Soluble total antigen derived from *Toxoplasma gondii* RH strain prevents apoptosis, but induces anti-apoptosis in human monocyte cell line

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**Abstract:** Apoptosis plays crucial role in the pathogenesis of toxoplasmosis, as it limits further development of the disease. The current study aimed to investigate the effects of different concentrations of soluble total antigen (STAg) of *Toxoplasma gondii* (Nicolle et Manceaux, 1908) on the apoptotic and anti-apoptotic pathways. PMA-activated THP-1 cell line was sensed by *T. gondii* STAg and the expression patterns of caspase-3, -7, -8, -9, Bax, Bcl-2, and Mcl-1 were evaluated. The results showed statistically significant concentration-dependent overexpression of both Bcl-2 (*P*-value < 0.0001) and Mcl-1 (*P*-value = 0.0147). The cas-7 showed overexpression in all concentrations (*P*-value < 0.0001). The cas-3 was suppressed in concentrations 100, 80, and 40 µg, but statistically significant downregulated in concentrations 10 and 20 µg. The Bax was suppressed in concentrations 100 to 20 µg, while it slightly downregulated 1.42 fold (*P*-value = 0.0029) in concentration 10 µg. The expression of cas-8 and -9 was suppressed in all concentrations. Our results indicated that *T. gondii* STAg downregulated and suppressed apoptotic and upregulated anti-apoptotic pathways. The upregulation of cas-7 in this study may indicate the role of *T. gondii* STAg in activation of inflammatory responses.

**Keyword:** *Toxoplasma gondii*, Soluble total antigen, Apoptosis, Mcl-1, Bcl-2.

Toxoplasmosis is a parasitic disease caused by *Toxoplasma gondii* (Nicolle et Manceaux, 1908), which usually causes no obvious symptoms in adults (Pappas et al. 2009). *Toxoplasma gondii* can infect all mammals; however, only felids are definitive hosts (Tenter et al. 2000). It is a cosmopolitan protozoan parasite, infecting more than one billion people of which most of them have chronic asymptomatic form (Aguirre et al. 2019). In individuals who have compromised immune systems, toxoplasmosis may lead to serious health problems, severe symptoms, and eventually death (Xiao and Yolken 2015). Infection during the pregnancy may affect offspring, which is known as congenital toxoplasmosis (Lourido 2019).

Apoptosis is the physiological cell death, which leads to elimination of unwanted or useless cells (D’Arcy 2019). This process plays a crucial role in the pathogenesis of toxoplasmosis, as well. During toxoplasmosis, apoptosis can play as a double edged sword (Mammari et al. 2019). Although this phenomenon may lead to cell death and limits further development of toxoplasmosis, *T. gondii* can use apoptosis to suppress the immune responses (Lüder et al. 2001). Ultrastructure evidence indicated that during the infection with *T. gondii*, tachyzoites were found in the cytosol of HeLa cells 10 min after infection and started to replicate 2–5 h after the infection (Toula et al. 2011). After the entry, *T. gondii* modulates cell survival and cell death cycles to dysregulate the immune responses, which leads to unrestricted parasitemia and destruction of host tissues (Opferman and Korsmeyer 2003, Payne et al. 2003, Mammi et al. 2019).

In mammalian cells, there are at least two main pathways that lead to apoptosis, i.e., intrinsic pathway (mitochondria-mediated events) and extrinsic pathway (death receptor-mediated events) (Elmore 2007, Obeng 2021). B-cell lymphoma (Bcl) 2 family proteins, cytochrome C, adaptor proteins, and caspases are the components of intrinsic pathway. The intrinsic pathway is stimulated by producing intracellular signals such as radiation (DNA
damage), absence of certain growth factors, hormones, and cytokines. These stimuli cause changes in the mitochondrial outer membrane permeabilisation (MOMP) and release of cytochrome c (pro-apoptotic proteins) from the intermembrane space into the cytosol. In addition, caspase (cas)-9 activation leads to cas-3 activation and cell death (Green et al. 2004, Jin and El-Deiry 2005, Akhtar and Bokhari 2021).

Death receptors and ligands, adaptor proteins, and caspases are the components of extrinsic pathway. Until now, 14 caspases are known in mammals of which ten major caspases have been identified and broadly categorised into signaling/initiator (2, 8, 9, and 10), effector-executioner (3, 6, and 7), and inflammatory caspases (1, 4, and 5) (Boatright et al. 2003, Yanumula and Cusick 2021).

The regulation and control of apoptotic mitochondrial events occur through members of the Bcl-2 family. Bcl-2 family are key regulators in the molecular mechanisms of apoptosis. This family includes anti-apoptotic proteins, such as Bcl-2, Bcl-x, Bcl-XL, and Bcl-w, and pro-apoptotic proteins, i.e., Bax, Bak, Bid, Bad, Bim, and Bik. These proteins either are free in cytosol or attached to cell membrane. Bcl-2 family regulates the release of cytochrome C from the mitochondria via alteration of mitochondrial membrane permeability (Cheng et al. 2001, Akhtar and Bokhari 2021).

The exact effect of *T. gondii* or its antigens on the apoptotic and anti-apoptotic pathways are still unknown and previous study revealed controversial results (Goebel et al. 2001, Hwang et al. 2010, Li et al. 2016). In the current study, the effects of different concentrations of soluble total antigen (STAg) of *T. gondii* tachyzoites on the apoptotic and anti-apoptotic pathways were evaluated on human monocytes cell line.

**MATERIALS AND METHODS**

**Parasite preparation**

To provide sufficient number of *Toxoplasma gondii, 1 \times 10^8* tachyzoites of the virulent RH strain were inoculated to the peritoneal cavity of white laboratory mice, and were harvested from infected mice after 3–7 day of inoculation. Briefly, peritoneal fluid of infected mice was aspirated and centrifuged at 10,000 g for 10 min. Supernatant was discarded and the pellet containing *T. gondii* tachyzoites were washed twice in phosphate-buffered saline (PBS; pH = 7.4). To prepare STAg, tachyzoites were suspended in 1 mL of lysis buffer (1% sodium dodecyl sulfate and 5mM EDTA) and sonicated (30 pulses for three times and 1 min intervals) on ice as described by Pazoki et al. (2021). After centrifugation for 10 min at 10,000 g, total protein concentration was measured using the Bicinchoninic acid (BCA) assay (Parstous, Mashhad, Iran).

**THP-1 cell line cultivation**

The human monocyte THP-1 cell line was propagated in 1640 medium (RPMI 1640; Biosera, Arya Tous, Tehran, Iran) supplemented with 10% (v/v) fetal bovine serum (FBS; Gibco, Thermo Fisher Scientific, Massachusetts, USA), 1% penicillin/streptomycin (Thermo Fisher Scientific) at 37 °C and 5% CO_2. THP-1 cells were seeded by hemocytometer slide and 5 \times 10^4 of cells were seeded in each well of a 12-well cell culture plate. To differentiate monocytes to macrophage M0, 30 ng/ml of phorbol 12-myristate 13-acetate (PMA; Santa Cruz Biotechnology Cat. No. sc-3576) were added to the cells. The plates were incubated for 36 h at 37°C in 5% CO2. Prior to the experiments and after morphological confirmation, supernatant medium containing non-adherent cells was discarded and replaced with PMA- and LPS-free cell culture medium (RPMI 1640 medium with 10% FBS and 1% antibiotic). M0 macrophages were rested for 24 h prior to experiments.

**Treatment of THP-1 cell line with different concentrations of Toxoplasma gondii STAg**

THP-1 cells were treated with 100, 80, 40, 20, and 10 µg/ml STAg. A well was treated with 20 ng/ml LPS (Santa Cruz Biotechnology Cat No. sc-3535) to compare its induction pattern with *T. gondii* STAg. A well full of PMA-activated THP-1 cell without any treatment either by *T. gondii* STAg or LPS, was considered as the control group. All groups were duplicate and evaluated 18 h after exposure.

**RNA isolation, cDNA synthesis, and Real-time RT-PCR**

Total RNA was extracted from THP-1 cells by the total RNA extraction kit (Yekta Tajhiz Azma, Tehran, Iran) according to the manufacturer’s protocol. DNase (Thermo Fisher Scientific™) treatment was performed to remove the probable residual DNA and to improve the quality of the extracted RNA according to the manufacturer’s protocol. The RNA solution was finally collected and stored at -70 °C until cDNA synthesis. Prior to cDNA synthesis, the concentration of purified RNAs was determined by NanoDrop (NanoDrop Technologies, Wilmington, Delaware, USA) and RNA adjustment (normalisation) was performed. cDNA was constructed using cDNA synthesis kit (Yekta Tajhiz Azma, Tehran, Iran) according to the manufacturer’s protocol.

To study the expression levels of the cas-3, 7, 8, 9, Bax, Bcl-2, and Mcl-1, qRT-PCR using specific primers (Table1) was employed. Rotor-Gene Q (Qiagen, Hilden, Germany) thermocycler was employed. The standard amplification program was initial denaturation 95 °C for 10 min followed by 40 cycle of denaturation at 95 °C for 20 s, annealing at 59–61 °C for 30 s, and extension at 72 °C for 20 s. Melt curve analysis was performed to determine the specificity of each reaction. Reactions were carried out in duplicate and the transcriptional level of each gene was treated with β-actin as housekeeping gene. The relative expression level of each gene was analysed using the 2^(-ΔΔCt) method and relative expression software tool (REST).

**Statistical analysis**

ANOVA test was applied to analyse the protein fold expression at different concentrations. Student’s t-test was applied to analyse the real-time PCR data. *P*-value < 0.05 was considered statistically significant. Statistical analysis was performed using GraphPad Prism software version 8.3.0.538.

**RESULTS**

The results of real-time PCR revealed increased expression of Bcl-2 and Mcl-1, as anti-apoptotic factors, in...
Fig 1. The comparison of the expression levels of Bcl-2 (A), Mcl-1 (B), Cas-7 (C), Cas-3 (D), and Bax (E) in THP-1 cell line regarding different concentrations of the soluble total antigen of Toxoplasma gondii (Nicolle et Manceaux, 1908) STAg. NS: not significant; * P value < 0.05; ** P value < 0.01; *** P value < 0.001.

all concentrations of T. gondii STAg. Regarding the concentrations, the highest expression level in Bcl-2 was seen in 100 µg/ml of T. gondii STAg (25.55 fold; P-value = 0.0008). In addition, a different expression was observed in Bcl-2 mRNA in other concentrations. Except concentration 100 µg, the Bcl-2 showed a statistically significant overexpression from 80 µg/ml to 10 µg/ml concentrations of T. gondii STAg (P-value < 0.0001) (Fig 1A).

Mcl-1 mRNA statistically overexpressed after exposure with T. gondii STAg in all concentrations (P-value = 0.0147). The highest expression was seen in concentration 20 µg/ml of T. gondii STAg (5.57 fold; P-value = 0.0103). Similar to Bcl-2, except the concentration 100 µg/ml, increased expression was observed in low concentrations of T. gondii STAg (Fig 1B).

Cas-7 showed overexpression in all concentrations. The highest expression was observed in test wells sensed by
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100 µg T. gondii STAg (19.53 fold; P-value = 0.002). In contrast to Bcl-2 and Mcl-1, a downregulation was seen in low concentrations (P-value < 0.0001) (Fig 1C).

The expression of cas-3 was suppressed in wells, which were sensed with concentrations 100, 80, and 40 µg/ml of T. gondii STAg, whereas it showed statistically significant downregulation in concentrations 10 and 20 µg/ml of T. gondii STAg with folds 4.53 (P-value = 0.01) and 2.21 (P-value = 0.028), respectively (Fig. 1D). The expression of Bax (as pro-apoptotic protein) was suppressed in concentrations 100 to 20 µg/ml of T. gondii STAg, whereas it was slightly downregulated 1.42 fold (P-value = 0.0029) in concentration 10 µg (Fig. 1E). Interestingly, the expressions of cas-8 and -9 were suppressed in all concentrations of T. gondii STAg.

**DISCUSSION**

At the beginning of the infection, Toxoplasma gondii provokes innate immunity system that leads to secretion of many different effectors. The host innate immunity is known to play a critical role in defense against intracellular infection like toxoplasmosis (Sasai et al. 2018, Lima and Lodoen 2019). Followed by the innate immunity, the cell-mediated immune responses and released cytokines limit parasite replication and control toxoplasmosis (Sasai et al. 2018). Upon entrance of T. gondii, the host’s cell em-
pleys immune responses, signalling pathways, transcriptional changes, and apoptosis mechanisms to prevent the development of the infection (Friedrich et al. 2017, Lima and Lodoen 2019). Besides the regulation of the immune responses, apoptosis acts as an effector mechanism of immune cells such as, natural killer (NK) cells and cytotoxic T lymphocytes, to eliminate infected target cells (Lieberman 2003, Opferman and Korsmeyer 2003).


As a results, cas-3 and 9 were suppressed in *T. gondii* infected THP cells and cas-7 showed an increased expression at all concentrations. The expression of the Bax gene, as pro-apoptotic factors, was inhibited. In addition, the expression of Bcl-2 and Mcl-1, as anti-apoptotic factors, were increased upon treatment with *T. gondii* STAg. In the line of our study, the experimental evidence suggests that *T. gondii* inhibits apoptosis of host cells with manipulating of the apoptosis pathways. Goebel et al. (2001) reported a downregulation of cas-9 and cas-3, and blocked release of cytochrome C from mitochondria in HL-60 and U937 cell lines treated by *T. gondii*.

Furthermore, evidence indicated that the anti-apoptotic Mcl-1 and Bcl-2 increased after *T. gondii* infection (Goebel et al. 2001, Molesta et al. 2003). In addition, Hwang et al. (2010) suggested that *T. gondii* inhibited cas-3, but up-regulated anti-apoptotic pathways via cytochrome C-mediated caspase activation. Although there are few data on the apoptosis and anti-apoptosis pathways induced by *T. gondii* in animal models, Kim and Denkers (2006) demonstrated that *T. gondii* provokes Gi-protein-dependent PI 3-kinase signaling to preclude apoptosis in host cell.

In the line of our findings, Payne et al (2003) suggested inhibition of cas-8 activity by *T. gondii* in murine fibroblasts. Caspases are important effectors of programmed cell death and their roles in apoptosis have been investigated (Hotchkiss and Nicholson 2006). Caspase activation is stimulated by initiators cas-8 and -9 that activate executioner cas-3. Activation of cas-3 and cas-7 ultimately leads to apoptosis (Fig. 2). Lakhani et al. (2006) showed that although cas-3 and cas-7 are key mediators during the apoptosis process, cas-3 knockout mice died prematurely, but the cas-7 knockout mice represented no phenotypic abnormalities. As well, cas-3–cas-7 knockout mice suffered from early perinatal death.

Evidence points out the non-apoptotic functions of cas-3 and cas-7 in different cell processes like cell proliferation, cell cycle regulation, cell differentiation, and inflammation. Indeed, cas-3 and cas-7 seem to have independent functions during apoptosis (Lakhani et al. 2006, Lamkanfi et al. 2009, Lamkanfi and Kannegeanti 2010). Actually, executioner caspases (cas-3 and/or cas-7) would be activated under inflammatory conditions, upon the activation of cas-9 and -1, as initiator caspases in the apoptosome and inflammasome complex, respectively (Hotchkiss and Nicholson 2006, Lamkanfi et al. 2009).

Unlike cas-3, cas-8, and cas-9, cas-7 was upregulated in our study. Cas-7 was suggested to be involved in inflammation. Lamkanfi et al. (2008) showed that upon the LPS stimulation, cas-7 is activated by cas-1 proposing an apoptosis-independent regulatory role for cas-7 in pro-inflammatory genes expression. In addition, it was suggested that macrophages are able to restrict *Legionella pneumophila* via NLRC4/IPAF inflammatory factors and cas-7 activation (Akhter et al. 2009).

These results are also supported by experimental studies that have shown that *T. gondii* or its proteins increase the expression levels of inflammasomes and inflammation (Witola et al. 2011, Chai and Shi 2014, Wang et al. 2019). However, the main differences between results most probably back to either the time points of experiments or the inducer. For example, Wang and Gao (2016) demonstrated that tachyzoites of *T. gondii* RH strain prevented the proliferation and induced apoptosis in HCC H7402 cells. In contrast, Wu et al. (2020) showed that rhoptry protein 16 (ROP16) inhibited programmed cell death in human 293T cell. These studies may help interpretation of our results. Actually, in the present study we used *T. gondii* STAg, which contains a broad spectrum of proteins that unlike the alive tachyzoite, these proteins may manipulate apoptosis pathways via different manner.

Some studies also proposed that strains of *T. gondii* and time points affect the expression of genes which are involved in the apoptosis pathway. Angeloni et al. (2009) reported different results, for example, in another study, the strain of *T. gondii* RH strain prevents apoptosis (Ngobili et al. 2015, Tomita et al. 2003), while Pld-1 and Pld-2 were increased upon treatment with *T. gondii* RH strain (Rosato et al. 2007).

### Table 1. Primers used in this study.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Sense primer (5′-3′)</th>
<th>Antisense primer (5′-3′)</th>
<th>Refs</th>
</tr>
</thead>
<tbody>
<tr>
<td>Caspase 3</td>
<td>CAGAAACTGAGACTGGCCAAGT</td>
<td>GCTTGGCCAGCATCCTGGTTCA</td>
<td>Xu et al. (2012)</td>
</tr>
<tr>
<td>Caspase 7</td>
<td>GAGAAGAGCCCTAGGTGCCAGTGG</td>
<td>TCCCCGTGCTGTTGTGTGTT</td>
<td>Tomita et al. (2003)</td>
</tr>
<tr>
<td>Caspase 8</td>
<td>TCTGAAATCTGATAGACATGAC</td>
<td>AGACTGAGATTGTCGATACCTAC</td>
<td>Ngobili et al. (2015)</td>
</tr>
<tr>
<td>Caspase 9</td>
<td>GTCCTCAAAACTCTGGTGGGAC</td>
<td>GCCCAAGCTCTTTTTCATCC</td>
<td>Ngobili et al. (2015)</td>
</tr>
<tr>
<td>Bax</td>
<td>ATGTGTCCTGACGGCAACTCC</td>
<td>AGTCCAAATGTCAGCCCA</td>
<td>Buskaran et al. (2021)</td>
</tr>
<tr>
<td>Bcl-2</td>
<td>AAGGCCGCGCCAGCACTTCT</td>
<td>GGTGCGGTCGTTCTGACGTTCA</td>
<td>Laffon et al. (2001)</td>
</tr>
<tr>
<td>Mcl-1</td>
<td>GGGACGAGGTCTGACTCTCATT</td>
<td>GATGCACCTTCTGGTTTATG</td>
<td>Rosato et al. (2007)</td>
</tr>
<tr>
<td>β-act</td>
<td>ATGTGCGCGGAGATTGGTCA</td>
<td>AGTGGGTGGTCCTTTAGAAGTG</td>
<td>Fu et al. (2010)</td>
</tr>
</tbody>
</table>
suggested that *T. gondii* strains affect the apoptosis. They showed that strain ME49, a moderately virulent strain, induced higher apoptosis index compared to RH strain, which is known as a highly virulent strain. Interestingly, our study supports this result that hypervirulent strains such as RH, are probably able to inhibit programmed cell death upon the infection to increase their replication in host cell. In the line of our findings, Contreras-Ochoa et al. (2013) showed that *T. gondii* tachyzoite RH strain controlled the apoptosis pathways in host cells and except the early stage of infection, apoptosis decreased up to 24 h after infection.

Therefore, based on our results, it seems that *T. gondii* STAG probably induces inflammatory responses and suppresses apoptotic pathways in monocyte cell line, although alive tachyzoite may activate apoptosis pathways in host cells. However, we employed RH strain which is a hyper-virulent strain of *T. gondii*, therefore, the results of interaction between host cell apoptosis and *T. gondii* should be interpreted according to different strains of the parasite.

In conclusion, our results indicate that *T. gondii* STAG downregulated and suppressed apoptotic and upregulates anti-apoptotic pathways in human monocyte cell line. In addition, the upregulation of cas-7 in this study may indicate the role of *T. gondii* STAG in activation of inflammatory responses. However, the results of this and previous studies suggest that alive tachyzoite, STAG, and/or specific proteins derived from *T. gondii* may interplay with host cells via different manners.

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**Ethics approval.** All procedures performed in this study were in accordance with the ethical standards (IR.SBMU.RIGLD.REC.1398.035) released by Ethical Review Committee of the Research Institute for Gastroenterology and Liver Diseases, Shahid Beheshti University of Medical Sciences, Tehran, Iran.

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