Occurrence of Atypical and New Genotypes of *Toxoplasma Gondii* in Free-Range Chickens Intended for Human Consumption in Brazil

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Received: 30 July 2019 / Accepted: 20 February 2020 / Published online: 5 March 2020
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Abstract
Purpose The aim of the present study was to detect and genotype *T. gondii* in free-range chickens destined to human consumption in Alagoas state, Brazil.

Methods Two hundred blood samples were collected from free-range chickens and submitted to indirect immunofluorescence antibody test (IFAT). Brain tissue from 14 animals randomly selected were subjected to mouse bioassay. Positive samples in mouse bioassay were submitted to PCR and genotyped by PCR–RFLP.

Results Out of two hundred blood samples from chickens, 72 (36%) samples were considered positive by IFAT. Two *T. gondii* strains were isolated, both being characterized as atypical and classified as #146 and a new genotype, named #279 in ToxoDB database.

Conclusions Our results showed a sero-occurrence of *T. gondii* in free-range chickens intended for humans, and the genetic diversity of the parasite in Brazil, with a new genotype described.

Keywords Toxoplasmosis · Genotyping · PCR–RFLP · Zoonosis

Introduction

*Toxoplasma gondii* is a protozoan parasite that can infect all warm-blooded animals and is widely prevalent in humans and animals worldwide; usually the infections is asymptomatic; however, congenital infection can cause damage to the fetus and reproductive disorders in humans and animals [1]. Humans mainly become infected by ingesting undercooked or raw meat contaminated with tissue cysts, or by ingesting food or water contaminated with *T. gondii* oocysts [2].

Due to their feeding habits, free-range chickens are considered good indicator of soil contamination by *T. gondii* oocysts and an important source of infection for humans, being used as sentinel animals in regions with high prevalence of human infection [3].

Nowadays, there is a higher demand by consumers for differentiated products with superior quality, which has been influencing changes in the animal production systems. Although there is an increase in production systems handling with animal welfare, extensive system may present risks if it is not well managed, as the animals are reared loosely and may have more contact with potential pathogenic agents such as *T. gondii* [4].

In the past, *T. gondii* population structure was considered clonal, being *T. gondii* strains classified as Types I, II or III based on SAG2 marker [5]. However, after developing of other molecular markers, studies from different regions of the world, especially in Central and South America, have shown that *T. gondii* has a high genetic diversity, with high prevalence of atypical strains [6, 7]. In mice, the
Clonal genotypes type I are classified as high virulence ($LD_{100}=1$), while types II and III are classified as medium to low virulence ($LD_{100} \geq 10^3$) [5]; however, atypical genotypes are highly virulent to mice, with correlation of ROP18 and ROP5 alleles with virulence [8]. In humans, atypical genotypes have been reported as cause of congenital ocular toxoplasmosis and acute systemic toxoplasmosis in immunocompetent individuals, showing a potential association between the atypical genotype and pathogenicity in human hosts [8–10].

The aim of the present study was to evaluate the occurrence of anti-\textit{T. gondii} antibodies and genetically characterize isolates obtained from free-range chickens from Brazil.

**Material and Methods**

**Animals and Study Area**

Two hundred free-ranging chickens (\textit{Gallus gallus domesticus}) from seven different farms belonging to municipality of Viçosa, state of Alagoas, northeast Brazil, had their blood taken, between February and July 2016. Blood samples were collected by puncture of the brachial vein and serum samples obtained were stored at $-20$ °C until further analysis. Fourteen chickens from all seven farms were randomly selected and had their brain tissue collected aseptically, followed by mouse bioassay (Table 1). All procedures involving animals were conducted in accordance with good animal practices and approved by Animal Ethics Committee (number 43/2015).

**Serology**

Serum samples obtained from chickens and mice from mouse bioassay were subjected to immunofluorescence antibody test (IFAT) according to methodology previously described [11]. Tachyzoites of RH strain were used as antigen, and serum samples from chickens known as positive and negative for \textit{T. gondii} were used as controls and included in all slides. Only serum samples that showed fluorescence over the whole parasites’ surface were considered positive. Samples with titers with $\geq 16$, for both chickens and mice, were considered positive.

**Mouse Bioassay**

Mouse bioassay was performed according to methodology previously described [12]. Brain tissue (30 g) from each animal was macerated, homogenized with 0.85% NaCl solution (saline) 1:5 (w/v), filtered through gauze and centrifuged at 700×g for 10 min. The supernatant was discarded, the pellet resuspended with 50 ml of saline solution and centrifuged again. Next, the supernatant was discarded, the pellet resuspended in 2 ml of antibiotic saline solution (1000 IU of penicillin and 100 µg of streptomycin per ml) and inoculated intraperitoneally into two Swiss Webster (SW) mice. Mice were observed daily and those that showed clinical signs of infection were euthanized for collection of peritoneal liquid for detection of tachyzoites. Mice who survived for 45 days after inoculation were euthanized and blood and brain tissue samples were taken for antibodies and tissue cyst detection, respectively. Positive samples for presence of \textit{T. gondii} were inoculated in MARC-145 cells for growth of the parasite [13].

**DNA Extraction and PCR**

DNA extraction was performed using a commercial kit (DNeasy Blood and Tissues Kit, Qiagen®, Germany) following the manufacturer’s instructions. \textit{T. gondii} DNA was detected by a nested PCR, using the external primers TgNN-TgNN2 and internal primers TgNP1-TgNP2, amplifying a fragment of 227 bp of the ITS1 region of the parasite [14]. DNA of RH strain tachyzoites and ultrapure water were used as positive and negative controls, respectively, and included in all PCR run. PCR products were subjected to electrophoresis on 1.5% agarose gel stained with Sybr Safe DNA Gel Stain (Invitrogen®, USA), and visualized and photographed under UV light.

**Multilocus PCRP–RFLP**

Genetic characterization of the isolates was performed by polymerase chain reaction—restriction fragment length polymorphism using 11 genetic markers (SAG1, SAG2, alt. SAG2, SAG3, BTUB, GRA6, c22-8, c29-2, L358, PK1 and Apico) as previously described [7]. All products were subjected to electrophoresis on 3% agarose gel stained with Sybr Safe DNA Gel Stain (Invitrogen®, USA), visualized and photographed under UV light. The results obtained were

| Farms | Blood samples | Brain samples |
|-------|---------------|---------------|
| A     | 10            | 1             |
| B     | 10            | 1             |
| C     | 5             | 1             |
| D     | 38            | 2             |
| E     | 43            | 4             |
| F     | 51            | 3             |
| G     | 43            | 2             |
| Total | 200           | 14            |
compared and classified according to the genotypes present in ToxoDB in (https://toxodb.org/toxo/). Phylogenetic tree comparing the isolates with the reference strains was constructed using the Split Tree software version 4.13 [15].

Results and Discussion

Antibodies against *T. gondii* were observed in 36% (72/200) according with IFAT. Animals presented titers of 16 (8.22%), 32 (10.96%) 64 (17.8%), 128 (13.69%), 256 (19.17%) and 512 (30.14%). Previous studies performed in free-range chickens in different regions from Brazil have shown a prevalence ranging from 10 to 100% [16, 17]. In the northeast region of Brazil, studies conducted with backyard chickens have described a prevalence of *T. gondii* up to 40.56% by means the IFAT, and 94.8% of studied farm with at least one positive animal [18, 19], similar to our results, where 36% of animals and 100% of farms were considered positive.

In the mouse bioassay, *T. gondii* was isolated from two seropositive chickens, with titers of 16 and 64. The obtained isolates, named as TgCkAL01 and TgCkAL02, were obtained from tissue cyst of euthanized mice. One mouse developed clinical signs of chronic infection at 37 days post-inoculation, being euthanized for collection of peritoneal liquid and brain tissue. These biological samples were inoculated in MARC-145 cells for parasite growth; however, only in the cell culture inoculated with brain tissue was observed parasites. According with genetic characterization, two atypical genotypes were observed: one matched with ToxoDB genotype #146 and the other one has not been described before (classified now as ToxoDB #279) (Table 2).

In our study, we had an isolation rate of 14.3%, with two isolates obtained. Previous studies with free-range chickens form Brazil have described an isolation ranging from 15.12 to 87.87% [20–22]. In the present study, isolates was obtained from seropositive animals; however, previous studies have described isolation of the parasite from seronegative animals, including chickens and doves [23, 24]. In the past, *T. gondii* was considered to be parasite with low degree of genetic diversity [5] with three clonal lineages; however, studies performed later, mainly in South American countries, a higher genetic diversity was observed, with occurrence of many atypical genotypes [6]. The isolates obtained in our study were considered of low to moderate pathogenicity, since no animal showed clinical signs of acute infection; however, since in the mouse bioassay the infectious dose is unknown, additional studies aiming to determine the virulence patterns of these isolates should be performed [25].

Genotype #146 has already been described only in chickens, cattle egret (*Bubulcus ibis*) and feral cats (*Felis catus*) [1, 26, 27], all of them being obtained of animals from Fernando de Noronha Island, in Brazil. Since Viçosa and Fernando de Noronha Island are apart by more than 800 km, we can observe that *T. gondii* genotypes are spread over the country. Besides the isolate #146, we detected a new genotype, now classified as #279, showing a high genetic diversity of *T. gondii* in Brazil, as previously observed in different isolates from Brazilian animals [23, 28]. The phylogenetic tree (Fig. 1 shows that the genotypes observed in the present study are not related with any clonal strains, being more related with Brazilian typical strains BrII and BrIII, which have been isolated in different animals in Brazil; these strains have also shown similar patterns of pathogenicity with our isolates, since BrII, and BrIII are considered as intermediate and non-virulent strains, respectively [29].

| Strain ID | Genetic markers | References |
|-----------|-----------------|------------|
| GT1       | I               | #10 [30]   |
| PTG       | II or III       | #1 [30]    |
| CTG       | II or III       | #2 [30]    |
| MAS       | u-l             | #17 [30]   |
| TgCgCal   | I               | #66 [31]   |
| TgCtBr5   | III             | #19 [29]   |
| TgCtBr64  | I               | #111 [29]  |
| TgRsCrl   | u-l             | #52 [32]   |
| Present study |              |            |
| TgCkAL01  | I               | #146       |
| TgCkAL02  | I               | New (#279) |

Table 2 PCR–RFLP genotypic profiles from *Toxoplasma gondii* isolates obtained from free-range chickens (*Gallus gallus domesticus*) intended for human consumption in Brazil
Conclusion

The presence of anti-\textit{T. gondii} antibodies and the presence of viable parasite in chicken tissues indicate the existence of environmental contamination in the region and demonstrate the circulation of new genotypes of the parasite. Further studies evaluating the pathogenic patterns of these isolates should be done to improve the understanding of the epidemiology and potential association with human toxoplasmosis in this region.

Acknowledgements

We would like to thank the Conselho Nacional de Desenvolvimento Científico e Tecnológico (CNPq) for financial support.

Compliance with Ethical Standards

Conflict of Interest On behalf of all authors, the corresponding author states that there is no conflict of interest.

Ethical Approval All procedures performed in studies involving animals were in accordance with ethical standards of the Animal Ethics Committee at which the studies were conducted (protocol number 43/2015).

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