Prevalence and Risk Factors of Toxoplasma gondii Infection in Domestic Cats from the Tropics of Mexico Using Serological and Molecular Tests

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1. Introduction

Toxoplasmosis is a major parasitic zoonoses of worldwide distribution caused by the intracellular protozoa Toxoplasma gondii. Felines, in particular domestic cats, have an important role in the epidemiology of the disease because the sexual (and asexual) reproduction takes place in them and they excrete a large number of infective and environmentally resistant oocysts [1, 2]. This parasite infects most mammals (including humans) and birds which are intermediary hosts. Among vertebrates, food animals are particularly important because they are intended for human consumption [3, 4]. Cats primarily become infected when they ingest the encysted tissue cyst from the intermediate hosts like rodents and birds. Human acquire infection when they ingest sporulated oocysts from the soil, contaminated water sources, tissue cysts from raw or undercooked meat, or transplacentally [2, 5]. Infection in cats frequently takes place as subclinical or asymptomatic disease and is very rare to develop evident clinical signs [6]. Toxoplasmosis in cats is reported worldwide with seroprevalences ranging from 60 to 90%. It is probably due to different diagnostic techniques used in each of the reported studies, cultural factors and environment of every region promoting the persistence of infective oocysts [2, 7]. In Mexico, the reported prevalence of cat toxoplasmosis varies from 9.3% to 70.8% [8–11] depending on the region. Associated risk factors for infection in cats included feeding with raw meat [8, 10], age and sex.
and access to hunting [12]. In Mexico the seroprevalence of human toxoplasmosis is estimated to be 25% [13]. In Yucatan, Mexico, no epidemiological information about toxoplasmosis in cats is available despite the presence of the infection in humans, with a seroprevalence of 57.5% [14] and the tragic consequences associated with toxoplasmosis in AIDS patients [15].

There are several diagnostic methods for determining infection of T. gondii. The most widely used serological test for diagnosis of toxoplasmosis is the indirect ELISA (IgM and IgG). Moreover, the polymerase chain reaction (PCR) is a molecular test which allows detection of parasite DNA; it is highly specific and sensitive and very useful together with serological tests to differentiate the chronic, acute or reactivated infections. PCR is also an important diagnostic method when the immune system of the patient is compromised or when antibody titers have no reach threshold levels of detection [16, 17]. In domestic cats, several diseases may affect the overall health and cause immunosuppression such as infection with Bartonella spp. feline immunodeficiency virus (FIV) and feline leukemia virus (FeLV) [18, 19] that may increase the effects of acute toxoplasmosis [20, 21].

The information on the prevalence of toxoplasmosis in cats is useful for evaluating environmental contamination with the protozoa and to evaluate the risk it poses to public health. The aim of this study was to estimate the prevalence of toxoplasmosis in domestic cats in Merida, Yucatan, Mexico, using serological and molecular methods, and to determine possible risk factors (sex, age, diet, hunting, body condition, and number of cats per household).

2. Material and Methods

2.1. Study Area and Population. A cross-sectional study was performed in the city of Merida, Yucatan, (19° 30’ and 21° 35’ north latitude, 87° 30’ and 90° 24’ west longitude). The city’s climate is characterized by warm subhumid with summer rainfall and average temperature of 26°C. The relative humidity is maximum 83% and minimum 61% with a height of 6 meters above sea level [22]. The population of interest were cats (Felis silvestris catus) living in Merida. The inclusion criteria were domiciled cats, of any race or sex, and older than 3 months of age.

2.2. Sample Size. A pilot study in 40 cats measuring IgG antibodies specific for T. gondii using an indirect ELISA test showed a seroprevalence of 82.5% which was used as the expected prevalence in the calculation of sample size. It was considered an infinite population of cats, a confidence level of 95% and an absolute accuracy of 5%, calculated a sample size of 220 cats [23].

2.3. Collection of Samples. A total of 231 blood samples (1–3 mL) were taken by puncturing the jugular vein and deposited in two vacutainer tubes, with and without anticoagulant, the first for the PCR test and the second for serological testing. DNA extraction was performed with the commercial kit DNeasy Blood and Tissue (QIAGEN, cat no. 69506). Before extraction, a prelysis of blood was conducted as suggested by Jalal et al. [24]. The mean volume of DNA extraction from each cat was 100 μL. Samples were stored at −20°C until further PCR assay. From the collected samples, 11 were not viable for diagnostic testing (negative samples by PCR for beta-globin constitutive gene) and were discarded leaving a total of 220 samples.

2.4. Risk Factors. Through a questionnaire survey to each owner, information was obtained to assess the following risk factors.

Age. Age was determined based on the review of the animal or based on their dentition [25, 26]. The animals were considered in two groups: ≤1 year and >1 until 7 years.

Sex. Male or female.

Type of diet. Two categories were considered: cats fed only with commercial food and cats fed with commercial food and/or raw meat.

Access to the hunting. If the cat had any access to hunting.

Number of cats per household. Categorized as one or more than one cat at home.

Body condition. It was divided into two groups: cats with good body condition and body condition regular to bad [27].

2.5. Indirect ELISA Test. The presence of specific IgM and IgG antibodies against T. gondii was determined separately by the use of indirect ELISA tests (Human-GmbH, Wiesbaden, GER). The technique used was adapted to that described by Figueiroa et al. [28], but using anti-IgM and IgG cat antibodies labeled with horseradish peroxidase (HRP) (Santa Cruz Inc., CA, USA) on 96-well plate coated with sonicated parasite proteins from tachyzoites of T. gondii. Serum samples were diluted to a ratio of 1:100 in phosphate-buffered saline (PBS; pH 7.2). The secondary goat anti-IgG and anti-IgM cat antibody HRP labeled were used at a dilution of 1:5,000. Sera from cats showing high anti-IgG antibodies titer by ELISA (1:1024) and positive results to PCR against T. gondii were used as positive controls, and sera pool from 10 healthy cats previously tested by triplicate with ELISA IgM, IgG and PCR, were used as negative controls. On the basis of the indirect ELISA results, subjects were diagnosed as either positive/negative for specific IgG and IgM antibodies to T. gondii. The optical density (OD) was measured in a spectrophotometer at 450 nm (Multiskan Multisoft Primary EIA) and was used to compute the percent positivity (PP) using the formula mean OD (sample or negative control) divided by the mean OD value positive control multiplied by 100. Percent positivity of 15% or above was considered as positive.

2.6. Polymerase Chain Reaction. The PCR was performed as described by [24]. Primers Tg1 (5’-AAAAATGTGGGAATGAAAGAGACGG-3’) and Tg2 (5’-AGCAATCAACGGAGACGGTGT-AAT-3’) that amplify a fragment of 469 base pairs (bp) from the B1 gene of T. gondii were used. Each PCR reaction was carried out at a final concentration of 0.4 mM dNTPs, 2.5 μmol of each primer, 1.5 mM MgCl2, and 1U of GoTaq
used and the reaction was completed in a final volume of 25 μL. Amplification conditions were 95 °C for 10 minutes for denaturalization, followed by 35 cycles of 94 °C for 1 minute, 52 °C for 30 seconds and 72 °C for 1 minute, and a final extension cycle of 72 °C for 7 minutes. The PCR products were analyzed by electrophoresis on agarose gels and stained with 1.8% ethidium bromide. As a positive PCR control a plasmid clone pMOSBlue/Toxo 469/3, containing a sequence of 469 pb, amplification product of Tg1 and Tg2 primers of parasite genome was used.

2.7. Statistical Analysis. Prevalence for each serological or molecular result and its combinations was estimated. Because of the small number of cases, Chi-square test and binomial logistic regression were carried out only for IgM, IgG and PCR positive cases. Significant risk factors, under the Chi-square tests, were further investigated using binomial logistic regression. Values were considered significant when \( P < 0.05 \). Odds ratio (OR) and confidence interval (CI) at 95% were also estimated. Statistical analyses were carried out using the SPSS package v 17.0 (SPSS Inc., Chicago, Il, USA).

3. Results

3.1. Prevalence. Serological prevalence of \( T. gondii \) was 91.8% and 75.5% for IgG and IgM respectively. One hundred seventy-three samples of DNA (79%) tested amplified a 469 bp fragment corresponding to the expected size of the B1 gene sequence specific for \( T. gondii \). An example of amplification is shown in Figure 1. The result of the different serological, molecular, and combination of positive and negative cases is shown in Table 1. Cases were categorized as acute when were only PCR positive or when the patient showed positive titers of IgM + PCR. Chronic cases were considered when patients were only IgG positive, IgG + PCR positive, IgM + IgG positive, or IgM + IgG + PCR positive. Positive IgM + IgG + PCR cases (61.4%), positive IgM + IgG (11.8%) and positive IgG + PCR (13.2%) were the most frequent cases.

3.2. Risk Factors. The binomial logistic regression for cats with positive IgM + IgG + PCR cases showed association with body condition (fair—poor) and the presence of more than one cat at home (Table 2). However, there was no association of the variables sex and type of diet with cats infected with \( T. gondii \). Also preliminary Chi-square tests showed not association of the factor cat having access to hunting and toxoplasmosis infection.

4. Discussion

The results of this study indicate a high rate of infection in cats with \( T. gondii \), which suggests a high environmental contamination with infective oocysts. There is a wide variation in reported prevalence of feline toxoplasmosis, which is probably due to environmental, cultural, and diversity of diagnosis techniques and type of molecule measured in each study (IgA, IgM, IgG, genome, etc.). In tropical countries like Colombia, Thailand, and Brazil, reported IgG seroprevalences of toxoplasmosis in cats are 45.2% [29], 11.0% [30], and 40% [31], respectively. In some regions of Mexico the reported seroprevalence in cats is 28.8% in Colima [9], 21.9% in Mexico City [10] and 70.8% in Guadalajara [8]. This study has recorded the highest seroprevalence (91.8%) of IgG antibodies towards \( T. gondii \) in cats, even higher than that found in Tehran, Iran, where an overall prevalence of 63% \( (n = 100 \) cats) reported and from which feral cats had a 90% infection [32]. In Parana, Brazil, it was reported that 84.4% of cats were seropositive for IgG [33], which is close to the results obtained in this study. The rainfall and subtropical climate conditions with hot summers of Parana is similar to the tropical climate of the city of Merida, which is favorable to maintain viable oocysts in the environment for long periods of time.

In this study the prevalence of cats seropositive for IgM was 75.5%. In a study by Kodym et al. [34] it was reported that IgM antibodies against \( T. gondii \) may remain in circulation for 12–18 months, so that the titers of IgM cannot by themselves be indicators of recent infection, which is contrary to what is believed. In our study, the IgM and IgG serological status was complemented with molecular studies for a better determination of the infection status.

The percentage of acute cases of toxoplasmosis found in this study (5.9%) considering only PCR positive cases (1.8%) plus IgM, PCR positive cases (4.1%) are lower than the 8.3% found by Galván et al. [8] in Guadalajara, Mexico, considering only IgM antibodies. Acute cases are not commonly found in survey studies [35] and their presence indicates a constant dynamic of the disease and the high risk of contact of cats with the protozoan. This prevalence of cases defined as acute may depend on the chance of cats to be in contact with infecting cyts through the hunting infected prey or food provided to them by their owners, hence the importance consider adequate food for cats. Early-acute cases are those that were positive by PCR before the production of any immune response. In this study 79% of cats were positive to PCR, but only 1.81% were in the initial stage of infection where there is not production of antibodies [36] or those cats that may have compromised immune system. Suh and Joo [37] reported 5.3% of cats PCR positive to \( T. gondii \). However, Lee et al. [38] more recently reported
Table 1: Type and frequency of *T. gondii* infection in cats according to serological and molecular results.

<table>
<thead>
<tr>
<th>Type</th>
<th>IgM</th>
<th>IgG</th>
<th>PCR</th>
<th>n</th>
<th>%</th>
</tr>
</thead>
<tbody>
<tr>
<td>Acute</td>
<td>−</td>
<td>−</td>
<td>+</td>
<td>4</td>
<td>1.8</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>−</td>
<td>+</td>
<td>9</td>
<td>4.1</td>
</tr>
<tr>
<td></td>
<td>−</td>
<td>+</td>
<td>−</td>
<td>12</td>
<td>5.5</td>
</tr>
<tr>
<td></td>
<td>−</td>
<td>+</td>
<td>+</td>
<td>29</td>
<td>13.2</td>
</tr>
<tr>
<td>Chronic</td>
<td>+</td>
<td>+</td>
<td>−</td>
<td>26</td>
<td>11.8</td>
</tr>
<tr>
<td></td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>135</td>
<td>61.4</td>
</tr>
<tr>
<td>Negative</td>
<td>−</td>
<td>−</td>
<td>−</td>
<td>5</td>
<td>2.2</td>
</tr>
<tr>
<td>Total</td>
<td>220</td>
<td></td>
<td></td>
<td>100</td>
<td></td>
</tr>
</tbody>
</table>

Table 2: Binomial logistic regression to evaluate the association between studied variables and *T. gondii* infection cases from chronically reactivated cats. Chronically reactivated cases (positive to IgM, IgG, and PCR).

<table>
<thead>
<tr>
<th></th>
<th>β</th>
<th>SE</th>
<th>OR</th>
<th>CI</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age (&gt;1–7 Years)</td>
<td>0.489</td>
<td>0.294</td>
<td>1.63</td>
<td>0.916–2.903</td>
<td>0.097</td>
</tr>
<tr>
<td>Cats per household (&gt;1)</td>
<td>−0.714</td>
<td>0.292</td>
<td>0.49</td>
<td>0.276–0.869</td>
<td>0.015</td>
</tr>
<tr>
<td>Body condition (Regular-bad)</td>
<td>0.928</td>
<td>0.338</td>
<td>2.53</td>
<td>1.306–4.904</td>
<td>0.006</td>
</tr>
</tbody>
</table>


47.2% of positive cats by using nested PCR both in South Korean conditions.

A high number of chronic cases should be expected while studying cat populations. Traditionally, cats seropositive for IgG are considered as chronically infected [6]. In this study 91.9% of cats were in the chronic stage of infection but with different serological and/or molecular status. When considering a constant exposure to the protozoa the immune response and circulation of the genome in the host may differ giving as result the variety of diagnostic situations here reported.

Cases when cats were IgG positives but with high levels of IgM are an indicator of revival [21]. But they also may indicate a switching from IgM to IgG weeks after infection. Reactivation in chronically infected stages involves reactivation of cysts and conversion of bradyzoites to tachyzoites and not necessarily involves a new enteropetitelial phase and excretion of oocysts [6]. However, IgM not necessarily indicates a recent infection but the outcome of the combination of different antibodies and molecular studies may be useful to confirm the chronically reactivated stage; the recirculation of the parasite in blood and serological tests indicate that there was a previous immune response.

Most cats in the chronic stage found in this study were IgM + IgG + PCR positive and are proposed to be named as chronic reactivated cases; this means that not only the parasite is circulating in the environment producing constant contact with the cats, but there are periods of reactivation with or without excretion of oocysts depending on its immune system. In chronically infected hosts with tissue cysts, these may be rupture and release bradyzoites to the circulation [39] which are eventually destroyed by the immunocompetent host or reactivate the infection in the immunosuppressed animal [40].

It is expected that cats having access to the streets be more likely to hunt and become infected with *T. gondii*. However, in the present study no association was found between cats gaining access to the streets and hunting than those kept indoors. The nocturnal gecko *Hemidactylus frenatus* is commonly found inside all households in the area of the study which are predated by cats and ten infected; these in part may explain the results founded.

In chronic reactivated cases, a significant protective association was found with more than one cat per household. This result is difficult to explain; however, it indicates that when living with other cats, they are less likely to be infected and reinfected with *T. gondii*, similarly as reported in cats from México city which indicates a change in environmental conditions and different immune status of cats living together [10].

Body condition has been used as an indicator of poor nutrition in small animals and consequently a weakened immune status, increased susceptibility to disease and longer periods of parasitemia including those caused by protozoa [40, 41]. In the present study regular-to-poor body condition was significantly associated with chronic reactivated cases, indicating some degree of release and circulation of *T. gondii* oocysts. Cats with immunosuppressive diseases, poor body condition are those suffering with periods of *T. gondii* reactivation [42, 43].

Finally, it is expected that with age, the risk of contact with the agent and infection occurs, but results from the present study indicates that cats become infected very early during their life.

It is concluded that *T. gondii* is widely widespread in the domestic owned cat population from Mérida city with various periods of reinfection and therefore there is a high environmental contamination with infective oocysts. Cats over one year old, with a poor body condition, increase the risk of becoming infected with *T. gondii*. 
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