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## Research Article

### DIAGNOSIS OF TOXOCARA CANIS IN DOGS IN BAGHDAD BY PCR TECHNIQUE

Afkar M. Hadi<sup>1\*</sup> and May H. Kawan<sup>2</sup>

<sup>1</sup>Natural History Museum and Researches Center/University of Baghdad

<sup>2</sup>Department of Parasitology/Veterinary Medicine Collage/University of Baghdad

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#### ABSTRACT

The study was determined the total infection rate of *Toxocara canis* 52% in the domestic and stray dogs (42.4% , 65.5%) respectively; at 215 fecal samples were collected from five different areas of Baghdad city that distributed as north, south, west, east and center, from March to October 2015.

Statistical analyses appeared that no significant differences  $p > 0.01$  among the areas of Baghdad city, no significant differences  $p > 0.01$  between sexes of dogs, there are significant differences  $p < 0.01$  between the ages of dogs that little puppies more sensitive for infection; and no significant differences  $p > 0.05$  among stray dogs.

Solid tissue and protocol procedure was applied to *T. canis* adult worm due to isolate a pure DNA. The result of DNA extraction showed that fresh tissue of adult worm yielded enough DNA concentration for PCR amplification that 100 ng with purity 1.6. The PCR was used for amplify the target fragment by using specific primers were designed manually in the *T. canis* specific part of ITS-2 Sequences. This study reported the development of sensitive and specific PCR assay allowing rapid and reliable identification of *T. canis* by the fragment size amplified was 380 bp in ITS-2 gene.

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#### INTRODUCTION

Pet dogs and cats can play an important role in transmission of zoonotic nematodes such as *Toxocara canis* and *Toxocara cati*, by excreting eggs directly into the human environment, without the involvement of vectors or intermediate hosts. Human toxocarosis remains a hazard despite the availability of highly effective anthelmintics for dogs and cats. A closer collaboration between veterinary and public health professionals within the 'One Health' concept is also required.[1]The close relationship of people with their companion animals, known as the human-animal bond, provides benefits with regard to socialisation, mental health and even physical wellbeing[2]. *Toxocara canis*, have an oral-faecal transmission cycle and humans can be infected by ingestion of infective eggs from contaminated soil (gardens, sandpits and play-grounds); from unwashed hands or raw vegetables, or by direct contact with pets [3,4].*T.canis* originally a parasitic nematode of canine, bitches and their puppies [5]. However, *T. canis*eggs can also hatch in a large number of non-canid species, including human beings [6]. Larvae in non-canid species hatch and distribute themselves throughout the viscera; most often these larvae become

encapsulated in a granulomatous response [7] or accumulated in the brain, where they elicit little or no histological reaction[8].The environmental contamination with *Toxocara canis* in a dog feces, especially in public parks, children playgrounds and streets [9,10,11,12].

The aim of present study is to determine the prevalence of *T. canis* in domestic and stray dogs with different ages and both sexes, in Baghdad city the capital of Iraq, include five regions: north, south, east, west and center. And Molecular diagnosis of infection of *T. canis* among dogs by PCR techniques.

#### MATERIALS AND METHODS

**Samples collection:** 215 fecal samples were collected from stray and domestic dogs randomly, in Baghdad city, which were divided into five areas: north area include Al-Shaab and Al-Rashdia; south area include Al-Ciadia and Al- Doraa; west area include Al- Kadhimiya and Al- Shulla; east area include Al-Ghadeer and Al-Sadder; center area include Al- Adamiyah and Bab Al-Muadam; from March to October 2015.

\*Corresponding author: Afkar M. Hadi

Natural History Museum and Researches Center/University of Baghdad

**Concentration technique**

**Sheather's solution preparation**

Sugar solution that was prepared as follow: 454 gm sucrose was dissolved in 355 ml distilled water [13]. This solution will grow mold if left out, so it would be added 6.5 gm phenol. Flotation method was done for all the fecal samples as previously described by [14].

**DNA extraction and PCR**

DNA extraction was performed according to manufacture's instruction kit using gSYNC DNA Extraction Kit (Geneaid), as the following:

**Sample preparation:** Adult worms of *T. canis* were obtained from infected dogs by treatment with anthelmintic drugs. Individual parasites were repeatedly washed in physiological saline, pH 7.3; A bout one centimeter was cut from *T. canis* worm tissue and put in centrifuge tube. 200 µl of GST Buffer was added and 20 µl of proteinase K, vortex thoroughly as suggested by [15].

**Genotyping:** PCR reaction was performed using specific primers were designed for the ITS-2 genes, primers were designed depending on their nucleotide sequence by Alpha DNA Company as a lyophilized product of different picomol concentrations. Lyophilized primers have been dissolved in a free DNase/RNase water to give a final concentration of (100 pmol/µl) (as stock solution), to prepare 10µM concentration as work primer resuspended 10 pmol/µl in 90 µl of deionized water to reach a final concentration 10µM. The sequences of these primers have been listed in (table 1) as [16].

**Table 1** Sequences of primers used in this study.

Gene marker	Primer name	Primer (5' → 3')	Primer length	Acc No. (NCBI)
Tcan1		(F): 5'-AGTATGATGGGCGCGCAAT-3'	20	AB110034
		(R): 5'-TAGTTTCTTTCCGCT-3'	19	AB110034

\*Primer from ALPHA Canadian company.

**PCR Programs**

The sample was put in the Thermocycler device for amplification of DNA and set the device program as follow: Denature template 94C at 30 sec.; Initial denaturation 94C at 60 sec.; Annealing 58C at 30 sec.; Extension 72C at 30 sec. that 35 cycle; Final Extension 72C at 10 min.

**Analysis of PCR Products**

The PCR products and the ladder marker have been resolved by electrophoresis. 3 µl of loading buffer was loaded on 1 % agarose gel (1g agarose/100 ml 1X TBE buffer) and run at 75 volt for 3-4 hours. The gel was stained with ethidium bromide for 15 minutes. In addition, bands were visualized on Gel documentation system with specific camera. Molecular marker ladder (100 bp) made by Bionner company has been used to estimate the molecular size of the bands.

**RESULTS**

The current study recorded high rate of infection 52% with *T. canis* in domestic and stray dogs. Out of 215 fecal samples

examined, 112 samples were positive by using sucrose flotation method.

High rate of infection with *T. canis* in dogs was recorded from different areas of Baghdad (north, south, west, east and center), out of 125 fecal samples were collected from domestic dogs, the rate of infection was 42.4%, and from 90 fecal samples of stray dogs, 65.5% was the rate of infection, (Table 2).

**Table 2** Total rate of infection with *T. canis* in dogs in Baghdad.

Canids	No. of fecal samples examined	No. of positive samples	%
Domestic dogs	125	53	42.4
Stray dogs	90	59	65.5
Total	215	112	52

All the study areas in Baghdad city showed high rates of infection with *T. canis* in domestic dogs, the high rate 47.6% in west of Baghdad, and the low rate 32% in the east of Baghdad. (Table 3).

**Table 3** Rates of infection in domestic dogs with *T. canis* in Baghdad according to areas.

Areas	No. of samples examined	No. of positive samples	%
North	24	11	45.8
South	23	9	39.1
Center	32	15	46.8
West	21	10	47.6
East	25	8	32
Total	125	53	42.2

Domestic dogs showed that total rate of infection in male with *T. canis* was more than female infection (46.2%, 38.6%) respectively. However, statistical analysis showed that there were no significant differences  $p > 0.01$  between sexes of dogs (Table 4).

**Table 4** Rates of infection in domestic dogs with *T. canis* in Baghdad according to sexes.

Areas	No. of male samples	No. of male positive	No. of female samples	No. of female positive
North	8	4	13	5
South	13	5	7	3
Center	12	6	10	4
West	8	5	8	3
East	13	5	13	5
Total	54	25 (46.2%)	51	20 (38.6%)

$\chi^2$  2.6\* No significant differences  $p > 0.01$  between sexes of dogs.

**Table 5** Rates of infection in domestic dogs with *T. canis* in Baghdad city according to ages.

Areas	No. of examined samples for (3-5) months puppies	No. of positive samples for (3-5) months puppies	No. of examined samples for (6-12) months dogs	No. of positive samples for (6-12) months dogs
North	2	2	1	-
South	2	-	1	1
Center	6	3	4	2
West	3	-	2	2
East	3	3	3	-
Total	16	8 (50%)	11	5 (45.4%)

$\chi^2$  9.4\* There are significant differences  $p < 0.01$  between the ages of dogs.

Furthermore, little puppies (3-5) months showed that total infection rate with *T. canis* more than total infection rate of young dogs (6-12) months (50%, 45.4%) respectively. Thus, statistical analysis showed that there are significant differences  $p < 0.01$  between the ages of dogs (Table 5).

Stray dogs showed high total infection rate 65.5%, and center of Baghdad showed the highest rate 100% while south of Baghdad showed the lower rate 52.5%. However, statistical analysis showed that there were no significant differences  $p > 0.05$  among stray dogs (Table 6).

**Table 6** Rates of infection in stray dogs with *T. canis* in Baghdad city.

Areas	No. of examined samples	No. of positive samples	%
North	16	9	56.25
South	19	10	52.6
Center	17	17	100
West	22	12	54.5
East	16	11	68.7
Total	90	59	65.5

$\chi^2$  0.8 \*No significant differences  $p > 0.05$  among stray dogs.

All the study areas of Baghdad city showed high rates of infection, the highest rate was 65.3% in center area, and the lowest rate was 45.2% in the south area for both domestic and stray dogs. Statistical analysis showed that there were no significant differences  $p > 0.01$  in the rates of infection among areas (Table 8).

**Table 8** Total rates of infection in domestic and stray dogs with *T. canis* in Baghdad city.

Areas	No. of examined samples	No. of positive samples	%
North	40	20	50
South	42	19	45.2
Center	49	32	65.3
West	43	22	51.1
East	41	19	46.3
Total	215	112	52

$\chi^2$  1.7 \*No significant differences  $p > 0.01$  among the areas.

### Genomic DNA extraction from worm

DNA extraction was done by using gSYNC DNA Extraction Kit (Geneaid) from a piece of *T. canis* worm which obtained 100 ng a pure DNA that purity was 1.6 for PCR amplification.

### PCR technique

The region spanning the ITS-2 was amplified from the gDNA (100ng) by PCR using. The amplicon sizes for *T. canis* were 380bp, (fig.1) primers were purchase from Alpha DNA, Canadian company.

## DISCUSSION

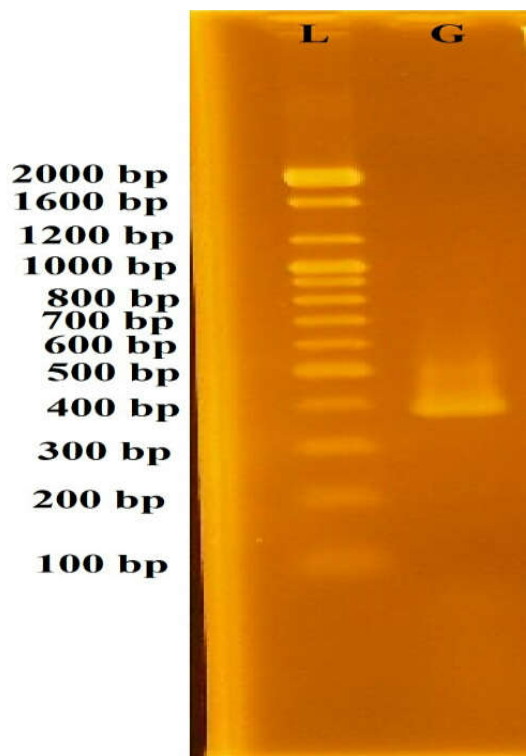
### Prevalence of *Toxocaracanis* in dogs in Baghdad city

*T. canis* infection in dogs is a public health problem in most countries, although it has been poorly documented in many of them, even Iraq. The current study recorded to high rate of infection with *T. canis* in dogs, the total rate of infection was 52%, in Baghdad city capital of Iraq, indicates that there is high risk for human toxocariasis and there should be a prompt evaluation of the significance of *T. canis* for public health. The prevalence of *T. canis* depend on many factors that can influence in the rate of infection including age, sex, source of

dogs, aggregation of stray dogs, high offspring and distribution puppies.

The current study determined the prevalence of *T. canis* in dogs in Baghdad city that include five areas: north, south, west, east and center, the high rate of infection was 65.3% in the center area of Baghdad, and the low rate 45.2% in the south area of Baghdad for both domestic and stray dogs, that may be due to the difference in density stock and the number of dogs, however, statistical analysis showed that there were no significant differences in the rates of infection among areas, that similar to [17] who denied the differences between Urban and rural areas in Basrah city. Furthermore, [18] reported to the number of dogs infected with *T. canis* that varied at three main stations in Basrah city, the highest was shown in the north station and there were no significant differences among three stations in Basrah city.

Stray dogs showed high rate of infection with *T. canis* 65.5% that acceptance with [19] who recorded 67.5% in Baghdad city. While, in Sulaimani province, Kalar city, [20] found the rate of infection with *T. canis* was 36% in stray dogs. In Mosul province [21] Pointed out that the percentage of infection of 35 stray dogs with *T. canis* was 25.7%, while, [22] reported that 19 dogs from 58 examined dogs (32.6%) were infected with *T. canis* in Mosul city, and in Najaf province [23] showed that 46% of dissection stray dogs were infected with *T. canis*. [17] During his survey on intestinal nematodes found that the percentage of infection with *T. canis* of 51 dissected stray dogs in Basrah was 23.5%. [24] Recorded 6% rate of infection rate with *T. canis* between stray dogs in Konya province/ Turkey. While, [25] reported that 66.6% was found to be infected with *T. canis* from 21 stray dogs examined in Izmir/ Turkey. These differences in rates of infection were due to the difference of climates (months of study) and number of samples collected.



**Fig.1** PCR product (380 bp) of fragment of ITS-2 gen of *T. canis* adult worm.

The infection rate in stray dogs was 65.5% higher than the infection rate in domestic dogs 42.4% that was similar to [26] in New Zealand, who showed that higher rate of infection with *T. canis* among stray and unwanted pets than in well cared dogs. 42.4% rate of infection in domestic dogs that recorded in current study disagreement with [27] who showed that the percentage of infection with *T. canis* eggs was 6.55% when they examined 305 fecal samples from pet dogs in Tehran/ Iran; that may be due to varied from one country to another depending on the distribution of veterinary hospitals, monthly examination of dogs and dog's treatments.

No significant differences were recorded in the rates of infection between male and female dogs (46.2%, 39.2%) respectively in current study, this result dislike with [17] results, who revealed that female dogs were more infected than male in Basrah this was due to the differences in samples that collected. The results were similar to [28] in California, who reported that sex of the host had no significant influence on the degree of parasitism with *T. canis* infection. Also, [29] showed that the prevalence of *T. canis* infection was similar in both sexes in Kinshasa, Zaire. Furthermore, [30] showed that male's dogs were more often infected than females and no significant differences between them in Calabar, Nigeria.

According to age our study recorded that *T. canis* infection in puppies (3-5) months of age was higher than older puppies (6) months of age and up, that because the close contact between small puppies and their mothers that transport the infection either by: transplacental, transmorary, or direct contact. This result was similar to [31] who examined 76 puppies aged between 2-4 months in Japan and showed that 68.4% were infected with *T. canis*. While, [32] pointed out that *T. canis* was the major parasite in all age groups of dogs after they examined 1126 dogs in Japan.

#### Molecular study

Solid tissue and protocol procedure was applied to *Toxocaracan* is adult worm to isolate a pure DNA, for the first time in Iraq. The result of DNA extraction showed that fresh tissue of adult worm yielded enough DNA concentration for PCR amplification that 100 ng with purity 1.6 this result was similar to the quantification of DNA by spectrophotometer which revealed that the DNA concentration ranged between 6-10 ng /  $\mu$ l and purity rang was between 1.6 – 2 [33].

This study reported the development of sensitive and specific PCR assay allowing rapid and reliable identification of *T. canis* by the fragment size amplified was 380 bp in ITS-2 gene as shown in figure 1; by using specific primers, the targeted fragment was determined according to Gen bank, [AB110034]. This result was acceptant with [16] who revealed that the positive control of *T. canis* consist of 380bp. While, the result was not acceptance with [34] who represented PCR with 14ng of *T. canis* DNA, isolated from dog; and 10ng of *T. canis* DNA, isolated from fox with 587bp for both. The causes of these difference were either due to different amount of DNA or differences of primers. The second internal spacer (ITS-2) of gDNA had proven in current study was particularly valuable in this context for molecular studying of *T. canis*, that similar to [15, 34, 35, 36] whom extracted genomic DNA from adult worms that collected from dogs, foxes and cats. Furthermore,

[37, 38] proved that the advantage of DNA-based methods was a high genetic variability within molecular markers such as ITS2 for the discrimination of *T. canis* and *T. cati* from their closely-related neighbors: *T. leonina*, *T. vitulorum* and *T. malaysiensis*. Previously, [39] compared between target gene ITS-1 and ITS-2; thus, they revealed that the higher degree of within – species variation in the ITS-1 sequences compared with the ITS-2 suggests that the later sequences may provide more reliable species markers for Ascaridoid nematodes. [15] Revealed that the PCR method has been a potential as a supporting technique for the diagnosis of human toxocariasis and infected liver of Mongolian gerbils experimentally infected by. Furthermore, [40] described a novel real-time PCR that, the availability of a validated direct molecular assay for *Toxocara* spp. DNA in BAL will improve diagnosis of human toxocarosis, particularly for patients with respiratory distress. Recently, [41] compared among the performance of three diagnostic tests, that was, histopathology, enzyme-linked immunosorbent assay (ELISA), and polymerase chain reaction (PCR), for the diagnosis infection of *Toxocaracati* in the orbital tissue of Mongolian gerbils and Wistar rats experimentally. That of the three evaluated methods, the PCR could be recommended for scientific and laboratory diagnoses of ocular toxocariasis in experimentally infected animals.

The current study revealed that Molecular techniques provide useful tools for the genetic characterization of *T. canis* in Iraq for first time, and can use for diagnosis of Toxocariasis in human tissues in the future.

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