**Toxoplasma gondii**: isolation, biological and molecular characterisation of samples from free-range *Gallus gallus domesticus* from countryside Southeast Brazil

**Resumo**

Toxoplasma gondii apresenta alta prevalência mundial, capaz de infectar diversos animais. Felinos são considerados os hospedeiros definitivos e entre os hospedeiros intermediários destacamos os mamíferos e as aves. O homem pode se infectar ingerindo cistos teciduais presentes na carne das aves e mamíferos. O conhecimento dos aspectos biológicos e moleculares do parasito possibilitam melhor entendimento da epidemiologia da toxoplasmose. Neste trabalho foi realizada triagem sorológica por hemaglutinação indireta (HI) em 58 galinhas caipiras (*Gallus gallus domesticus*) utilizadas para consumo humano, provenientes do estado do Espírito Santo, Brasil. Treze galinhas apresentaram sorologia positiva para *T. gondii*. O coração e o cérebro de cinco galinhas positivas foram colhidos, tratados com pepino e inoculados separadamente, em dois camundongos Swiss, por via intraperitoneal. Observou-se taquizoítos no peritônio entre 7 e 10 dias após o inóculo. Foram obtidos 10 novos isolados de *T. gondii* os quais foram estudados em camundongos BALB/C inoculados com 10^1 a 10^4 taquizoítos por animal. Todos os isolados foram considerados virulentos ou de virulência intermediária. A caracterização molecular dos isolados, realizada por PCR-RFLP, demonstrou a
ocorrência de três genótipos distintos. Nenhum isolado apresentou genótipo clonal ou linhagem clonal do Brasil. Não foi observada diferença molecular (PCR-RFLP) entre os isolados obtidos a partir do cérebro ou do coração da mesma ave. Dois isolados já haviam sido relatados na literatura como causadores de doenças em humanos.

**Palavras-chave:** Toxoplasma gondii, isolamento, caracterização biológica, genotipagem.

**Introduction**

*Toxoplasma gondii* is a pathogenic agent with significant impacts in human and veterinary medicine. According to estimates, one-third of the human population is infected with the parasite (WEISS & DUBEY, 2009). Felids are the only known definitive hosts of *T. gondii*, which means that the sexual cycle of the parasite occurs in only these animals. Among the most important intermediate hosts are mammals, including humans, and birds. Free-range chickens (*Gallus gallus domesticus*) habitually scratch the soil in search of food and are thus prone to acquire infection in an environment potentially contaminated with oocysts, one of the infective forms of *T. gondii*. For this reason, this bird species is considered an excellent indicator of environmental contamination (DUBEY et al., 2007b).

*Toxoplasma gondii* has a highly clonal population structure in the Northern Hemisphere (DARDE, 2008), whereas nonclonal strains, *i.e.*, atypical genotypes, are predominant in South America (SILVA et al., 2014; SHWAB et al., 2014). Thus, the genetic diversity of the parasite typically follows a geographical distribution. While few genotypes dominate in the Northern Hemisphere, hundreds of genotypes coexist in the Southern Hemisphere. Despite none being notably dominant over others, some genotypes have a higher relative frequency (SHWAB et al., 2014). In Brazil, more atypical strains exist (DUBEY & SU, 2009; CARNEIRO et al., 2013; SILVA et al., 2014), a situation related to several factors, such as geographical range, tropical climate, rich fauna and diverse transmission routes (FERREIRA et al., 2006). Furthermore, four genotypes with wide circulation and described in different hosts in Brazil have been proposed as Brazilian clonal lineages. These lineages are termed BRI, BRII, BRIII and BRIV and are distinct from the archetypal lineages described in different hosts in Brazil have been proposed as Brazilian clonal lineages. These lineages are termed BRI, BRII, BRIII and BRIV and are distinct from the archetypal lineages.

One classic means of categorising *T. gondii* virulence is by inoculating the parasite in laboratory mice. According to mortality in mice, *T. gondii* is classified as virulent, intermittently virulent or avirulent. Clonal strains of type I are virulent, independent of the inoculated dose, whereas the avirulent strains (type III) establish a chronic infection at doses below $10^3$ tachyzoites (SILVA et al., 2014).

Thus, the objective of the present study was to isolate *T. gondii* from the tissues of free-range chickens from Espírito Santo State (ES), Southeast Brazil, and to follow up by assessing the virulence and genotyping from the isolates. The knowledge of the biological and molecular aspects of the parasite is important for the epidemiology of toxoplasmosis disease.

**Materials and Methods**

A total of 58 chickens (*Gallus gallus domesticus*) grown for human consumption in Espírito Santo State from the municipalities of Cariacica (*n* = 8), Viana (*n* = 16) and Venda Nova do Imigrante (*n* = 34), ES, Brazil, were evaluated in the period of January to October 2014. Chickens sera were obtained from whole blood samples, collected in tubes without anticoagulant. First, each sample was submitted to indirect haemagglutination assays (IHAs) (Imuno-HAI Toxoplasmosse WAMA®, São Carlos, São Paulo, Brazil) according to the manufacturer’s instructions. Seropositive chickens were sacrificed by cervical dislocation, and the brains and hearts were collected. Organs were stored at 4 °C until used. *Toxoplasma gondii* strains were isolated in a bioassay of the brains and hearts of five seropositive chickens. Isolation was not successful for the eight remaining chickens. Slaughtering was made by the chicken’s producer for their own consumption. After that, heart and brain were removed, packed in individual plastic bags, and transported on ice to the Laboratory of Toxoplasmosis, located at the Instituto de Ciências Biológicas of the Universidade Federal de Minas Gerais (ICB - UFMG). Full organs were individually macerated and digested with pepsin solution (pepsin, 2.6 g; NaCl, 5.0 g; HCl, 7.0 mL; and water to make 500 mL). The homogenate was incubated at 37 °C in a shaking water bath for 1 h, washed twice by centrifugation (1200×g for 10 min) with sterile phosphate-buffered saline (PBS, pH 7.2). The final sediment was suspended with 2 mL of PBS, pH 7.2 containing 1000 units of penicillin and 100 μg of streptomycin per milliliter and inoculated intraperitoneally into two Swiss mice (DUBEY, 1998; CARNEIRO et al., 2013). Toxoplasmosis in mice was confirmed when tachyzoites were found in the lungs, or cysts in the brain by optical microscopy.

*T. gondii* virulence in mice was determined in accordance with Ferreira et al. (2001). Four groups of five mice (twenty female BALB/c mice infected with each *T. gondii* isolate) were infected intraperitoneally with increasing doses of $10^1$, $10^2$, $10^3$ and $10^4$ tachyzoites of each isolate in a volume of 0.2 mL. To obtain tachyzoites, Swiss mice were infected intraperitoneally with 300-500 brain cysts with each *T. gondii* isolate. Seven days after inoculation, the parasites were washed and collected from the peritoneal cavity with PBS pH 7.2. The exsudate were centrifuged and tachyzoites filtered in 3 μm polycarbonate membrane. The number of tachyzoites was adjusted and used for virulence assay (fresh tachyzoites) or stored as frozen pellets at −20 °C, until DNA extraction for genotyping study. Five control animals were inoculated with 0.2 mL of sterile PBS pH 7.2. Mice mortality was daily observed throughout 30 days, when the survivors were tested for anti-*T. gondii* IgG antibodies by enzyme-linked
immunosorbent assay (ELISA) and brain cysts, as described by Ferreira et al. (2001).

Toxoplasma gondii isolates were genotyped by means of polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) analysis using primer pairs for 12 different loci: SAG1, SAG2-5+5', SAG2-new, SAG3, BTUB, GRA6, c22-8, c22-9, CS3, L358, PK1 and Apico (SU et al., 2010). The RH (clonal type I), ME49 (clonal type II) and VEG (clonal type III) strains were used as controls (FUX et al., 2003). The profiles of the bands obtained after digestion with specific restriction endonucleases (SU et al., 2010) were analysed in polyacrylamide gels stained with silver nitrate and then compared with the profiles of reference strains. The obtained results were analysed in the virtual database ToxoDB (www.toxodb.org) and compared with the genotypes of the deposited strains (SU et al., 2010).

**Ethics**

The present study has been approved by the Animal Ethics Committee of the Federal University of Espírito Santo (Comitê de Ética em Experimentação Animal, Universidade Federal do Espírito Santo – CEUA/UFES 080/2011).

**Results**

**Serology**

All chicken from the municipalities of Cariacica and Viana exhibited negative IHA results. Of the 34 chickens from Venda Nova do Imigrante, 13 (38%) tested positive for anti- *T. gondii* IgG.

**Isolation and characterisation of *T. gondii* strains**

The isolation of *T. gondii* was successful; all inoculated Swiss mice died and/or developed ascites with tachyzoites at 7 to 15 d post-inoculation. Five isolates were obtained from the brains (TgCkBrEs1b to TgCkBrEs5b) and five from the hearts (TgCkBrEs1h to TgCkBrEs5h). Among the isolates obtained in this study, none were considered avirulent, and only two of the heart samples (TgCkBrEs1h and TgCkBrEs5h) were considered immediately virulent (Table 1), even though survival only occurred at the lowest dose (10^1 tachyzoites).

**Genotyping of *T. gondii* isolates**

All ten isolates obtained were successfully genotyped by the 12 molecular markers described by Su et al. (2010). Three different genotypes have already been described in the literature (Table 2). The isolates TgCkBrEs1b and TgCkBrEs1h exhibited genotype #206 (ToxoDB). The TgCkBrEs2b, TgCkBrEs2h, TgCkBrEs3b and TgCkBrEs3h isolates exhibited genotype #36 (ToxoDB), and the TgCkBrEs4b, TgCkBrEs4h, TgCkBrEs5b and TgCkBrEs5h isolates exhibited genotype #6 (ToxoDB). No difference was observed in the genotypes between the isolates obtained from the brain or heart of the same bird.

**Discussion**

Of the free-range chicken farms in rural areas of Espírito Santo State, no cases of seropositive animals were found in the municipalities of Cariacica and Viana. The present study was developed in the mountainous region of the state (Venda Nova do Imigrante) for which no previous data on avian toxoplasmosis were available. In Venda Nova do Imigrante, 38% of the evaluated birds exhibited specific antibodies against *T. gondii*, our results were similar compared to results from backyard chickens in the metropolitan area of Recife (FERNANDES et al., 2016) and in the semiarid region of the state of Pernambuco (SÁ et al., 2017). Venda Nova do Imigrante also has a high prevalence of ocular toxoplasmosis, with 11.27% positive cases (ABREU et al., 1998), and is likely a favourable site for the presence and transmission of the parasite, as the mean prevalence of ocular toxoplasmosis in Brazil is 1.6% (ZANETTI & PLETSCH, 2007).

**Table 1.** Site of isolation and classification of virulence by mortality in BALB/c mice inoculated with decreasing doses of *Toxoplasma gondii* tachyzoites isolated from free-range chickens in Espírito Santo State, Brazil.

<table>
<thead>
<tr>
<th>Isolates</th>
<th>Site of isolation</th>
<th>10^1 tach.</th>
<th>10^2 tach.</th>
<th>10^3 tach.</th>
<th>10^4 tach.</th>
<th>Classification of virulence</th>
</tr>
</thead>
<tbody>
<tr>
<td>TgCkBrEs1b</td>
<td>VNI/I</td>
<td>5/5**</td>
<td>4/4</td>
<td>5/5</td>
<td>2/2</td>
<td>Virulent</td>
</tr>
<tr>
<td>TgCkBrEs2b</td>
<td>VNI/II</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>Virulent</td>
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<tr>
<td>TgCkBrEs3b</td>
<td>VNI/II</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>Virulent</td>
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<tr>
<td>TgCkBrEs4b</td>
<td>VNI/IV</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>Virulent</td>
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<tr>
<td>TgCkBrEs5b</td>
<td>VNI/IV</td>
<td>5/5</td>
<td>5/5</td>
<td>5/5</td>
<td>4/4</td>
<td>Virulent</td>
</tr>
<tr>
<td>TgCkBrEs1h</td>
<td>VNI/I</td>
<td>5/5</td>
<td>5/5</td>
<td>0/2</td>
<td></td>
<td>Intermediate Virulent</td>
</tr>
<tr>
<td>TgCkBrEs2h</td>
<td>VNI/II</td>
<td>5/5</td>
<td>5/5</td>
<td>2/2</td>
<td>2/2</td>
<td>Virulent</td>
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<tr>
<td>TgCkBrEs3h</td>
<td>VNI/II</td>
<td>5/5</td>
<td>5/5</td>
<td>3/3</td>
<td></td>
<td>Virulent</td>
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<tr>
<td>TgCkBrEs4h</td>
<td>VNI/IV</td>
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<td>5/5</td>
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<td>Virulent</td>
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<tr>
<td>TgCkBrEs5h</td>
<td>VNI/IV</td>
<td>5/5</td>
<td>5/5</td>
<td>1/3</td>
<td></td>
<td>Intermediate Virulent</td>
</tr>
</tbody>
</table>

* TgCkBrEs1b: Tg = *Toxoplasma gondii*; Ck = chicken; Br = Brazil; Es = Espírito Santo State; b = brain; h = heart; VNI/I = Venda Nova do Imigrante, holding I; VNI/II = Venda Nova do Imigrante, holding II; VNI/IV = Venda Nova do Imigrante, holding IV; **Number of BALB/c mice that died/total number of mice inoculated and confirmedly infected (by means of parasitological diagnosis and/or ELISA).
Table 2. Multilocus genotyping of *Toxoplasma gondii* isolates from free-range chickens slaughtered in Espírito Santo State, Brazil, by polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) analysis.

<table>
<thead>
<tr>
<th>Isolates</th>
<th>Virulence</th>
<th>Genetic markers</th>
<th>CS3</th>
<th>ToxoDB Genotype</th>
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<tr>
<td></td>
<td></td>
<td>SAG1 SAG2 alt SAG2 SAG3 BTUB GRA6 c22-8 c29-2 L358 PK1 Apico</td>
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<tr>
<td>RH88</td>
<td>Virulent</td>
<td>I I I I I I I I I I</td>
<td></td>
<td>#10</td>
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<tr>
<td>ME49</td>
<td>Intermediate</td>
<td>II/III II II II II II II II II II</td>
<td></td>
<td></td>
</tr>
<tr>
<td>VEG</td>
<td>Avirulent</td>
<td>II/III III III III III III III III III</td>
<td></td>
<td></td>
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<tr>
<td>TgCkBrs1b</td>
<td>Virulent</td>
<td>u-1 II III III II III I III I I I</td>
<td></td>
<td>#206</td>
</tr>
<tr>
<td>TgCkBrs1h</td>
<td>Intermediate</td>
<td>u-1 II III III II III I III I I II</td>
<td></td>
<td>#206</td>
</tr>
<tr>
<td>TgCkBrs2b</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
<td></td>
<td>#24</td>
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<tr>
<td>TgCkBrs2h</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
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<td>#24</td>
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<tr>
<td>TgCkBrs3b</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
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<tr>
<td>TgCkBrs3h</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
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<tr>
<td>TgCkBrs4b</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
<td></td>
<td>#6</td>
</tr>
<tr>
<td>TgCkBrs4h</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
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<tr>
<td>TgCkBrs5b</td>
<td>Virulent</td>
<td>I I I III I I I I I I I I I I I</td>
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<tr>
<td>TgCkBrs5h</td>
<td>Intermediate</td>
<td>I I I III I I I I I I I I</td>
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The IHA method was chosen for the detection of specific antibodies against *T. gondii* because it exhibits good agreement with the modified agglutination test (MAT) (82%) (BELTRAUME et al., 2012). IHA is considered a reliable test, in addition to the advantages of low cost and easy application compared to other methods (CASARTELLI-ALVES et al., 2014).

*T. gondii* was isolated from all chickens submitted to the bioassay. According to Dubey et al. (1998), *T. gondii* cysts are more abundant in neural, muscle and cardiac tissue. Accordingly, the heart of the chickens was chosen for isolation due to the heavy parasitism and because it is considered a culinary delicacy in several regions of Brazil. To ensure higher chances of isolation, the brains of the animals were also used. Aigner et al. (2010) have shown by means of real-time PCR that no significant difference exists in the number of parasites per gram of cardiac and neural tissue in seropositive birds.

According to Dubey et al. (2005), the success of isolation depends on the number of inoculated animals, the amount of tissue used and the concentration of the parasite in the tissues. The success of isolation of the parasites in the present work corroborates these data. The frequency of positivity for isolation of *T. gondii* from chickens typically ranges from 28.4% (OLIVEIRA et al., 2009) to 70% (DUBEY et al., 2002). Cases with a 100% success rate of isolation, as in the present work, are rare.

Depending on its lethality in mice, *T. gondii* can be considered virulent, avirulent or intermittently virulent. Type I clonal strains, such as the RH strain, are always virulent, independent of the dose. Mice die even with an inoculum of a single tachyzoite. On the other hand, type II clonal strains establish a chronic infection by producing tissue cysts, even at doses as high as 10⁷ tachyzoites. Type II clonal strains exhibit intermediate virulence (FERREIRA et al., 2001). All isolates from the present work were virulent, except for TgCkBrs1h and TgCkBrs5h, which were considered intermittently virulent. These results agree with the observations of Dubey et al. (2003) and Beltrame et al. (2012), who have shown that mouse-virulent *T. gondii* strains circulate in asymptomatic vertebrate hosts in Brazil.

According to Gilbert et al. (2008), a predominance of virulent *T. gondii* strains exist in Brazil compared to Europe, where clonal genotypes are predominant. That finding agrees with the results found in the present work. The authors also reported that the severity of cases of ocular toxoplasmosis is higher in Brazil than in Europe, which suggests that the observations of the experimental model could be applied to humans.

The isolates obtained from the different organs (heart and brain) within the same chicken did not differ with respect to molecular characterisation. Slightly different phenotypic differences were found, i.e., earlier death in animals inoculated with chicken brain (data not shown) and all isolates from the brain being classified as virulent. Thus, due to phenotypic similarity and genetic identity, we can simplify the nomenclature of the isolates and describe them as TgCkBrs1, TgCkBrs2, TgCkBrs3, TgCkBrs4 and TgCkBrs5. For a deeper understanding of the isolates, Dubey et al. (2008a) have suggested that cases where phenotypic characterisation is different from the molecular characterisation be submitted to DNA sequencing to determine whether the isolates are, in fact, identical or to spot where the differences occur. Another possibility is to assess the diversity of the ROP5, ROP16, ROP17 and ROP18 alleles with respect to virulence, as suggested by Shwab et al. (2016).

Despite the possibility of co-infection by two different *T. gondii* strains in the same host, as reported by Dubey et al. (2007a), such co-infection was not observed in the present study. The genotypes isolated from different chickens within the same municipality exhibited different genotypes. This result is similar to showed in St. Kitts, West Indies, that reveal a greater genetic diversity of strains circulating on the island (HAMILTON et al., 2017). The isolate TgCkBrs1 exhibited the same genotype as TgCkBrsRj3 (according to the ToxoDB), which was obtained from a free-range chicken from the municipality of Rio Bonito, Rio de Janeiro State, Brazil. This same genotype (#206 ToxoDB) has also been described in a
free-range chicken from the municipality of Colatina in Espírito Santo State (PENA et al., 2013) and in an isolate from the blood of a new-born with congenital toxoplasmosis in Minas Gerais State (CARNEIRO et al., 2013). Considering the distance between the cited regions, this occurrence provides evidence of a common strain circulating throughout the Southeast Region of Brazil. The TgCkBrs2 and TgCkBrs3 isolates are similar, and their genotype (#6 ToxoDB) has already been described in G. gallus in Rio de Janeiro State (DUBEY et al., 2008b).

Two further isolates described in the present study, i.e., TgCkBrs4 and TgCkBrs5, are considered similar genotypes (genotype #6 ToxoDB). This genotype, also known as Brl, was previously identified in chickens, cats and dogs in Brazil (PENA et al., 2008). Ferreira et al. (2011) have described an isolate with a genotype like that of the two isolates mentioned above, which were obtained from a 45-year-old HIV-positive patient with neurotoxoplasmosis characterised by diffuse encephalitis. Even when not clear whether a given T. gondii isolate exhibits the same virulent or avirulent behaviour in humans as in mice (DARDÉ, 2008; BOOTHROYD & GRIGG, 2002), the discovery of these isolates in humans deserves attention, as these are atypical strains and can generate new mechanisms of pathogenicity. Since undercooked heart from chicken is a delicacy enjoyed both in Espírito Santo and other Brazilian states, it is probable that these animals may be carriers of highly pathogenic strains for humans in the region.

As for the CS3 marker, all five isolates exhibited allele I or II, and with respect to virulence, none were considered avirulent, thus suggesting a relation between the presence of allele I and II of the CS3 marker and virulence in mice, as described previously (KHAN et al., 2005; PENA et al., 2008; SILVA et al., 2014).

Conclusion

Free-range chickens were successfully used as sentinels of environmental infection by T. gondii. The animals exhibited high tissue parasitism, which facilitates isolation via bioassays in mice. Genotypes were found that have been described previously from human infections in the same region of Brazil (Southeast), and their biological characterisation revealed that these strains were virulent or intermediately virulent. No preference of the different strains for different tissues was found in the chickens.

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References


