysiensis* (from cats) was more closely related to *T. cati* (from cats) than to *T. canis* (from dogs) (Zhu et al., 1998; Le et al., 2016; Li et al., 2008). In addition, Wickramasinghe et al. (2009), using mt DNA sequences, reported that *T. vitulorum* was genetically more similar to *T. malaysiensis* than to *T. canis* and *T. cati*, and Sultan et al. (2015) indicated that *T. canis* and *T. cati* were more closely related to one another than to *T. vitulorum* following a phylogenetic analysis of ITS-1 sequence data. Moreover, in the latter study, *T. tanuki* from raccoon dogs did not cluster with any other species of *Toxocara* included. We searched international databases for molecular investigations of *Toxocara* spp. in different hosts and in soil and found only a small number of studies (Table 2), indicating a major need for comprehensive molecular epidemiological studies of *Toxocara* species in different host species and in public areas in different geographical regions.

Table 2
List of studies reporting the molecular identification of *Toxocara* species in different host species and in soil.

Country	Origin of sample	Number of samples tested by molecular means	<i>Toxocara</i> spp.			References
			<i>T. canis</i>	<i>T. cati</i>	<i>T. malaysiensis</i>	
Switzerland	Dog	35	24	11	–	Fahrion et al. (2011)
Iran	Dog	28	9	–	–	Mikaeili et al. (2015)
Vietnam	Dog	14	14	–	–	Le et al. (2016)
Turkey	Cat and Dog	20	17	3	–	Oguz et al. (2018)
Pakistan	Dog	26 ^a	–	–	–	Shah et al. (2018)
India	Dog	30 ^a	25	–	–	Suganya et al. (2019)
Switzerland	Cat	36	–	36	–	Fahrion et al. (2011)
Iran	Cat	6	–	6	–	Mikaeili et al. (2013)
Iran	Cat	32	–	32	–	Mikaeili et al. (2015)
Vietnam	Cat	15	–	–	15	Le et al. (2016)
Iran	Chicken	6	5	1	–	Zibaei et al. (2017)
Japan	Soil	102 ^a	–	66	–	Macuhova et al. (2013)
Iran	Soil	71	12	59	–	Khademvatan et al. (2014)
Iran	Soil	24	1	23	–	Chooibneh et al. (2019)

^a Samples that amplified by PCR.

5. Need for public awareness and intervention programs

The distribution and seroprevalence estimates for human toxocarosis worldwide raises considerable public health concerns. We believe that community awareness about toxocarosis is still inadequate, and that improved education will be critical to enhancing a community's understanding of this disease, and its prevention and control (Fig. 3; Holland, 2015; Ma et al., 2018). Some www-based educational resources, such as the Centers for Disease Control and Prevention (CDC, <http://www.cdc.gov>) and the American Association of Veterinary Parasitologists (AAVP, <http://www.aavp.org/>) contain comprehensive information on *Toxocara*/toxocarosis. Veterinary and medical practitioners need to be responsible for educating pet owners about the medical importance of toxocarosis and about how to minimise the risks of zoonotic transmission (Smith et al., 2009). Importantly, veterinary practitioners are commonly exposed to and/or infected by *Toxocara* due to their occupational contact with small animals (Deutz et al., 2005; Poepl et al., 2013). In contrast to most veterinary courses in parasitology, especially helminths are rarely, if ever, covered in many curricula provided to medical students; therefore, knowledge about toxocarosis is usually quite limited in the medical profession. Since the need for a differential diagnosis of *T. canis* infection may not be considered in a clinical context, patients who present with pulmonary dysfunction (CT), asthma-like symptoms and/or cognitive deficits (NT) are often not tested for the possibility of a parasitic infection. Hygiene, preventing children from ingesting infective *Toxocara* egg-

contaminated soil or faecal remnants from carnivores and avoiding the ingestion of uncooked meat or liver are central to preventing toxocarosis in endemic countries (Fan et al., 2013; Moreira et al., 2014). In addition, although asymptomatic infections are often self-limiting, human toxocarosis cases must be treated to prevent NT and OLM (Othman, 2012).

Furthermore, the relationship among animals, humans and the environment must be kept in mind. Since numerous transmission routes are available to *Toxocara* species, a *One Health* approach is central to the prevention and control of human toxocarosis (Fig. 3; Holland, 2015; Ma et al., 2018). Reducing *Toxocara* infection in definitive host populations needs to be a priority to significantly decrease the number of infective *Toxocara* eggs in public places (Palmer et al., 2008). In this context, measures to prevent dog fouling have been recommended (Atenstaedt and Jones, 2011).

Nijse et al. (2015) indicated that the removal of faeces and inclusion of anthelmintic treatment strategies (four times per year) would reduce egg output and resultant environmental contamination. Particularly puppies of < 12 weeks of age and kittens (Morgan et al., 2013; Nijse et al., 2014) as well as working/stray dogs and cats should be dewormed with an effective anthelmintic drug (El-Tras et al., 2011; Roddie et al., 2008). Such treatment of pregnant animals is not effective at preventing transplacental transmission (ante-partum) or subsequent transmammary transmission (post-partum), and thus requires further research efforts (cf. Overgaauw and van Knapen, 2013). Controlling transmission is more challenging when wild animals (e.g., foxes) are involved (Holland, 2015; Morgan et al., 2013).

In endemic settings (such as underprivileged, rural communities in tropical towns or cities with poor sanitation and large stray dog or cat populations), intervention needs to be based on a sound understanding of the epidemiology of toxocarosis, human populations at most risk of infection (e.g., age-groups, socioeconomic status, and rural versus urban), and the cost and feasibility of control and surveillance.

6. Progress through 'omics and informatics– toward new diagnostic and therapeutic approaches

The *T. canis* genome project (Zhu et al., 2015) represented a major step toward an improved understanding of the molecular biology of the parasite as well as host-parasite relationships, and might support work to develop novel intervention and diagnostic methods (Gasser et al., 2016). The draft genome is 317 Mb in size and was predicted to encode > 18,500 genes; 78.4% of these genes could be annotated, and 5406 (29.1%) were inferred to have homologues in known biological (KEGG) pathways (Gasser et al., 2016; Zhu et al., 2015). Specifically, 870 ES proteins (including proteases, cell adhesion molecules, lectins,

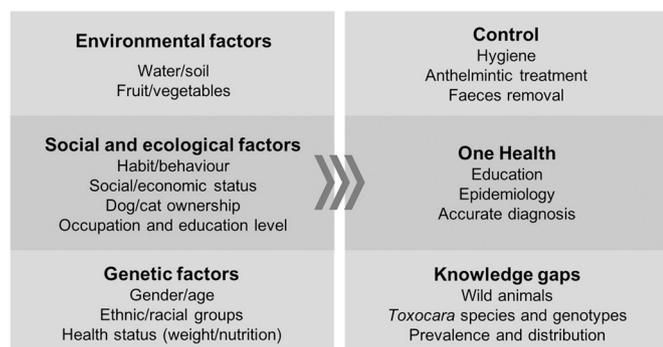


Fig. 3. Epidemiological risk factors and determinants of human toxocarosis. Human toxocarosis can be influenced by numerous factors including contamination of the environment by *Toxocara* eggs, social/ecological aspects (animal ownership, education and/or occupation) and genetic factors (e.g., age, gender and/or health status). A *One Health* strategy should improve our knowledge base and enhance the prevention and control of toxocarosis.

SCP/TAPS proteins, and mucins) were predicted to be involved in host invasion and in parasite-host 'cross-talk', such as immune evasion and/or immune modulation; 458 kinases, 408 phosphatases and 127 GTPases were predicted to have important roles in embryonic, larval development and reproduction; and some of the 156 GPCRs, 268 ion channel proteins and 530 transporters might be drug targets. Additionally, intestine-, stage- and gender-enriched molecules/pathways were defined, indicating essential involvement in survival, development and/or reproductive processes (Gasser et al., 2016; Zhu et al., 2015).

Utilising these genomic and transcriptomic resources (Gasser et al., 2016; Zhu et al., 2015), we explored the transcription profiles of small RNAs in adult *T. canis* (see Ma et al., 2016). The results suggested that miRNAs *Tc*-miR-6090 and *Tc*-miR-2305 are involved in embryonic and larval development as well as reproduction in *T. canis*, and that *Tc*-miR-100, *Tc*-miR-34 and *Tc*-let-7-5p appear to relate to host-parasite interactions. Other miRNAs, including *Tc*-miR-5126, *Tc*-miR-2861 and *Tc*-miR-2881, might be anthelmintic targets and/or link to drug resistance. Such molecular investigations are now starting to enhance our understanding of molecular processes in *T. canis*, and should provide a foundation for investigations of the parasite's developmental biology, host-parasite cross-talk and disease processes; such studies might also help discover new intervention and diagnostic approaches (cf. Britton et al., 2015; Hoy et al., 2014; Manzano-Román and Siles-Lucas, 2012).

7. Prospects for immunoprotection

There is an increasing interest in vaccines against zoonotic parasitic diseases. However, for many such diseases, protective immunity is often 'blocked' by a parasite's ability to inhibit and/or evade the host immune system (McSorley and Maizels, 2012). For *Toxocara* species, immune evasion is mediated by surface-coat and excretory/secretory (ES) molecules (Maizels, 2013), which might be vaccine targets for protective type 2-like immune responses (Długosz et al., 2015; Loukas and Maizels, 2000). TES components, originally explored by Maizels et al. (1984), include the C-type lectins TES-32 and -70 and glycosylated mucins which represent a surface-coat of the larval stage (Długosz et al., 2015; Gems and Maizels, 1996; Kennedy et al., 1987; Loukas et al., 1999, 2000; Loukas and Maizels, 2000; Maizels et al., 2000). Recently, based on analyses of the genome and transcriptome of *T. canis* (see Zhu et al., 2015), it was suggested that a vast array of proteases, adhesion molecules, SCP/TAPS proteins and lectins play roles in the host-parasite 'cross-talk' (cf. Gasser et al., 2016).

The host immune response to *Toxocara* includes a dominant CD4⁺ T-helper type 2 cell (Th2) activity, eosinophilia and production of specific antibodies (Del Prete et al., 1991; Maizels, 2013). TES antigens drive the production of type 2 cytokines (e.g., interleukin-4, IL-5, -10, and -13) from peripheral T-cells in exposed individuals, resulting in eosinophilia, enhanced cytokine expression and the production of IgE antibodies (Del Prete et al., 1991; Maizels, 2013; Mazur-Melewska et al., 2016). These factors might contribute to airway hypersensitivity, linking chronic *Toxocara* infection to allergic diseases, such as asthma and allergic rhinitis (Cooper, 2008; Maizels, 2013; Yariktas et al., 2007). Interestingly, respiratory signs, such as wheezing, in people (particularly children) can be associated with VLM and CT due to pulmonary migration of *Toxocara* larvae (e.g., Mendonça et al., 2012; Taylor et al., 1988). Whether children develop asthma later in life is unclear, but there is a possibility that they might be protected from atopic asthma as a result of an immunoregulatory response induced by infection (Cooper, 2008; Yariktas et al., 2007). Although many parasitic helminths can induce immunoregulatory cell populations (e.g., regulatory T cells and alternative-activated macrophages) (Hewitson et al., 2009), it is possible that *T. canis* has evolved to be more immunogenic in accidental hosts, including humans. Nevertheless, some anti-inflammatory effects can be mediated by TES, resulting in, for example, an inhibition of Toll-like receptor signalling and nitric oxide production

(Hewitson et al., 2009).

Immunomodulation is a characteristic of *Toxocara* infections. For example, increased protective pro-inflammatory cytokines (IFN-gamma, IL-6 and IL-13) and an anti-inflammatory cytokine (IL-10) can be detected in sera from *Toxocara*-infected children (Nagy et al., 2012), and macrophages from infected mice produce more IL-10/TGF- β and less IL-12/TNF compared with uninfected-control mice (Kuroda et al., 2001). There are various mechanisms contributing to the activation and suppression of immune responses. Firstly, the surface-coat shed by the worm enables migrating larvae to escape from eosinophils that 'stick' to their surface, thus neutralising antibody-dependent elimination (Fattah et al., 1986). Secondly, TES antigens (Th2-stimulator) can also modulate systemic and local immune responses (Maizels, 2013). For example, C-type lectins, particularly TES-32 and TES-70, have marked homology to low affinity IgE and macrophage mannose receptors in mammals (Loukas et al., 1999), capable of targeting host pathways involved in innate immunity (Hewitson et al., 2009). Additionally, extracellular vesicles (exosomes) might also represent other means of *Toxocara* being able to modulate immune responses, as such vesicles have been shown to transfer ES antigens and non-coding RNAs to host cells (Buck et al., 2014; Quintana et al., 2015). However, this new area should be explored in detail. An improved understanding of specific immune responses and modulatory mechanisms during toxocarosis, and defining immunogenic molecules, could assist significantly in developing an anti-*Toxocara* vaccine.

8. Conclusions and future focus

Toxocarosis is a neglected disease which affects millions of people, pets and stray companion animals around the world. Comprehensive meta-analyses of published seroprevalence and environmental studies suggest that exposure to *Toxocara* is common, particularly in children who live in subtropical/tropical and underprivileged regions of the world. Improved techniques (immunological, genetic and/or imaging) are needed for the diagnosis of the main clinical forms of toxocarosis (VLM, OLM, CT and/or NT), and need to be accessible to the estimated 750 million people who are at the highest risk of infection and who survive below the World Bank poverty level. There is a clear need to better understand the pathogenesis and impact of CT and NT in people, particularly in disadvantaged community settings; these two clinical conditions might be linked to major causes of pulmonary dysfunction and cognitive delays, respectively, but there are major knowledge gaps, and there are no formal research projects on this topic.

It is possible that the geographic distribution of *T. canis*, and possibly other species, is expanding as a consequence of increased human and animal mobility, together with the effects of climatic alterations, such as global warming. The human population has doubled in the last 50 years, and, today, more than half of the world's human population live in urban areas. The population of dogs and cats has also grown substantially (Gompper, 2013), increasing the density of these definitive hosts. Much of this growth and concentration has occurred in cities in the tropics and subtropics, and means that without specific prevention and control programs, the prevalence and socioeconomic impact of toxocarosis is likely to increase.

There is a clear need for a better understanding of the molecular epidemiology and ecology of *Toxocara*/toxocarosis (Gasser, 2013; Ma et al., 2018). Future work might focus on the use of improved molecular tools for specific parasite identification and genetic analysis and for the accurate diagnosis of current infection in people. It would be particularly important to explore, in detail, the zoonotic importance of *T. cati* and whether *T. malaysiensis* and/or whether other congeners other than *T. canis* and *T. cati* also play a significant role in human toxocarosis. Highly sensitive and specific molecular technologies could help establish what role these other ascaridoids play in the epidemiology of toxocarosis, to reveal the true extent and, thus, the public health significance of *Toxocara*, and would likely underpin new and improved

management strategies for the prevention, treatment and control of toxocarosis. Developing an effective ovicidal anthelmintic against *Toxocara* eggs and/or a vaccine to protect young canids and felids, or people, against toxocarosis, which have long been goals of the scientific community, would enhance the likelihood of controlling this neglected zoonotic disease. It is also hoped that progress in the areas of genomics, transcriptomics, proteomics and bioinformatics as well as recently established molecular resources will enable the discovery of new intervention targets for *T. canis* and its congeners.

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Appendix A. Supplementary data

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