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Evaluation of biological behavior of *Toxoplasma gondii* atypical isolates # 14 and # 163

Mariana Bernstein, Lais L. Pardini, Lucía M. Campero, M. Elisa Helman, Juan M. Unzaga, María C. Venturini, Gastón Moré

PII: S0014-4894(19)30321-2

DOI: https://doi.org/10.1016/j.exppara.2020.107860

Reference: YEXPR 107860

To appear in: Experimental Parasitology

Received Date: 19 July 2019

Revised Date: 11 February 2020

Accepted Date: 12 February 2020

Please cite this article as: Bernstein, M., Pardini, L.L., Campero, Lucí.M., Helman, M.E., Unzaga, J.M., Venturini, Marí.C., Moré, Gastó., Evaluation of biological behavior of *Toxoplasma gondii* atypical isolates # 14 and # 163, *Experimental Parasitology* (2020), doi: https://doi.org/10.1016/j.exppara.2020.107860.

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Credit author statement for "Evaluation of biological behavior of *Toxoplasma gondii* atypical isolates # 14 and # 163"

Mariana Bernstein: Investigation; Methodology; Writing - Original Draft; Writing - Review & Editing

Lais L. Pardini: Investigation; Validation; Writing - Review & Editing

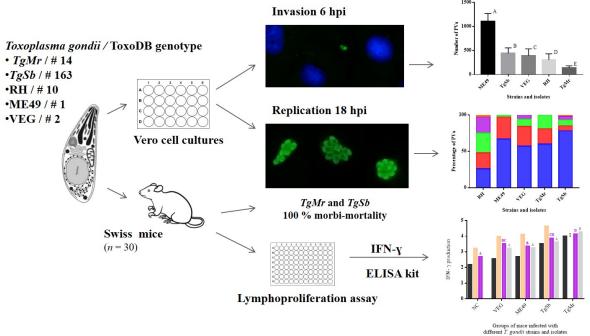
Lucía M. Campero: Investigation

M. Elisa Helman: Formal analysis

Juan M. Unzaga: Funding acquisition

María C. Venturini: Funding acquisition; Supervision

Gastón Moré: Conceptualization; Supervision; Writing - Review & Editing



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1	Evaluation of biological behavior of <i>Toxoplasma gondii</i> atypical isolates # 14 and # 163
2	Mariana Bernstein <sup>1, 2*</sup> , Lais L. Pardini <sup>1, 2</sup> , Lucía M. Campero <sup>1, 2</sup> , M. Elisa Helman <sup>1, 2</sup> , Juan
3	M. Unzaga <sup>1</sup> , María C. Venturini <sup>1</sup> , Gastón Moré <sup>1, 2</sup>
4	
5	<sup>1</sup> Laboratorio de Inmunoparasitología (LAINPA), FCV-UNLP, La Plata, Bs. As., Argentina
6	<sup>2</sup> Consejo Nacional de Investigaciones Científicas y Técnicas, Buenos Aires, Argentina
7	*Corresponding author:
8	Mariana Bernstein, Laboratorio de Inmunoparasitología, Universidad Nacional de La Plata,
9	Facultad de Ciencias Veterinarias, calle 60 y 118 (1900) La Plata, Argentina.
10	E-mail address: mbernstein@fcv.unlp.edu.ar
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## 23 ABSTRACT:

Toxoplasma gondii is an obligate intracellular protozoan parasite capable of infecting 24 warm-blooded animals, including humans. A highly diverse genetic population has been 25 reported in Central and South America, predominating mainly atypical genotypes. Different 26 genotypes showed different biological behavior in mice. The aim of this study was to 27 evaluate the biological behavior of T. gondii isolates obtained from Macropus rufogriseus 28 (TgMr) and Saimiri boliviensis (TgSb) identified as atypical genotypes # 14 and # 163, 29 respectively. Strains RH, ME49 and VEG were used as reference for clonal types I, II and 30 III, respectively. In vitro invasion and replication capacity assays were analyzed at 6 and 18 31 hpi, respectively. In vivo assay was performed in Swiss mice (n = 30) using  $1 \times 10^2$  and 32  $1 \times 10^3$  parasites/mouse as infective doses (ME49, VEG, TgMr, TgSb and negative control). 33 Morbi-mortality and tissues PCR were assessed. Lymphoproliferation assays were 34 performed and gamma interferon was measured by ELISA. The ME49 strain showed the 35 highest invasion, followed by TgSb and VEG, while RH and TgMr presented the lowest 36 invasions. The RH strain and the TgSb isolate showed more endodyogeny events (fastest 37 doubling times) than VEG and ME49 strains and the TgMr isolate. Both atypical isolates 38 39 showed high virulence (100 % morbi-mortality, at 8-10 dpi) and parasite DNA was detected in all tissue samples. Splenocytes from mice inoculated with TgMr and TgSb 40 registered the highest values of gamma interferon. An in vitro invasion-replication index 41 was established which correlates inversely with virulence in mice. In conclusion, T. gondii 42 atypical isolates # 14 and # 163 showed a different in vitro behavior than clonal strains, 43 with low invasion-replication indexes but being highly virulent in mouse model. 44

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### 49 1. INTRODUCTION

Toxoplasma gondii is an obligate intracellular protozoan parasite capable of infecting 50 warm-blooded animals, including humans (Dubey, 2010). Felines are the definitive hosts 51 and several species of mammals and birds act as intermediate hosts presenting different 52 clinical signs (Dubey, 2010). Toxoplasmosis in humans is generally asymptomatic, 53 however, it may cause a variety of neurological signs and multi-organic failure in 54 immunocompromised patients and fetal lesions in the retina (ocular toxoplasmosis), central 55 nervous system injuries and even abortion in primarily infected pregnant women (Weiss 56 and Kim, 2014). In the northern hemisphere (Europe, USA and Canada) a predominance of 57 clonal population of T. gondii has been reported, with clonal type II as dominant (Weiss 58 and Kim, 2014). However, a highly diverse population has been reported in Central and 59 60 South America (especially Colombia, Brazil and Argentina), being detected non-clonal or atypical genotypes with combinations of types I, II and III alleles, or even new alleles 61 (Bernstein et al., 2018; Pena et al., 2008; Weiss and Kim, 2014). Different genotypes 62 showed different biological behavior analyzed mainly in mice model (Dubremetz and 63 Lebrun, 2012). 64

*Toxoplasma gondii* survival relies on its capacity of invasion of new host cells (Weiss and Kim, 2014). The protozoan multiplies asexually in virtually all nucleated host cells producing tachyzoites (fast division) or bradyzoites (slow division). The tachyzoites enter the host cell by an active invasion mechanism, releasing enzymes which produce the invagination of the host cell membrane. This invagination surrounds the tachyzoites and forms the parasitophorous vacuole (PV) (Weiss and Kim, 2014). After the formation of the

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PV, the tachyzoites multiply by endodyogeny, in which two daughter cells are formed 71 inside the mother cell. The divisions continue resulting in a geometric expansion of the 72 parasites until the host cell is destroyed (Dubey et al., 1998). The first in vitro experiments 73 showed that the tachyzoites of the RH strain (clonal type I) have the highest multiplication 74 rate while the tachyzoites of the ME49 strain (clonal type II) have the highest invasion rate 75 (Cañedo-Solares et al., 2013; Contreras-Ochoa et al., 2012; Saadatnia et al., 2010). Some 76 authors suggest that strain virulence depends on replication capacity more than on invasion 77 78 capacity (Cañedo-Solares et al., 2013; Dubremetz and Lebrun, 2012). Similar studies have 79 been carried out with the protozoan Neospora caninum, closely related to T. gondii (Dellarupe et al., 2014b). It has been shown that in vitro invasion and replication capacity 80 can be used as virulence traits in N. caninum and correlated with virulence in mice model 81 (Dellarupe et al., 2014a, b). 82

The *T. gondii* invasion and replication experiments are very heterogeneous and most of them use clonal isolates (Alomar *et al.*, 2013; Cañedo-Solares *et al.*, 2013; Contreras-Ochoa *et al.*, 2012; Cuellar *et al.*, 2012; Malkwitz *et al.*, 2018; Saadatnia *et al.*, 2010). Therefore, a standard protocol is needed to make results comparable (Contreras-Ochoa *et al.*, 2012).

*Toxoplasma gondii* virulence has been defined as the mortality rate in different mice models (Dubremetz and Lebrun, 2012; Saraf *et al.*, 2017; Sibley and Boothroyd, 1992). The clonal type I (ToxoDB # 10) is highly virulent in mice, with a  $LD_{100} = 1$  parasite (Sibley and Boothroyd, 1992). In contrast, clonal type II is considered as intermediate virulence with a  $LD_{50} \ge 10^3$  parasites (ToxoDB # 1), and clonal type III (ToxoDB # 2) is considered non-virulent with a  $LD_{50} \ge 10^5$  parasites (Sibley and Boothroyd, 1992). Until now, non-clonal genotypes have shown different virulence in mouse model, leading to

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difficulties on predict virulence based only on genotypes (Dubey, 2010; Dubremetz and 95 Lebrun, 2012; Weiss and Kim, 2014). The morbidity rates, different degrees of 96 histopathological lesions, the presence of T. gondii DNA and tissue cysts are associated 97 with virulence in mouse models (Costa et al., 2018; Pereira et al., 2017; Pinheiro et al., 98 2015). Immune response in experimentally infected mice is characterized by the production 99 of IFN- $\gamma$  (gamma interferon), which induces the development of Th1 profile and 100 macrophages activation helping to control the parasite proliferation, however paradoxically, 101 102 its overproduction has been associated with acute virulence (Chen et al., 2016; Mordue et 103 al., 2001; Rodgers et al., 2005; Wang et al., 2016).

*Toxoplama gondii* genotype # 163 has been frequently detected in Brazil and considered of
variable virulence in mice model (Gennari *et al.*, 2015; Silva *et al.*, 2014). Genotype # 14
has been reported in several South American countries (Argentina, Brazil, Colombia, Chile
and Venezuela) and the USA showing a wider distribution than genotype # 163 (Pardini *et al.*, 2019). In Brazil, this genotype generated 100 % mortality in mice model, indicating a
highly virulent phenotype (Pena *et al.*, 2008; Rajendran *et al.*, 2012).

The biological behavior of non-clonal *T. gondii* genotypes *in vitro* model (expressed as invasion and replication) and its relation with the virulence in mouse model (expressed as mortality and morbidity) are still not clear (Dubey, 2010; Dubremetz and Lebrun, 2012; Weiss and Kim, 2014). It is necessary the establishment of accurate and reliable *in vitro* protocols which allow the parasites behavior evaluation and help to reduce, replace and refine the number of animals for *in vivo* assays.

116 The aim of this study was to evaluate the biological behavior of *T. gondii* isolates identified

as atypical genotypes # 14 and # 163 by *in vitro* and *in vivo* studies. Additionally, this study

- aimed to establish a relation between the *in vitro* and *in vivo* assays results.
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## 20 2. MATERIALS AND METHODS

Toxoplasma gondii isolates were obtained from fatal cases of toxoplasmosis in zoo animals 121 Saimiri boliviensis (Bolivian squirrel monkey) and Macropus rufogriseus (Bennett's 122 wallaby) in Argentina (Basso et al., 2007; Pardini et al., 2015), named in this study as TgSb 123 (genotype # 163) and TgMr (genotype # 14), respectively (Bernstein et al., 2018). 124 Tachyzoites were maintained in liquid nitrogen and re-isolated in mice deficient for IFN- $\gamma$ 125 (GKO; C.129S7 (B6) -ifng<tm1Ts> / J, The Jackson Laboratory, UK) to reduce number of 126 passages and adaptation to cell culture (Contreras-Ochoa et al., 2012; Dellarupe et al., 127 2014b). Strains RH, ME49 and VEG were used as reference for clonal types I, II and III, 128 129 respectively.

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## 2.1 In-vitro invasion and replication assay

This assay was designed based on previous studies from Alomar *et al.*, 2013 and Dellarupe *et al.*, 2014b.

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## 2.1.1 VERO cell cultures

VERO cells were cultivated with RPMI-1640 medium with glutamine (Gibco, USA),
supplemented with 10 % fetal bovine serum (FBS) (Natocor, Argentina) and 1 % of a
solution of antibiotics and antimycotic (Gibco BRL, UK). All cultures were incubated at 37
°C with 5 % CO<sub>2</sub>.

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## 2.1.2 Preparation of tachyzoites

Tachyzoites were harvested when cell monolayer was 80 % infected and the parasites were inside the PVs. Cell monolayer was detached with a cell scraper (GBO, Germany), and the suspension was passed through the 22G, 25G and 27½ G needles, to release the parasites

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- 142 inside the PVs. Tachyzoites were resuspended in the required infection dose to inoculate
- the VERO cultures. All infections were made within 1 h post released of the parasites.
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## 2.1.3 In vitro assay in 24-well plates

The invasion and the amount of produced tachyzoites (replication) were determined in a 145 24-well plate assay. Circular glass coverslips of 15 mm (MATSUNAMI, Micro Cover 146 Glass, USA) were placed in 24-well plates, and a concentration of 1x10<sup>5</sup> VERO cells *per* 147 well was added with the conditions described in section 2.1.1. The cell cultures were left 148 overnight (16 to 18 h), to obtain an 80 % confluent monolayer, and the tachyzoites were 149 prepared as described in section 2.1.2 and seeded in an infection dose of  $1 \times 10^{5}$ /well. Plates 150 were placed on ice for 10 min to synchronize cell invasion (Alomar et al., 2013). One hour 151 after infection, the culture medium was changed. Plates were washed with sterile phosphate 152 buffer solution (PBS) and fixed with cold methanol for 5 min at -20 °C at 6 and 18 hours 153 post infection (hpi). Fixed cultures were preserved with sterile PBS at 4 °C. Three 154 independent experiments were performed for each cut-off time, with 3 replicates each one. 155 Tachyzoites identification was performed by indirect fluorescent antibody test (IFAT). The 156 cultures were permeabilized with 0.25 % Triton X100 (Promega, USA) in PBS for 10 min. 157 Then, PBS with 5 % FBS was added as blocking solution for 30 min. As a primary 158

antibody a serum from a *T. gondii* naturally infected goat (with IFAT titer higher than 800),
diluted 1:100 in PBS-0.5 % Tween20 (Metraquímica s.r.l., Argentina) with 0.5 % bovine
serum albumin (BSA, Sigma, USA) was used (Gos et al ., 2017). As secondary antibody
anti-goat IgG conjugated with Alexa 488 fluorochrome (Life technologies, USA) was
diluted 1: 500 in PBS-0.5 % Tween20 and 0.5 % BSA and incubated in the dark. As nuclei
staining, DAPI (4', 6-Diamidino-2-Phenylindole, Dihydrochloride, Molecular Probes,
Invitrogen, USA) was used diluted 1: 2000 in PBS, incubated for 5 min at room

temperature. All antibodies incubations took place in humid chamber at 37 °C for 45 min 166 and 3 washes were made after incubations with sterile PBS. Finally, cultures were mounted 167 with a drop of ~10 µl of mounting fluid MOWIOL 4-88 (Sigma Aldrich, USA) inverting 168 each coverslip over the mounting fluid, air dried at room temperature overnight and stored 169 at -20 °C until microscope examination. 170

Fifty microscopic fields (40X objective) were randomly selected from each coverslip and 171 the PVs were counted by discriminating among PVs with 1, 2, 4, 8, and 16 tachyzoites. The 172 173 invasion was analyzed by the total number of PVs produced by each strain and isolate at 6 174 hpi. The replication was analyzed by calculating the total amount of parasites that showed at least one event of endodiogeny (PVs with 2 or more tachyzoites) at 18 hpi for each strain 175 and isolate. 176

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## 2.2 Mice bioassay

The experimental design was evaluated and approved by the CICUAL (Institutional 178 Committee for the Care and Use of Laboratory Animals), Faculty of Veterinary Sciences, 179 UNLP (protocol 42.6.14T). According their recommendations and to reduce the number of 180 animals, the RH strain was excluded from the mice bioassay due to its known  $LD_{100} = 1$ 181 parasite (Sibley and Boothroyd, 1992). The infective doses of isolates TgMr and TgSb 182 were defined by a previous trial were all inoculated mice with  $1 \times 10^3$  and  $1 \times 10^4$  tachyzoites 183 184 died with acute toxoplasmosis. A total of 30 specific pathogens free female Swiss mice, of 8 weeks age and 23-29 gr average weight, were used for the assay. Water and food were 185 provided ad libitum and animals were handled to minimize stress conditions throughout the 186 experiment. 187

Five groups (6 mice/group), were subcutaneously inoculated with the following infection 188 doses: 1x10<sup>2</sup> tachyzoites (3 mice/group) and 1x10<sup>3</sup> tachyzoites (3 mice/group) in each 189

190 group for TgSb (# 163), TgMr (# 14), ME49 and VEG, and with 0.5 ml PBS for the 191 negative control group (NC). The parasites were maintained in VERO cell cultures as 192 explained in sections 2.1.1. and 2.1.2.

193 The mice were monitored daily for 4 weeks, if compatible toxoplasmosis signs were 194 observed were sacrificed according to the requirements established by the CICUAL. 195 Autopsy was performed and samples of central nervous system (CNS) for histopathology 196 and PCR, lung for PCR and spleen for lymphoproliferation assay were obtained as detailed 197 below. Blood samples were obtained by cardiac puncture and sera were stored at -20 °C 198 until used.

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## 2.2.1 T. gondii DNA identification by PCR

Total DNA extraction was performed from CNS and lung samples with the Wizard® genomic DNA purification kit, according to the manufacturer's instructions (Promega, USA). PCR amplification was performed using TOX5-TOX8 primers as previously described (Pardini *et al.*, 2015).

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2.2.2 Histopathology

A portion of the CNS from each mouse was preserved in 10 % buffered formalin. Samples 205 were processed routinely, stained with hematoxylin and eosin (H&E) and examined 206 microscopically according to the routine protocols of LAPAVET "Dr. Bernardo Epstein ", 207 208 FCV, UNLP. The histopathological lesions were evaluated with the following scoring scheme: grade 1: mild non-suppurative meningoencephalitis, scarce focal gliosis and scarce 209 neuronal degeneration; grade 2: moderate non-suppurative meningoencephalitis, 210 perivascular cuffs; non-suppurative multifocal 211 grade 3: and necrotizing meningoencephalitis (Venturini et al., 1996). 212

213 2.2.3 Anti-T. gondii antibody detection

Mice sera were diluted in PBS from 1:25 in base 2 and IFAT was performed to final titer. Samples were considered positive until the last dilution where complete peripheral fluorescence was observed (Gos *et al.*, 2017).

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2.3 Lymphoproliferation assay

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## 2.3.1 Splenocytes cultures

Splenocytes were obtained from a third of each spleen. The organ was smear under laminar flow and purified by removing the red blood cells using 1 ml of lysis buffer solution (0.83 % ammonium chloride). Cells were counted with 0.5 % trypan blue viability stain in the Neubauer chamber and seeded onto 96-well culture plates at a concentration of 2 x  $10^5$ viable cells *per* well in 200 µl of culture medium (Wang *et al.*, 2016).

224 2.3.2 T. gondii total lysate antigen, stimulation and measure of IFN-y

The total lysate antigen (TLA) was produced from the *T. gondii* RH, ME49 and VEG reference strains and from isolates TgMr and TgSb, maintained in cell cultures as described in sections 2.1.1 and 2.1.2. The tachyzoites of each strain and isolate were centrifuged (3000 g, 10 min) and resuspended in 500 µl of sterile PBS. The suspension was sonicated in 3 cycles, 1 min / cycle, at 45W/s (Omni International, USA). The protein content of each TLA was quantified with a commercial kit (BCA, Pierce, USA) according to the manufacturer's instructions.

The following treatments were performed in triplicate for each mouse splenocytes
according previous studies (Chen *et al.*, 2016; Wang *et al.*, 2016):

A- Splenocytes seeded in medium with 10 % FBS as negative control. B- Stimulation with concanavalin A (5  $\mu$ g / ml; Biorad, USA) as positive control. C- Stimulation with TLA from the RH strain (10  $\mu$ g / ml). D- Stimulation with TLA from homologous tachyzoites (10  $\mu$ g / ml) (Rodgers *et al.*, 2005). Cultures were maintained at 37 °C with 5 % CO<sub>2</sub> for 72 h, the supernatants were collected and stored at -20 °C until use. The IFN- $\gamma$  was measured using a commercial ELISA kit (Mouse IFN- $\gamma$  ELISA Kit, Catalog n°: BMS606; Thermo Fisher Scientific Inc., USA) according supplier's instructions, on the pool of the triplicates of each treatment *per* mouse splenocytes.

#### 243 *2.4 Statistical analysis*

*Invasion and replication assay:* A Generalized Linear Mixed Model (GLMM) with log link
function was applied. Fixed factor was defined as: strains and isolates, and the response
variables were defined as: number of PVs at 6 hpi and total number of tachyzoites at and
18 hpi. The experiment effect was considered as a random parameter. When significant
differences were detected, the LSD Fisher test was performed (Balzarini *et al.*, 2008).
Percentages of PVs with 1, 2, 4, 8, and 16 tachyzoites were calculated for the different *T*. *gondii* strains and isolates at 18 hpi and were plotted as stacked column graphics.

*Mice virulence assay*: Morbi-mortality was calculated as: total of clinically ill and sacrificed animals / total of infected animals \* 100. Murine IgG anti-*T. gondii* titers among the groups were analyzed by Kruskal-Wallis test.

*Lymphoproliferation assay*: the IFN-γ production values obtained by ELISA were log10
transformed and analyzed by ANOVA and subsequent LSD Fisher.

The Pearson correlation coefficient was used to evaluate the association between the parameters evaluated in the *in vitro* assay (invasion and replication) and the morbimortality in the *in vivo* assay.

The InfoStat software version 2018 (Di Rienzo *et al.*, 2008) was used for all statistical analyzes. The degree of significance was established at p < 0.05 for all analyzes. All the 261 graphics were generated using the GraphPad Prism software version 7.00 (San Diego,262 USA).

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## **3. RESULTS**

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## 3.1 In-vitro invasion and replication assay

Homogeneity among replicates was observed: no significant differences were found among the replicates at 6 and 18 hpi (p > 0.05).

Significant differences were found in the invasion capacity between the reference strains and the atypical isolates analyzed (p < 0.0001). The ME49 strain showed the highest invasion capacity (p < 0.05), while the *TgMr* isolate presented the lowest invasion capacity of all parasites (**Figure 1A**).

272 Significant differences were observed in parasite replication, where the RH strain showed273 the highest replication capacity (Figure 1B).

Significant differences were observed in the load and number of PVs at 18 hpi (p < 0.0001) 274 according to different strains and isolates. Only the RH strain and the TgSb isolate 275 produced PVs with 16 tachyzoites (Figure 2). In summary, the RH strain and the TgSb 276 isolate showed high replication capacity (PVs of 16 tachyzoites = 4 endodyogeny events), 277 while the VEG and ME49 strains, and the TgMr isolate showed low replication capacity 278 279 (PVs of 8 tachyzoites = 3 endodyogeny events). The doubling times were on average  $\sim 4.5$ h for RH and TgSb (18 h / 4 division events) and ~ 6 h for VEG, ME49 and TgMr (18 h / 3 280 division events). 281

All strains and isolates had PVs with parasites that did not replicate or had fewer events of endodyogeny. The RH strain showed less than 30 % of PVs with 1 tachyzoite and equal percentages of PVs with 2, 4 and 8 parasites demonstrating high replication capacity as above-mentioned. Also, the *TgSb* isolate showed high replication capacity but very small percentages of PVs with 2 to 16 parasites (~ 25 % of PVs). The VEG and ME49 strains, and the *TgMr* isolate evidenced ~ 60 % of PVs without replication events (**Figure 2**).

An index between invasion and replication (invasion-replication index) was established for each strain and isolate, using the following formula: (average invasion \* 18 hpi average replication) / 1000. The resulting index values were: ME49: 790; VEG: 260; TgSb: 237; RH: 234; TgMr: 42.

*3.4 Mice bioassay* 

293 Clinical signs compatible with toxoplasmosis infection (xiphosis, hirsute hair, 294 conjunctivitis, decay and tachypnea) were observed in all the inoculated mice with the 295 isolates TgSb (sacrificed at 10 dpi) and TgMr (sacrificed at 8 dpi - inoculated with  $10^3$ 296 tachyzoites - and 9 dpi -inoculated with  $10^2$  tachyzoites -). Both atypical isolates showed 297 high virulence (100 % morbi-mortality) in mouse model. Reference strains ME49 and VEG 298 produced no morbidity, all mice were sacrificed at the end of the assay together with the 299 NC group.

No correlation was observed between *in vitro* invasion and mortality (Pearson coefficient 0.19, p > 0.05), nor between *in vitro* replication and mortality (Pearson coefficient 0.04, p > 0.05). The *in vitro* invasion-replication index was inversely related to the mortality recorded in this assay: the lower value of the *in vitro* index (isolate TgMr <strain RH < isolate TgSb) the higher mortality (higher virulence) in mice model.

Results from histopathological analysis and presence of *T. gondii* DNA in mice tissue samples are showed in **Table 1**. Out of all CNS from *T. gondii* inoculated mice (n = 24): 9 presented grade 1 lesions (5 *TgSb*; 4 ME49), 3 grade 2 (1 ME 49; 2 VEG), 2 grade 3 (2 VEG) and 10 did not present lesions. Different tissue lesions grades are shown in **Figure 3**.

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All the mice inoculated with *T. gondii* were seropositive. Antibody titers were higher in mice inoculated with ME49 (final titer 6400: *n* 4/6, and 3200, *n* 2/6) and VEG (final titer 12800, *n* 6/6), than mice inoculated with TgMr (final titer 50: *n* 4/6 and 25: *n* 2/6) and TgSb(final titer 100; *n* 4/6 and 50: *n* 2/6). Differences were significant (*p* < 0.05).

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## 3.4 Lymphoproliferation assay

The IFN- $\gamma$  values obtained from splenocytes of mice inoculated with the different strains 314 and isolates, unstimulated, stimulated with Concanavalin A, stimulated with TLA of RH, 315 316 stimulated with homologous TLA and its statistical comparison are shown in Figure 4. 317 Significance differences were observed among the splenocytes stimulated with RH TLA, showing TgMr and TgSb the highest values (p < 0.05). Also, TgMr splenocytes showed the 318 highest value (p < 0.05) when stimulated with homologous TLA. No significant differences 319 were detected between homologous TLA and RH TLA stimulation, only TgMr showed an 320 slightly higher IFN-y value when challenged with homologous TLA than with the RH TLA. 321 322 Results from all the assays are summarized in Table 1.

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#### 324 4. DISCUSSION

Most studies with T. gondii atypical genotypes biological behavior have focused the 325 analysis on mice virulence. In this study, we evaluated the in vitro capacity of invasion and 326 327 replication of T. gondii atypical isolates in correlation with an in vivo model. We adapted a methodology previously used in the biological characterization of reference genotypes of T. 328 gondii and the related protozoan N. caninum (Alomar et al., 2013; Dellarupe et al., 2014b). 329 The clonal reference strains in this study showed an *in vitro* behavior very similar to that 330 described by other researchers (Cañedo-Solares et al., 2013; Saadatnia et al., 2010). The 331 clonal type I (RH = virulent) showed low invasion capacity but high replication capacity, 332

while the clonal type II (ME49 = intermediate virulence) showed high invasion capacity, 333 but low replication capacity (Cañedo-Solares et al., 2013; Saadatnia et al., 2010). On the 334 other hand, clonal type III (VEG = non virulent), presented intermediate invasion and low 335 replication, similar to what was previously described (Malkwitz et al., 2018). The doubling 336 time of the RH strain tachyzoites was slightly faster than that recorded in previous studies 337 (Cañedo-Solares et al., 2013), which could be related to limited cell culture adaptation and 338 strain re-isolation in mice (Contreras-Ochoa et al., 2012; Dellarupe et al., 2014b). Malkwitz 339 et al., 2018, reported the ME49 replication values in Vero cells higher than NED strain 340 341 (clonal type III) after 12 hpi., using qPCR. In our study, the ME49 strain showed an active behavior at the beginning of infection (high invasion at 6 hpi), but low and slow replication 342 (most PVs only with 1 or 2 tachyzoites at 18 hpi), in contrast with the clonal type III (VEG 343 strain) that showed low replication capacity but PVs even with 8 tachyzoites. The in vitro 344 behavior we observed in the ME49 and VEG strains could be associated with its reported 345 346 mice virulence, where a low and slow replication would allow the infected cells survival 347 and the establishment of an effective immune response (Dubremetz and Lebrun, 2012; Mordue et al., 2001; Sibley and Boothroyd, 1992). The atypical isolate TgMr (# 14) had an 348 in vitro behavior similar to the RH strain with respect to its low invasion capacity (even 349 lower than RH), however, it had the lowest replication of all strains and isolates used in the 350 351 assay. On the other hand, the atypical isolate TgSb (# 163) had an intermediate invasion capacity (similar to VEG strain) and a high replication (similar to RH strain). Despite the 352 fact that the capacity and the intrinsic mechanism of invasion by reference strains (mainly 353 RH) has been extensively evaluated (Contreras-Ochoa et al., 2012; Dubey et al., 1998; 354 Dubremetz and Lebrun, 2012; Weiss and Kim, 2014), the results of the present study 355 suggest it is negatively related to virulence. In addition, considering our results, the 356

capacity of invasion and possibly its intrinsic mechanism, could not be fully extrapolated
between different *T. gondii* strains.

It has been stated that the clonal strains virulence correlates with high replication capacity 359 (Cañedo-Solares et al., 2013; Dubremetz and Lebrun, 2012; Malkwitz et al., 2018). Based 360 on our results, the RH strain and TgSb isolate showed the highest speeds (or the lowest 361 doubling times), the replication and the "speed" of division correlates positively with 362 virulence. Invasion and replication capacity have been used as phenotypic virulence traits 363 364 for N. caninum isolates (Dellarupe et al., 2014b) and were useful in this work to evaluate 365 the behavior of clonal and atypical T. gondii isolates. Nevertheless, the total amount of intracellular parasites for TgMr and TgSb were lower than the three reference strains, 366 suggesting that there may be other virulence factors not directly associated with invasion 367 and replication. In summary, the atypical isolates of T. gondii obtained from M. rufogriseus 368 and S. boliviensis have a different in vitro behavior than the clonal reference strains, 369 showing invasion and replication characteristics similar to the virulent RH clonal type, 370 371 respectively.

Both atypical isolates showed 100 % morbi-mortality, using doses as low as  $1 \times 10^2$ tachyzoites, similar as previously described using similar atypical isolates (Gennari *et al.*, 2015; Pardini *et al.*, 2019; Pena *et al.*, 2008; Rajendran *et al.*, 2012; Rego *et al.*, 2017; Silva *et al.*, 2014). Probably the use of a lower parasite dose could help to identify possible virulence differences, as suggested by other authors (Saraf *et al.*, 2017). On the other hand, mice infected with clonal strains (ME49 and VEG) had a 0 % morbi-mortality and were considered as non-virulent (Mordue *et al.*, 2001; Sibley and Boothroyd, 1992).

379 Mice infected with atypical genotypes, showed absence (TgMr) or low severity (TgSb) of 380 CNS lesions, possibly due to sacrifice in the acute stage of the infection (8-10 dpi) as

reported by others (Mordue et al., 2001; Pinheiro et al., 2015). Despite of this, T. gondii 381 DNA was detected in all CNS and lung samples, confirming the presence of the parasites. 382 Mice infected with ME49 and VEG strains showed no clinical signs, however, severe 383 lesions in the CNS were detected, which could be related to a cell-mediated response 384 controlling the excessive multiplication of protozoa (Weiss and Kim, 2014). These mice 385 were considered chronically infected as confirmed by the tissue cysts observation in CNS 386 (Dubey, 2010). In addition, 5/6 and 2/6 mice infected with ME49 resulted positive to T. 387 388 gondii DNA in CNS and lung, respectively, possibly due to the preferential location in 389 CNS of the chronic stages of the infection (Costa et al., 2018; Dubey, 2010). As expected for non-lethal clonal strains, tachyzoites multiplication was controlled by the host immune 390 system and protozoa were confined to CNS cells as bradyzoites (Pinheiro et al., 2015). 391 Additionally, the differences in the IgG titers obtained in this study could be associated 392 with the time of infection (Dubey, 2010). 393

In our study, the isolate with the genotype # 163, was characterized as virulent, both in *in vitro* and in the *in vivo* assays, similar as reported for one of the isolates from Rego *et al.*, (2017). It seems that the same genotype could present differences in virulence and therefore the biological behavior could not be 100 % predictable from the molecular markers used in this work (Rego *et al.*, 2017). Similar to other studies, our results indicate that isolate with the genotype # 14, is characterized as virulent (Pardini *et al.*, 2019; Pena *et al.*, 2008; Rajendran *et al.*, 2012).

401 As a final goal, mice cellular (IFN- $\gamma$ ) immune response was evaluated. In both cases, the 402 isolates *TgMr* and *TgSb* induced a higher IFN- $\gamma$  production than clonal strains ME49 and 403 VEG, similar to what has been described by other authors (Chen *et al.*, 2016; Wang *et al.*, 404 2016). The extremely high IFN- $\gamma$  production from infected mice splenocytes (especially

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with TgMr), could indicate a greater number of effector lymphocytes. These results agree 405 with those reported by other researchers in which high levels of IFN- $\gamma$  were detected in 406 lethal infections (RH strain), whereas moderate levels were detected in non-lethal infections 407 (Mordue *et al.*, 2001). Therefore, the adaptive immune response generated by these atypical 408 isolates could contribute to the pathogenesis and the fatal outcome of toxoplasmosis 409 (Mordue et al., 2001; Weiss and Kim, 2014). It has been suggested that the T. gondii strain 410 used to prepare the TLA affects the cytokines produced by lymphocytes in vitro (Rodgers 411 412 et al., 2005). Our results indicate that all TLA stimulated splenocytes produced similar 413 amounts of INF-D. In general, splenocytes stimulated with TLA from RH, produced slightly higher concentrations with the exception of those stimulated with homologue of 414 TgMr. It is probable that the TgMr isolate expresses different antigens recognized as 415 virulence factors, such as rhoptry proteins, which could induce an overproduction of IFN- $\gamma$ 416 in infected animals (Dubremetz and Lebrun, 2012; Rodgers et al., 2005). Splenocytes were 417 collected at the time of sacrifice which differed between atypical isolates and clonal strains 418 419 (ME49 and VEG) leading to potential differences. Probably more accurate results could be obtain performing this assay at fix time, however, it requires a higher number of infected 420 mice and the use of non-lethal dose or strains, which is not always possible with highly 421 virulent atypical strains. In summary, the overproduction of IFN- $\gamma$  in mice infected with 422 423 TgMr and TgSb could be related to the expression / secretion of immune response modulators by these protozoa and could be associated with the higher virulence detected in 424 the in vivo model (Dubremetz and Lebrun, 2012; Mordue et al., 2001; Rodgers et al., 425 2005). On the other hand, the evaluation of the humoral response (IgG titers) did not allow 426 comparisons since blood sampling was conducted at different times. Futures studies aiming 427

428 humoral response evaluation should considerer sampling at fix time (Chen et al., 2016; Tao429 et al., 2013).

Finally, an in vitro invasion-replication index was established that could correlate with 430 431 virulence and mice morbi-mortality inversely (Table 1). The RH strain and the atypical isolates, presented the lowest index values, mainly due to a low invasion. The lower 432 invasion-replication index was associated with the higher virulence, although a larger 433 number of strains and isolates should be evaluated to confirm this assumption. Other 434 435 factors not measured in the *in vitro* assay may possibly be influencing virulence such as the 436 expression and allelic profiles of ROP proteins (Dubremetz and Lebrun, 2012). In the same way, CS3 is a useful marker to predict the virulence of T. gondii in the "atypical" isolates 437 from Brazil (Silva et al., 2014). It would be interesting to analyze the mentioned molecular 438 markers in a higher number of isolates, including atypical isolates obtained from Argentina. 439 It would be important to evaluate the *in vitro* behavior of other phylogenetically related T. 440 gondii atypical isolates in order to confirm the usefulness of the assay presented here as 441 442 complement / substitute to mouse model experiments. Also, performing the protocols described in this study with different cell lines, could allow the improvement of virulence 443 prediction and the design of comparable studies (Contreras-Ochoa et al., 2012). The 444 validation of an in vitro invasion-replication index could be useful to indirectly predict 445 446 virulence in mice.

In conclusion, the *T. gondii* atypical isolates # 14 and # 163 resulted in a different *in vitro* behavior than clonal strains, with low invasion-replication indexes, but showing high virulence in mice model in association with high levels of INF- $\Box$ .

450

### 451 **5. ACKNOWLEDGEMENTS**

We would like to thank Dr. J. P. Dubey from the Animal Parasitic Diseases Laboratory,
Agricultural Research Center (USDA), Beltsville, Maryland, USA for providing the clonal
type III reference strain (VEG). Also, Dr. Alejandra Quiroga from LAPEVET "Dr.
Bernardo Epstein", FCV, UNLP, for the histopathological analysis and Isidoro Ercoli and
Selva Peñaloza for their excellent technical assistance.

457

#### 458 **6. FUNDING**

459 This work was partially supported by FONCyT through PICT 2015-0788, UNLP I+D

460 project V250 and the UNLP young researchers grant 1145/17. Mariana Bernstein was

461 awarded a doctoral fellowship from CONICET, and the present study is part of her doctoral

462 thesis.

463

## 464 **7. CONFLICT OF INTERESTS**

465 The authors declare no conflict of interests.

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- 566

## 567 Figure captions:

**Figure 1**. *In vitro* assay: Invasion values (number of parasitophorous vacuoles at 6 hpi, **A**) and replication values (number of tachyzoites at 18 hpi, **B**) for each *T. gondii* strain and isolate. Different letters indicate significant differences (p < 0.05, LSD Fisher). Reference: PVs = parasitophorous vacuoles.

Figure 2. *In vitro* assay: Comparison of the percentages of PVs of 1, 2, 4, 8 and 16
tachyzoites for each *T. gondii* strain and isolate at18 hpi. Reference: PVs = parasitophorous
vacuoles.

**Figure 3**. *In vivo* assay: Representative photomicrographs of *T. gondii* infected mice CNS sections showing different grades of lesions (H&E staining). Grade 1: Neural necrosis

577 (arrow), 40x (**A**, mouse inoculated with *TgSb*). Grade 2: mild non-suppurative focal 578 meningitis (arrow), 20x (**B**, mouse inoculated with VEG). Grade 3: numerous mononuclear 579 cells, focal gliosis (arrow), 20x (**C**, mouse inoculated with VEG). Two large cysts in 580 cerebellum without inflammatory reaction (arrow), 40x (**D**, mouse inoculated with VEG). 581 Scale bars = 100  $\mu$ m.

**Figure 4**. Lymphoproliferation assay: production values of IFN- $\gamma$  by splenocytes of mice inoculated with the different *T. gondii* strains and isolates, unstimulated (A, black), stimulated with Concanavalin A (B, orange), stimulated with TLA of RH (C, purple) and homologous TLA (D, gray). Values obtained by ELISA (pg/ml) were log10 transformed to apply an ANOVA. Different letters indicate significant differences among the columns with the same color (p < 0.05). <sup>1</sup> Note: stimulation with Concanavalin A of splenocytes from mice infected with *TgMr* exceeded the IFN- $\gamma$  detection values from ELISA kit.

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Table 1. Summary of the results from the in vitro and the in vivo assays with different T. gondii strains and isolates.

Isolates / Strains	Toxo DB- genotype			Invasion- replication index <sup>c</sup>	In vivo assay							
		Invasion <sup>a</sup>	Replication <sup>b</sup>		Morbi- mortality <sup>d</sup> (%)	Sacrifice	Lesions <sup>f</sup> (%)	<i>T. gondii</i> cysts <sup>g</sup> (%)	<i>T. gondii</i> DNA in CNS <sup>h</sup> (%)	<i>T. gondii</i> DNA in lung <sup>i</sup> (%)	IFN-□ Production <sup>j</sup> (pg/ml)	IgG titers <sup>k</sup> (n)
1-TgSb	# 163	443	535	237	6/6 (100)	10 dpi	5/6 (83,3)	0/6 (0)	6/6 (100)	6/6 (100)	3.86/3.64 (12542/8005)	100/50 (4/6;2/6)
2-TgMr	# 14	143	295	42	6/6 (100)	8-9 dpi <sup>e</sup>	0/6	0/6 (0)	6/6 (100)	6/6 (100)	4.15/4.26 (17333/24288)	50/25 (4/6;2/6)
3-RH	# 10	310	757	234	ND	ND	ND	ND	ND	ND	ND	ND
4-ME49	# 3	1108	713	790	0/6 (0)	30 dpi	5/6 (83.3)	0/6 (0)	5/6 (83.3)	2/6 (33.3)	3.35/3.27 (2977/2227)	6400/3200 (4/6;2/6)
5-VEG	# 2	394	662	260	0/6 (0)	30 dpi	4/6 (66.6)	4/6 (66.6)	5/6 (83.3)	5/6 (83.3)	3.51/3.22 (3811/1741)	12800 (6/6)

References:

<sup>a</sup>: Invasion: average number of PVs at 6 hpi.

<sup>b</sup>: Replication: average number of the total amount of parasites that showed at least one event of endodiogeny (PVs with 2 or more tachyzoites) at 18 hpi.

<sup>c</sup>: Invasion-replication index: (average invasion 6 hpi \* average replication 18 hpi) / 1000.

<sup>d</sup>: Morbi-mortality: total of clinically ill and sacrificed animals / total of infected animals \* 100.

<sup>e</sup>: Mice inoculated with 1x10<sup>3</sup> tachyzoites were sacrificed 8 dpi, while mice inoculated with 1x10<sup>2</sup> tachyzoites were sacrificed 9 dpi.

<sup>f</sup>: Number of mice with CNS lesions / number of mice inoculated.

<sup>g</sup>: Number of mice with *T. gondii* cysts in CNS / number of mice inoculated.

h: Number of mice with *T. gondii* DNA detection in CNS / number of mice inoculated.

<sup>i</sup>: Number of mice with detection of *T. gondii* DNA in lung / number of mice inoculated.

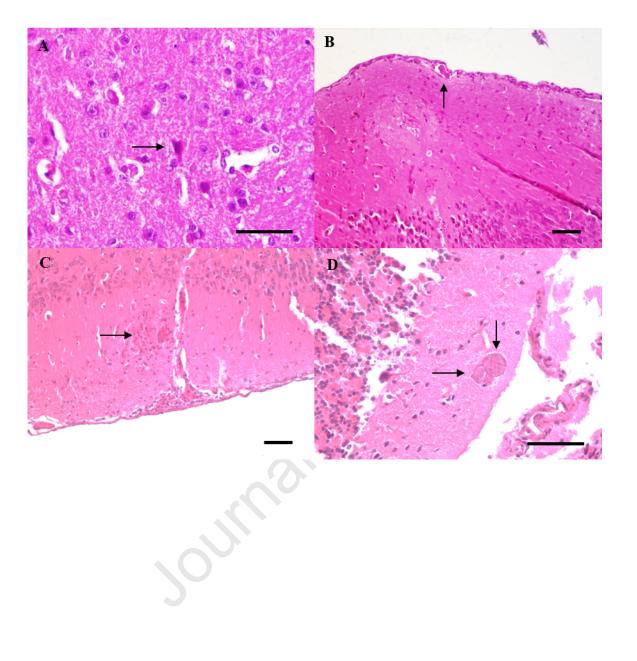
<sup>j</sup>: Average values obtained from splenocytes stimulated with TLA from RH/and homologous TLA. Values were log10 transform for statistical analysis. Original average values obtained by ELISA (pg/ml) between brackets.

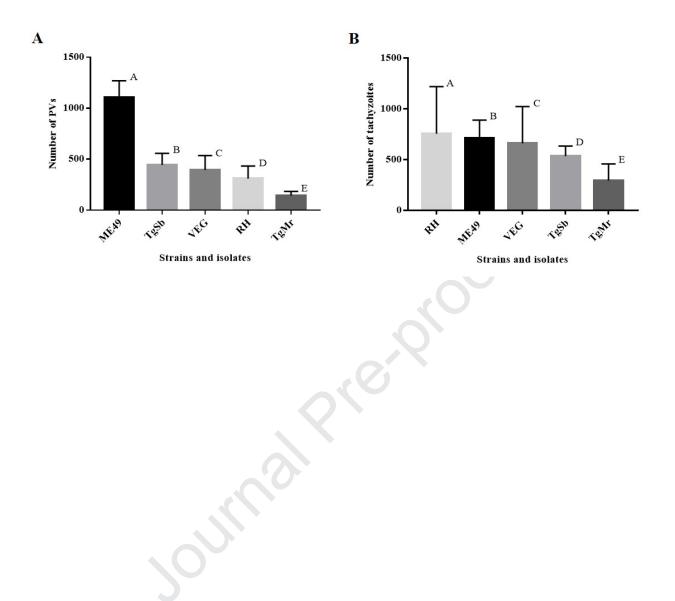
<sup>k</sup>: IFAT final titers: number of mice with IgG final titer / number of mice inoculated.

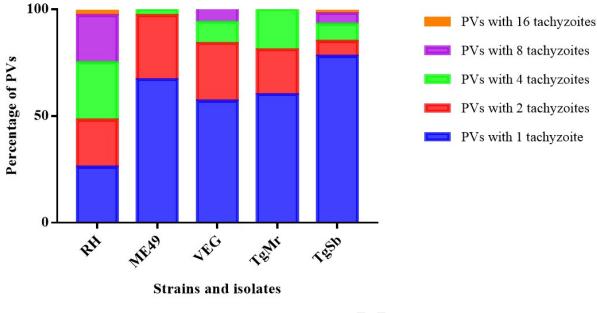
dpi: days post infection.

ND: no data.

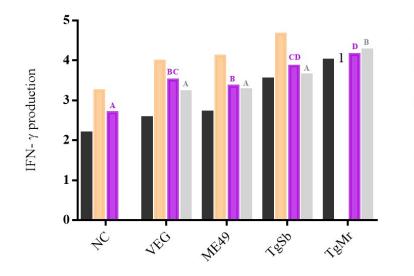
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Groups of mice infected with different *T. gondii* strains and isolates

A- Splenocytes non-stimulated B- Splenocytes stimulated with concanavalin A

C- Splenocytes stimulated with TLA from the RH strain

D- Splenocytes stimulated with TLA from homologous tachyzoites

## HIGHLIGHTS

- In vitro and in vivo behavior of 2 T. gondii atypical isolates was evaluated. •
- Atypical isolates showed similar in vitro behavior to the virulent type I strain. .
- Atypical isolates showed 100 % morbi-mortality in mice with a  $10^2$  parasites. .
- High virulence was associated with high IFN-□ in lymphoproliferation assay. .
- Low values of invasion-replication index correlated with high virulence in mice. .