A study of the prevalence and genotypes of *Giardia duodenalis* infecting kennelled dogs

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Abstract

*Giardia duodenalis* is a protozoan parasite of animals that is zoonotic. Given the capacity of this organism to spread via the faecal–oral route, animals held in overcrowded and unhygienic conditions are at high risk of infection. Faecal samples from dogs in three kennels in Rome were examined by microscopy and PCR for *G. duodenalis*, and the prevalence data generated were correlated with variables such as kennel identity, age of dog, length of time the dog had been kennelled and clinical signs.

The overall prevalence of the parasite in the faecal samples was 20.5% and was higher in samples from the largest kennel, which had the greatest turnover of dogs, and in faecal samples from younger animals. *Giardia* cysts were found more frequently in diarrhoeic animals but were also found in dogs with no clinical signs. Although the finding that the majority of isolates were dog-specific rather than zoonotic genotypes suggests that the zoonotic risk from this pathogen is less than previously thought, the higher prevalence of infection in younger dogs may pose a specific public health issue as such animals are more frequently re-homed with families.

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Introduction

In 1991, legislation introduced in Italy (L 281/1991) gave local authorities responsibility for the control of stray dogs. Captured strays are now kennelled under the supervision of veterinarians and are vaccinated and given anthelmintic treatments to prevent the spread of infectious and parasitic diseases. Given the capacity of the flagellate protozoan parasite *Giardia duodenalis* to spread via the faecal–oral route, such kennelled dogs can be at increased risk of this infection. Furthermore, some genotypes of *G. duode- nalis* are zoonotic (Hunter and Thompson, 2005).

The prevalence of canine giardiosis worldwide varies from 5% in well cared-for domestic pets to up to 100% in kennelled animals (Barr and Bowman, 1994; Bugg et al., 1999; Barutzky and Schaper, 2003; Dubnà et al., 2007). In Italy, sheltered animals have levels of *Giardia* infection of between 14% and 74% compared to levels of 4–19% in domestic pets (Giangaspero et al., 2002; Capelli et al., 2003; Berrilli et al., 2004; Bianciardi et al., 2004; Papini et al., 2005; Capelli et al., 2006; Giangaspero et al., 2007). Many factors affect the prevalence of the parasite including the age of the dog, living conditions, animal density, nutritional and immune status, and the methods used to diagnose infection.
Molecular studies have revealed that *G. duodenalis* has at least seven major genotypes or ‘assemblages’ designated A–G (Monis et al., 2003). Assemblages A and B have been found in a broad range of hosts including dogs and humans whereas genotypes C to G have distinct host associations and are not considered zoonotic (Monis et al., 1996; Monis et al., 1999; Hunter and Thompson, 2005). In addition to assemblages A and B, genotypes C and D are also commonly found in the dog (Monis et al., 1998).

The objective of this study was to determine the prevalence and genotypes of *G. duodenalis* infecting kennelled dogs in the Rome area using both microscopy and PCR and to evaluate the associated risk factors. Such information would be useful in the context of public health and might also lead to the development of novel ways of managing such dogs so as to reduce the incidence of this infection.

**Materials and methods**

**Study procedure**

A cross-sectional study was carried out on a population of 1400 dogs from three kennels, identified as K1, K2 and K3, in Rome. The number of animals to be tested was calculated as 124, on the basis of an expected prevalence of 15% (Berrilli et al., 2004), with a maximum sampling error of 6% and a confidence interval (CI) of 95%.

Animals were randomly faecal sampled on one occasion by a veterinarian between April 2005 and March 2006 and the samples were examined for the presence of *G. duodenalis* both by microscopy and by PCR and data relating to potentially significant variables were collected via questionnaire at the time of sampling. The questionnaire requested data in relation to kennel identity, sex of dog, age of dog (<1 year old or >1 year old), length of time the dog had been kennelled (0–3 months, 3–12 months or >12 months), access to common shared space, whether food was wet or dry, previous treatments (product used and date), and any clinical signs including details of faecal consistency.

**Kennel details**

The first kennel (K1) was newly built and housed an average of 900 dogs. Groups of up to 10 dogs had access to common, clay-based paddocks and the dogs were paired in boxes with washable, smooth cement floors. The second kennel (K2) housed 250 dogs on average and the dogs were kept in boxes in pairs or threes and had daily access to a common paddock. Box and paddock surface were similar to those in K1. The third kennel (K3) typically housed 230 dogs but had ceased to take animals in 2003 so that animal numbers had been steadily decreasing. In this kennel individual boxes held between 2 and 7 dogs. Standards of hygiene and animal management in the three kennels were similar and at each kennel, dogs were given anthelmintic treatment on arrival and were periodically re-treated.

**Microscopic examination of faeces for Giardia duodenalis**

Faecal samples were examined for *Giardia* cysts and/or trophozoites using the wet mount Lugol’s iodine staining method and the formol ethyl-acetate concentration technique (Ritchie, 1948; Young et al., 1979).

**Genotyping of Giardia duodenalis**

Genomic DNA was extracted directly from fresh or frozen faeces samples using QIAamp DNA Stool Mini Kit (Qiagen) and stored at 4 °C until molecular analysis was performed. All DNA extracts were subjected to a nested PCR to amplify a 130 bp region of the 18S-rRNA gene (Read et al., 2002). Sterile distilled water was included as negative control in each batch of DNA extraction and PCR reactions. Amplification products were visualised by 1% agarose gel electrophoresis stained with a solution of 6 μL/mL SYBR Safe DNA gel stain (Invitrogen) and visualised under ultra-violet light. PCR products were purified using a NucleoSpin Extract (Macherey–Nagel) purification kit and were sequenced. To determine the genotype of the sample, multiple alignment of sequences were obtained using ClustalW 1.8.

**Statistical analysis**

Data were statistically compared using the Chi-square and Fisher exact tests. Epi Info (2005) software was used to determine the Odds ratios (OR) of the associations between the various environmental and animal variables highlighted in the questionnaire and presence of Giardia organisms.

**Results**

*Giardia duodenalis* was detected by microscopy in 14 of the faecal samples (prevalence 11.0%, 6.2–17.8 CI 95%) and by PCR in 26 samples (prevalence 20.5%; 13.8–28.5 CI 95%). Given that the PCR detection method was more sensitive than microscopy, particularly when the level of infection was low (McGlade et al., 2003; Amar et al., 2004), only PCR detection data were used to evaluate associations with the other variables (Table 1).

There was a significant association between kennel identity and prevalence of *Giardia* infection ($\chi^2 = 18.5; P = 0.0001$), with a higher prevalence in K1 (46.9%) than in K2 (10.5%) or K3 (15.8%). Dogs housed in K1 had a much higher risk of acquiring *Giardia* infection than those in the other kennels (OR = 6.7; 2.4–19.2 CI 95%) and animals <1 year old had a significantly higher prevalence of infection (46.2%) than animals >1 year old (17.9%) (OR = 3.9; 1.0–15.0 CI 95%). When length of time kennelled was considered, the prevalence values for the three time categories of 0–3 months, 3–12 months, and >12 months, were 25.8%, 66.7% and 33.3%, respectively. The Fisher exact test indicated a significant difference between

**Table 1**

<table>
<thead>
<tr>
<th>Examined</th>
<th>Positive</th>
<th>Prevalence (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total number</td>
<td>127</td>
<td>26</td>
</tr>
<tr>
<td>Kennel identity</td>
<td>K1</td>
<td>32</td>
</tr>
<tr>
<td></td>
<td>K2</td>
<td>76</td>
</tr>
<tr>
<td></td>
<td>K3</td>
<td>19</td>
</tr>
<tr>
<td>Age</td>
<td>&lt;1 year</td>
<td>114</td>
</tr>
<tr>
<td></td>
<td>&gt;1 year</td>
<td>13</td>
</tr>
<tr>
<td>Length of time kennelled</td>
<td>&lt;3 months</td>
<td>31</td>
</tr>
<tr>
<td></td>
<td>3–12 months</td>
<td>9</td>
</tr>
<tr>
<td></td>
<td>&gt;12 months</td>
<td>6</td>
</tr>
<tr>
<td>Clinical signs</td>
<td>Yes</td>
<td>22</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>105</td>
</tr>
</tbody>
</table>

ND: not done.
the 0–3 month and 3–12 month categories ($P = 0.03$) and no significant differences between the remaining combinations of categories. *Giardia duodenalis* was detected in 13/22 (59.1%) of diarrhoeic dogs and in 13/105 (12.4%) samples of faeces of normal consistency indicating a significant association between the presence of infection and this clinical sign (OR = 10.2; 3.6–28.6 CI 95%). None of the other recorded variables (sex, access to common shared space, whether food was wet or dry, previous treatments) had any significant association with the prevalence of infection.

DNA from all samples found positive by PCR was successfully sequenced. Sequence analysis revealed 8 isolates of the zoonotic assemblage A (30.7%) and 18 of the host-specific genotypes C (14 isolates, 53.8%) and D (4 isolates, 15.3%). Isolates of different genotype were not found within samples. Because of the low numbers of genotypes detected it was no possible to test for significant associations between genotype and the specified variables.

**Discussion**

The overall prevalence of 20.5% *Giardia* infection identified in this study is similar to that previously reported in Denmark (17.1%) (Hansen et al., 2000), the former Yugoslavia (14.4%) (Nikolic et al., 2002), Norway (20.7%) (Hamnes et al., 2007) and Italy (15–32.1%) (Giangaspero et al., 2002; Capelli et al., 2003; Berrilli et al., 2004; Bianciardi et al., 2004; Capelli et al., 2006). Although Papini et al. (2005) and Széna´si et al. (2007) found high prevalences of infection of 55.2% and 58.8% in kennelled dogs using a *G. duodenalis*-specific coproantigen ELISA, the discrepancy found by Széna´si et al. (2007) between this study and the current study were the dog-specific genotypes C and D. Although this finding suggests that the zoonotic risk from this pathogen is perhaps less significant than previously thought, it must be remembered that the higher prevalence of infection in younger dogs may pose a specific public health issue as such animals are more frequently re-homed with families.

**Conclusions**

This study has identified *G. duodenalis* infection as a significant threat to the health of kennelled dogs. The prevalence of the infection may be underestimated given that it frequently occurs without concomitant clinical signs. The findings highlight the need for the regular surveillance of kennelled dogs for this parasite using specific diagnostic methods and with follow-up drug treatments and attention to kennel hygiene as appropriate.

**Conflict of interest statement**

None of the authors of this paper has a financial or personal relationship with other people or organisations that could inappropriately influence or bias the content of the paper.

**Acknowledgement**

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**References**


