Toxoplasma gondii in four captive kangaroos (Macropus spp.) in China: Isolation of a strain of a new genotype from an eastern grey kangaroo (Macropus giganteus)

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**ABSTRACT**

Marsupials are highly susceptible to *Toxoplasma gondii* infection. Here, we report *T. gondii* infection in four kangaroos from a zoo in China. Kangaroos were imported into China in 2000 and were since bred in zoo. In 2017–2018, four kangaroos died due to respiratory system disease or injury. The bodies were submitted to the laboratory to test for *T. gondii* infection. Antibodies to *T. gondii* were found in 75% (3/4) of the kangaroos via the modified agglutination test with the cut-off 1:25. Cysts were observed in the histopathological sections of tongue and diaphragm or squashes of fresh myocardium in two kangaroos. These cysts were confirmed as *T. gondii* by immunohistochemical staining and molecular biological analysis. One viable *T. gondii* strain was isolated from one kangaroo and designated as TgRooCHn1. DNA from *T. gondii* tachyzoites obtained from cell culture was characterized by 10 PCR-RFLP markers and the virulence genes ROP5 and ROP18. The genotype of this isolate did not match with any known genotypes; it was designated as ToxoDB#292. The virulence of TgRooCHn1 (10^6 tachyzoites) was non-lethal to mice, and it formed tissue cysts. To our knowledge, the present study is the first isolation of ToxoDB#292 strain from kangaroo. Improvements for captive settings were initiated, including greater attention being paid to birds and stray cats, fed frozen meat for carnivores.

1. Introduction

*Toxoplasma gondii* infects virtually all warm-blooded animals, including birds, humans, livestock, and wild animals (Dubey, 2010). Felids are the definitive hosts, and other warm-blooded animals can be intermediate hosts. Most animals are resistant to *T. gondii* infection, and most infections are subclinical; however, some species such as new world primates, squirrel monkeys, lemurs, marmosets, meerkats and kangaroos are very susceptible to the infection (Dubey et al., 1988; Epiphanio et al., 2003; Cenci-Goga et al., 2011; Nishimura et al., 2019). Horizontal transmission by *T. gondii* cysts or oocysts is likely the most important mode of transmission of *T. gondii* for these animals.

Initially, kangaroos (Macropus spp.) in zoos around the world were imported from Australia and New Zealand; some zoos then started to breed these animals locally. Mortality of captive marsupials infected with *T. gondii* has been reported, and some animals have also been reported to develop chronic infection (More et al., 2010; Guthrie et al., 2017; Dubey, 2010; Diaz-Ayala et al., 2016). There are few reports concerning genotyping of *T. gondii* from kangaroos and wallabies. Two *T. gondii* strains with ToxoDB #1 and #2 were isolated from kangaroos from zoos in Argentina (More et al., 2010). Three *T. gondii* strains isolated from wallabies were ToxoDB #2 (Dubey and Crutchley, 2008). Non-archetypal type II-like strains of *T. gondii* were found in marsupials from Australia (Parameswaran et al., 2010). ToxoDB#263 and ToxoDB#4 *T. gondii* strains were identified in wallaby tissues from the Virginia Zoo (Guthrie et al., 2017).

Kangaroos usually eat grass and grassroots, and infection of *T. gondii* is a useful indicator of soil and environmental contamination by

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Table 1: Background and isolation of *Toxoplasma gondii* from kangaroos in China.

| Case | Species | Sex | Age in year | Sample received | Clinical signs | Pathological findings | MAT | PCR | Bioassay | Date received | Date assayed | T. gondii cysts | Date Measured | Number of positive mice/number of inoculated mice |
|------|---------|-----|-------------|----------------|---------------|----------------------|-----|-----|----------|---------------|---------------|----------------|---------------|-----------------------------------------------|
| 1    | *Macropus rufus* | Female | 5 | 9/30/2017 | Nasal discharge, depressed, anorexia. | Interstitial pneumonia, necrotic splenitis, necrotic hepatitis, *T. gondii* cysts. | + | + | + | 6/24/2017 | 7/31/2017 | Not found | 9/12/2018 | 0/5 0/2 |
| 2    | *M. giganteus* | Female | 6 | 7/23/2017 | Without obvious clinical signs. | Bacteria, liver fatty degeneration, necrotic splenitis, *T. gondii* cysts. | + | + | + | 7/24/2017 | 7/24/2017 | Not found | 9/12/2018 | 0/5 0/2 |
| 3    | *M. giganteus* | Male | 5 | 7/26/2017 | Nasal injury, intermittent muscle spasms, dyspnea. | Neuronal necrosis. | + | + | - | 7/31/2017 | 8/3 | - | 9/12/2018 | nd |
| 4    | *M. rufus* | Male | 4 | 9/11/2018 | Depression, hematuresis, salivation, skin injury, anorexia. | Acute splenitis, interstitial pneumonia, glomerulonephritis. | + | - | - | - | nd | nd | nd | 0/2 |

**nd:** experiment not done.

Kangaroos were bioassayed in mice following previously published methods (Dubey, 2010).

**2.2. Detection of antibodies against *T. gondii* and *T. gondii* DNA in the serum and tissue digests of kangaroos**

Kangaroo meat juice was serially diluted from 1:25 to 1:200 (case 1–3) and 1:25 to 1:12,800 (case 4) and tested for antibodies to *T. gondii* using the modified agglutination test (MAT) as described by Dubey and Desmonts (1987). Whole formalin fixed *T. gondii* antigens were obtained from the University of Tennessee Research Foundation (Knoxville, TN, USA; https://utrf.tennessee.edu/). A titer of 1:25 was considered indicative of exposure to *T. gondii*. The sera of positive controls from mice inoculated with *T. gondii* Me 49 strain and negative controls from mice were included in the same 96-well plate. Digested tissues (myocardium, leg muscle, tongue, and diaphragm) from kangaroos were used to test *T. gondii* DNA. The DNA from the digested samples was extracted using a commercial DNA extraction kit (Tiangen Biotec Company, China, DP 304) and eluted in 50 μl deionized water. PCR assays were done to detect *T. gondii* using the specific primer pairs TOX5/TOX8 (5′-GCAGCGTACGACAGTGCATCTGGATT-3′ and 5′-CCCAGTCTGCGTCTGTCGGGAT-3′) (Schares et al., 2008). Quality control of PCR analyses including eliminating nucleic acid contaminating and running negative and positive controls was conducted (Burkardt, 2000). Negative PCR controls (not including template DNA) and the positive control DNA from *T. gondii* Me 49 were included in all batches. The PCR products from *T. gondii* were expected to be 450 bp in length.

**2.3. Histopathological analysis**

The tissues were processed using routine histological techniques and then embedded in paraffin. Paraffin sections (5 μm in thickness) of the samples were prepared and stained with hematoxylin and eosin (H&E). Immunohistochemistry (IHC) was conducted using rabbit anti-*T. gondii* serum as the primary antibody and mouse anti-rabbit IgG conjugated with HRP/DAB as the second antibody (IHC detection kit, Abcam, ab64264). Brain tissue sections of a VEG *T. gondii*-infected mouse were used as a positive control for IHC (kindly provided by Dr. Dubey, ARS, USDA).

**2.4. Isolation of viable *T. gondii* from kangaroo tissues by bioassay in mice**

Tissue samples (hearts, tongues, diaphragm, and leg muscle) of four kangaroos were bioassayed in mice following published methods (Dubey, 2010). Briefly, striated muscle (50 g) was homogenized, digested in pepsin (5.2 g pepsin, 10.0 g NaCl, and 14 mL HCl diluted to 1 L...
with deionized water, pH 1.1–1.2). The striated muscle homogenate was subsequently incubated at 37 °C in a shaking water bath for 60 min. After this, the sample was filtered through double gauze and centrifuged at 1200×g for 10 min. The supernatant was then removed, and the pellet was suspended in 0.01 M PBS (pH 7.2) and neutralized by mixing with 1.2% sodium bicarbonate. Following mixing, the sample was centrifuged at 1200×g for 10 min, after which the supernatant was removed and 5 mL of saline containing 1000 units penicillin and 100 μg of streptomycin per mL was added. Striated muscle digested liquid was then inoculated subcutaneously into mice (1 mL per mouse). The homogenates were inoculated subcutaneously into 2–5 BALB/C mice or/and gamma interferon (γ-IFN) knockout mice. Survivors were bled on day 60 post-inoculation (DPI), and 1:25 and 1:200 dilutions of mouse serum were tested for *T. gondii* antibodies using MAT. Mice were euthanized at 61 DPI, and brain cysts were examined and counted as described by Dubey et al. (2012). Smears of tissues of dead or euthanized mice were examined for *T. gondii* tachyzoites or cysts. If cysts or tachyzoites were not found in the mouse tissue, homogenized lung, brain, heart, and tongue were used for subpassage into new groups of mice.

2.5. *In vitro* cultivation and genotyping

Brain homogenates of *T. gondii*-positive mice were seeded into CV-1 cell culture flasks as described previously by Dubey (2010). DNA was extracted from cell culture-derived tachyzoites. Multiplex PCR of the *T. gondii* strain was performed using 10 PCR-RFLP genetic markers (SAG1, SAG2 (5′-3′ SAG2, alt.SAG2), SAG3, BTUB, GRA6, c22-8, c29-2, L358, PK1, and Apico) as previously described by Su et al. (2010). Genotyping of virulence genes ROP5 and ROP18 was also performed as previously reported (Rego et al., 2017; Cheng et al., 2017). Reference *T. gondii* DNA was included in all batches.

2.6. Evaluation of the pathogenicity of *T. gondii* tachyzoites isolated from kangaroo

Virulence of the *T. gondii* isolated from the kangaroo was evaluated in BALB/C mice. *T. gondii* tachyzoites collected from cell cultures were counted in a disposable hemocytometer and diluted in 10-fold steps from 10<sup>-1</sup> to 10<sup>-6</sup> to reach an end-point of < 1 tachyzoite. Then, < 1, 10<sup>0</sup>, 10<sup>1</sup>, 10<sup>2</sup>, 10<sup>3</sup>, and 10<sup>4</sup> tachyzoites were intraperitoneally inoculated into five BALB/C mice for each dilution. The clinical symptoms were recorded, and mice were monitored daily. The smears of
lung or mesenteric lymph nodes of sacrificed mice were examined for T. gondii tachyzoites. After 60 days, all surviving mice were bled and tested for IgG antibodies to T. gondii using the MAT assay with titers between 1:25 and 1:200. The mice were euthanized at 61 DPI, and cysts of T. gondii were counted in the brains of all mice using a squash preparation (Dubey et al., 2012). Mice were considered infected when either antibodies to T. gondii or parasites were detected in their sera or tissues.

2.7. Ethics

This study was approved by the Institutional Animal Use Protocol Committee of the Henan Agricultural University, China. The Beijing Association for Science and Technology (Approval SYXK [Beijing] 2007–0023) approved the protocol used in this study. We handled all mice in strict accordance with the good animal practices of the Animal Ethics Procedures, Recycle-Reduce-Reuse principle, and Guidelines of the People's Republic of China.

2.8. Statistical analysis

Statistical analysis was performed using Graph Pad Prism 4.0 software (Graph Pad Software Inc., San Diego, CA, USA). Data were analyzed with the chi-squared test or Fisher’s exact test. A value of p < 0.05 was considered statistically significant.

3. Results

3.1. Necropsy and pathologic findings

Four dead kangaroos were collected from the zoo during 2017–2018 (Table 1). Case 1 kangaroo presented signs of yellow nasal discharge, depression and anorexia. Postmortem findings were pulmonary congestion, spleen atrophy, and enlarged liver. Microscopic analysis showed interstitial pneumonia, necrotic splenitis and hepatitis. Case 2 kangaroo died suddenly without obvious clinical signs. The gross findings were pulmonary and liver congestion, pleural effusion, and spleen atrophy. Many rod-shaped bacteria were present in the alveolar edema fluid and blood in the lung. Liver fatty degeneration, necrotic splenitis, and T. gondii cysts were observed in the tongue and diaphragm (Fig. 1). Case 3 kangaroo displayed a history of nasal injury from fighting with other kangaroos. Intermittent muscle spasms and dyspnea were observed. Multifoial glial nodules with neuronal necrosis were present in the brain, along with alveolar expansion and rupture. Case 4 kangaroo showed signs of depression, salivation, skin injury, and anorexia. This kangaroo developed hematuria and failed to respond to penicillin, streptomycin or supportive therapy. Gross examination showed pulmonary and kidney congestion, as well as splenomegaly. The histopathological findings were acute splenitis, interstitial pneumonia, and glomerulonephritis.

3.2. Immunohistochemical staining and PCR

Under light microscopy, many oval-shaped cysts were observed in the skeletal muscle cells of the tongue and diaphragm of case 2 (Fig. 1A and B). T. gondii cysts were confirmed with IHC-staining (Fig. 1C and D). The average cyst load was 17.1 ± 10.8 per square centimeter, and the sizes of the cysts were within the range of 5.2 μm–8.0 μm × 2.4 μm–6.1 μm by H&E sections. No inflammation was found around the cysts. Round cysts were found in squashes of fresh myocardium from case 4 kangaroo (Fig. 1E). The average cyst load was 4.2 ± 2.6 per square centimeter, and the sizes of the cysts were 42.5 μm–68.0 μm × 47.6 μm–70.2 μm by squash sections. T. gondii DNA in the tissue digests from all kangaroo cases was checked. However, T. gondii DNA using the primers TOX5/TOX8 was detected only in case 2 and 4 (Table 1).

3.3. Serological examination and isolation of T. gondii in mice and cell cultures

Antibodies to T. gondii were found in three of the four kangaroos (75%), with titers 1:200 (Table 1). The striated muscle homogenates from all kangaroos were individually tested via bioassay in mice. One group of mice (5/5) was inoculated with tissue samples from case 2 kangaroo, and the mice were sacrificed after showing signs of toxoplasmosis 22–27 DPI. Tachyzoites were found in the lungs, and many small cysts (diameter around 5 μm–50 μm) were found in the brains of the mice (Fig. 1F). This isolate was successfully propagated in cell culture after seeding cell 14 days from T. gondii positive mouse tissues, and the strain was designated as TgRooChn1. Genetic typing using 10 genetic markers revealed that this isolate did not match with any known genotypes. This isolate was designated as a new ToxoDB number #292, atypical type III. In addition, the isolate showed ROP5 allele 3 and ROP18 allele 3 (Table 2).

3.4. Virulence evaluation of TgRooChn1

After inoculating BALB/C mice with different dosages of TgRooChn1 tachyzoites, all the mice survived and showed no clinical signs. No mice died before termination at 60 days (Table 3). Tachyzoite

| Isolated ID | SAG1 | (3’ + 5’) SAG2 | Alt SAG2 | SAG3 | BTUB | GRA6 | C22-8 | C29-2 | L358 | PK1 | Apico | ROP5 | ROP18 | Genotype ToxoDB |
|-------------|------|---------------|----------|------|------|------|-------|-------|------|------|-------|------|-------|----------------|
| GT1, reference | I    | I             | I        | I    | I    | I    | I     | I     | I    | I    | 1     | 1    | 1     | #10             |
| PGT, reference | II   | II            | II       | II   | II   | II   | II    | II    | II   | II   | 2     | 2    | #1               |
| CTG, reference | II/III | III          | III     | III   | III   | III   | III    | III    | III   | III   | 3     | 3    | #2              |
| TgCatBr64, reference | I    | II            | II       | III   | II    | II    | u-1   | I     | u-2   | 5     | 2    | #66             |
| TgCgCa1, reference | II   | II            | II       | III   | III   | III   | III    | III    | III   | III   | 3     | 3    | #17             |
| TgCgCa1, reference | u-1  | I             | II       | III   | III   | III   | III    | III    | III   | III   | 3     | 3    | #111            |
| TgRooChn1 | u-1  | I             | II       | III   | III   | III   | III    | III    | III   | III   | 3     | 3    | #52             |

Table 2: Genotypes of T. gondii isolates from kangaroos in China according to PCR-RFLP of 10 markers and virulence proteins.

Table 3: Virulence of T. gondii strain TgRooChn1 in BALB/C mice.

| No. of tachyzoites | No. of mice infected/No. of mice inoculated (%) | Brain cysts by day 61 DPI |
|--------------------|-----------------------------------------------|--------------------------|
| 10<sup>4</sup>     | 5/5(100%)                                     | 386.0 ± 97.0<sup>b</sup>  |
| 10<sup>3</sup>     | 5/5(100%)                                     | 502.5 ± 147.4<sup>b</sup>|
| 10<sup>2</sup>     | 5/5(100%)                                     | 224.4 ± 47.5<sup>a</sup> |
| 10<sup>1</sup>     | 3/5(60%)                                      | Not found                |
| 1                  | 2/5(40%)                                      | Not found                |
| < 1                | 0/5(0%)                                       | Not found                |
| Blank control      | 0                                             | Not found                |

<sup>a,b</sup> Different letters indicate significant differences (p < 0.05).
numbers above 10^2 induced T. gondii infection in all of the mice. T. gondii cysts were detected in the brains of mice when euthanized at 61 DPI. The number of brain cysts in the group of mice inoculated with 10^2 tachyzoites had 224.4 ± 47.5 cysts. The numbers were significantly increased in groups inoculated with 10^3 tachyzoites (502.5 ± 147.4) or 10^4 tachyzoites (386.0 ± 97.0) (P < 0.05). No obvious pathological lesions were observed in mice infected with the TgRooCHn1 strain at 61 DPI.

4. Discussion

In this study, T. gondii infection was identified in four kangaroos from a zoo in China. Three out of the four kangaroos were positive for T. gondii IgG. Meat juice from one kangaroo’s (case 4) heart was seronegative for T. gondii IgG (titer from 1:25 to 1:12,800); however, T. gondii DNA and T. gondii-like cysts were found in this animal. It may be because due to acute infection, and anti-T. gondii IgG antibody was not produced yet. Unfortunately, isolation T. gondii was unsuccessful from case 1, case 3 and case 4 kangaroos.

T. gondii cysts in the tongue and diaphragm of case 2 kangaroo were observed, and this was further confirmed by specific T. gondii PCR from digested tissues, by cysts reactive to T. gondii antibody by IHC in histopathological sections (Fig. 1 C and D), and bioassays in mice. Histopathological evidence of T. gondii in tissues has been found in the skeletal muscle, adrenal gland, brain, heart, liver, stomach, intestine, and mesenteric lymph node of marsupials (Dubey et al., 1988; Fernandez-Aguilar et al., 2013; Basso et al., 2007; Bermudez et al., 2009; More et al., 2010; Díaz-Ayala et al., 2016). A large number of T. gondii cysts were detected in the skeletal muscles of case 2 kangaroo and the brains of inoculated mice. This is in agreement with the findings of More et al. (2010). Furthermore, T. gondii-specific DNA was detected in the brain or tongue of all seropositive kangaroos, indicating a high level of infection by T. gondii in this species (Parameswaran et al., 2009; Guthrie et al., 2017). The infection level of T. gondii cysts may also be related to the genotype or strain of T. gondii, the infective dose, the host species, and the immunity of hosts. The heavy load of T. gondii cysts in kangaroo tissues may be an indicator of the susceptibility of this species. As kangaroo meat is available for human consumption, meat contaminated with T. gondii can be a risk factor for transmission of the parasite to humans.

T. gondii was successfully isolated from kangaroo by bioassay in mice. Genotyping revealed that TgRooCHn1 is a new type and designated as ToxoDB #292. Experimental infection in mice showed that TgRooCHn1 is avirulent and can readily establish chronic infection and form tissue cysts in mice. In this study, the ROP18/ROP5 genotype combination suggests this strain TgRooCHn1 is non-lethal to mice (Shwab et al., 2016), which matched with the mouse virulence test in this study. To date, only a few T. gondii genotypes have been identified in animals and humans in China. These genotypes include ToxoDB #1 (type II), #3 (type II variant), #4, #9, #10 (type I), #204, and #225, and the ToxoDB #9 is the dominant type in China (Dong et al., 2018). There was no T. gondii inspection when the kangaroos were imported from Australia to China. 45 distinct T. gondii genotypes were detected directly from tissues of 16 macropods (Shuting et al., 2012). However, few isolates of T. gondii from Oceania have been genetically characterized using multilocus PCR-RFLP genotyping. ToxoDB#3 and ToxoDB#1 are the predominant genotypes in animals from Oceania, with ToxoDB#10 also being detected (Brennan et al., 2016; Shwab et al., 2014). However, because these kangaroos were born in captivity, they may have ingested T. gondii oocysts shed by stray cats or captive felids in the zoo. The ToxoDB#292 genotype may have already been circulating in local source, as genetic diversity may exist in China. It is important to investigate T. gondii infection in more animals, especially cats and small mammals. It is not clear that the mortality of the kangaroos in this study was solely due to T. gondii infection or whether the infection was mixed with other pathogens. Improvements for the captive setting were implemented after this study; more attention was paid to birds and stray cats, and the frozen meat fed to carnivores.

Declarations

Abbreviations

MAT: modified agglutination test; H&E: hematoxylin and eosin staining; IHC: immunohistochemistry.

Consent for publication

All authors consent for publication of this report.

Availability of data and material

The data, materials, and protocols associated with this report may be shared by the readers without undue qualifications.

Competing interests

The authors declare no competing interests. None of the authors of this report have financial or personal relationships with other people or organizations that could inappropriately influence its content.

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Authors’ contributions

RJS performed laboratory tests and IHC analysis. HD performed laboratory tests and PCR-RFLP analysis. NJ participated in laboratory tests and PCR-RFLP analysis. YRY designed the study protocol, analyzed the results and wrote the manuscript. All authors have read and approved the final version of the manuscript.

Conflicts of interest

None.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ijppaw.2019.03.003.

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