

Differential Regulation of Proinflammatory Cytokine Expression by Mitogen-Activated Protein Kinases in Macrophages in Response to Intestinal Parasite Infection

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Blastocystis is a common enteric protistan parasite that can cause acute, as well as chronic, infection and is associated with irritable bowel syndrome (IBS). However, the pathogenic status of *Blastocystis* infection remains unclear. In this study, we found that *Blastocystis* antigens induced abundant expression of proinflammatory cytokines, including interleukin 1 β (IL-1 β), IL-6, and tumor necrosis factor alpha (TNF- α), in mouse intestinal explants, in mouse colitis colon, and in macrophages. Further investigation utilizing RAW264.7 murine macrophages showed that *Blastocystis* treatment in RAW264.7 macrophages induced the activation of ERK, JNK, and p38, the three major groups of mammalian mitogen-activated protein (MAP) kinases that play essential roles in the expression of proinflammatory cytokines. ERK inhibition in macrophages significantly suppressed both mRNA and protein expression of IL-6 and TNF- α and mRNA expression of IL-1 β . On the other hand, JNK inhibition resulted in reductions in both c-Jun and ERK activation and significant suppression of all three proinflammatory cytokines at both the mRNA and protein levels. Inhibition of p38 suppressed only IL-6 protein expression with no effect on the expression of ERK activation and significant suppressed. Our study thus demonstrated for the first time that *Blastocystis* could induce the expression of various proinflammatory cytokines via the activation of MAP kinases and that infection with *Blastocystis* could induce the expression of various proinflammatory intestinal diseases through the activation of inflammatory pathways in host immune cells, such as macrophages.

ctivation of the immune system in the gut by microbial infection or loss of tolerance for nonpathogenic commensal microbes leads to inflammation and contributes to the development of intestinal disorders, such as inflammatory bowel diseases (1, 2). Blastocystis is a common enteric protistan parasite and an emerging pathogen found in humans and many animals (3). It is the most frequently isolated protist reported in human fecal samples and has prevalences of up to 10% in developed countries and as much as 50 to 60% in developing countries (4, 5). At least 17 subtypes (STs) of Blastocystis have been identified, with 9 of them (subtype 1 [ST-1] to ST-9) detected in humans (6). Transmission occurs via the fecal-oral route in cyst form via contaminated food or water (4, 7). Higher incidences of Blastocystis infections are reported in immunocompromised patients, such as HIV-infected and cancer patients (8-10). Common clinical symptoms of human Blastocystis infection, blastocystosis, are abdominal pain, diarrhea, vomiting, and bloating (4). Blastocystis infections are frequently associated with irritable bowel syndrome (IBS), as well (11-15).

While the mechanisms of mucosal injury induced by *Blastocys*tis infection in the human gastrointestinal tract remain unclear (16), it has been shown that *Blastocystis* is able to disrupt barrier function and increase permeability in intestinal cells via rearrangement of F actin and ZO-1 distribution (17). Furthermore, *Blastocystis* could induce apoptosis of intestinal cells in a contactindependent manner by involving caspase 3, suggesting that the parasite may secrete virulence factors that initiate the apoptotic pathway (17). *Blastocystis* was able to induce the production of granulocyte-macrophage colony-stimulating factor (GM-CSF) and interleukin 8 (IL-8) in HT-29 and T-84 human colonic epithelial cells (18). The induction of IL-8 in T-84 cells is dependent on NF- κ B with the involvement of cysteine proteases (19). Upregulation of proinflammatory gene expression and intense infiltration of proinflammatory cells in the colon have been shown in rat and mouse models of *Blastocystis* infection (20, 21). A recent study also showed significantly higher levels of IL-17 and IL-23 in mice infected with *Blastocystis* (22). These studies demonstrate the pathogenic potential of *Blastocystis* despite a longstanding controversy as to whether it is an intestinal pathogen.

The uncertainty of *Blastocystis* pathogenesis and observations of differences in virulence have been attributed to different subtypes of the intestinal parasite. This is supported by studies revealing variation of cysteine protease activity between subtypes (23). Additionally, while ST-7 (B) was shown to be more sensitive to nitrosative stress than ST-4 (WR-1), it appeared to possess unique evasion strategies, with its ability to inhibit epithelial nitric oxide

Received 2 July 2014 Returned for modification 2 August 2014 Accepted 20 August 2014 Published ahead of print 25 August 2014 Editor: J. A. Appleton Address correspondence to Kevin Shyong Wei Tan, mictank@nus.edu.sg, or Yongliang Zhang, miczy@nus.edu.sg. Supplemental material for this article may be found at http://dx.doi.org/10.1128 /IAI.02279-14. Copyright © 2014, American Society for Microbiology. All Rights Reserved. doi:10.1128/IAI.02279-14 The authors have paid a fee to allow immediate free access to this article. (NO) production by downregulating epithelial inducible nitric oxide synthase (iNOS) expression (24). These studies suggesting subtype-dependent variation show the need to further investigate differences in pathogenic potential across different subtypes of *Blastocystis*. In addition, host genetic polymorphism could influence the expression of symptoms in response to *Blastocystis* infection. For instance, it has been reported that some IL-8 and IL-10 single-nucleotide polymorphisms (SNPs) could change individual susceptibility, increasing the relative risk of development of IBS in *Blastocystis* carriers (25).

The mitogen-activated protein kinase (MAPK) signaling pathways are evolutionarily conserved and important mediators involved in transducing extracellular signals, such as stress, growth factors, and cytokines, to intracellular responses in order to generate an appropriate physiological response to the stimuli (26, 27). Three major groups of MAPKs exist in mammalian species, namely, the extracellular signal-regulated protein kinases (ERKs), the p38 MAP kinases, and the c-Jun NH₂-terminal kinases (JNKs). MAPKs are activated through a core triple-kinase cascade whereby each MAPK is activated by its specific upstream MAPK kinase (MKK) via phosphorylation of both threonine and tyrosine residues in its activation loop. MKKs are activated after phosphorylation of their serine/threonine residues by MAPK kinase kinases (MKKKs) (28). Activated MAPKs phosphorylate a wide range of substrates, including transcription factors and cytoskeletal proteins (29).

MAPKs play crucial roles in innate immunity by triggering cellular responses and production of cytokines during pathogenic infections (see Fig. S1 in the supplemental material) (30). MAPK activation in macrophages in response to protozoan parasites has been reported extensively. Studies on Trypanosoma cruzi have revealed its ability to activate ERK and p38 with kinetics similar to those of lipopolysaccharide (LPS)-induced MAPK activation, leading to the production of IL-12 and tumor necrosis factor alpha $(TNF-\alpha)$ by macrophages (31). Infection with *Toxoplasma gondii* activates ERK, JNK, and p38 during invasion and proliferation of its tachyzoites (32). MAPK signaling pathways appeared to be involved in increased secretion and expression of monocyte chemotactic protein 1 (MCP-1), macrophage inflammatory protein 1α (MIP- 1α), cyclooxygenase 2 (COX-2), and prostaglandin E2 (PGE2) in RAW264.7 macrophages in response to T. gondii (32). Parallel studies with Entamoeba histolytica showed MAPK activation in RAW264.7 macrophages by its DNA and Gal/GalNAc adherence lectin (33, 34). Apart from cytokine production, p38 and JNK appeared to be involved in macrophage apoptosis induced by Trichomonas vaginalis (27). However, whether Blastocystis infection could induce MAPK activation in host cells is unknown.

In this study, we investigated host-pathogen interactions using the RAW264.7 murine macrophage line model. Using two *Blastocystis* subtypes, ST-4 (WR-1) and ST-7 (B), we demonstrated for the first time that *Blastocystis* can activate MAPK pathways and induce proinflammatory cytokines, including IL-1 β , IL-6, and TNF- α , in a subtype-dependent manner. In addition, we investigated the role of each MAPK signaling pathway in proinflammatory cytokine expression in murine macrophages in response to *Blastocystis* ST-7 (B) and found for the first time that ST-7 (B) serine proteases play a major role in the activation of MAPKs and the expression of proinflammatory cytokines in macrophages.

MATERIALS AND METHODS

Parasite culture and preparation of lysate. For this study, Blastocystis ST-4 (WR-1) and ST-7 (B) were used. ST-4 (WR-1) was isolated from a healthy rat during an animal survey (35). ST-7 (B) was recovered from a patient suffering from gastrointestinal symptoms at the Singapore General Hospital (23). All isolates were categorized according to a recent Blastocystis sp. classification system (36). The parasites were cultured axenically, as described previously (23, 24). In short, cultures of each isolate were maintained in 10 ml of prereduced Iscove's modified Dulbecco's medium (IMDM) containing 10% inactivated horse serum (Gibco) and incubated at 37°C in an anaerobic jar (Oxoid, United Kingdom) with an AnaeroGen gas pack (Oxoid, United Kingdom). Under these conditions, the morphology of the parasites is in the vacuolar state. Subculturing of parasites was performed at intervals of 3 or 4 days. Cultures harvested from log-phase in vitro cultivation at 24 hours postincubation were washed thrice in phosphate-buffered saline (PBS) at $500 \times g$ for 15 min at room temperature (23). The pellet was resuspended in PBS, and cells were counted with a hemocytometer. Parasitic lysates were then prepared by three freeze-thaw cycles in liquid nitrogen and a 37°C water bath. Saccharomyces boulardii was obtained from the ATCC (MYA-796), and cells were grown in 10 ml YPDA (1% [wt/vol] yeast extract, 2% [wt/vol] peptone, 2% [wt/vol] dextrose, 0.04% [wt/vol] adenine) liquid medium inside an incubator shaker at 28°C and 200 rpm. Cells were harvested at log phase by centrifugation at 800 \times g for 5 min and washed with 1 ml sterile water. The pellet was resuspended in 1 ml yeast lysis buffer (100 mM Tris-HCl, pH 7.4, 50 mM KCl, 1 mM EDTA, 0.1% [wt/vol] NP-40). Glass beads (100 mg, acid washed, 425 to 600 µm; Sigma; G8772) were added, and the lysate was prepared with a bead beater (Precellys 24; five times at 6,000 rpm for 30 s with 1 min cooling on ice during the intervals).

Experimental *in vivo* inoculations. C57BL/6 mice were purchased from the National University of Singapore (CARE) unit. All animals were housed in an animal biosafety level 2 (ABSL-2) conventional clean animal facility. The animal experiments were performed in accordance with the Singapore National Advisory Committee for Laboratory Animal Research guidelines. The protocol (R13-5890) was reviewed and approved by the National University of Singapore Institutional Animal Care and Use Committee.

Eight-week-old C57BL/6 male mice were first given 2% dextran sulfate sodium salt (DSS) in drinking water for 5 days (day 1 to day 5). On day 5, the mice were divided into two groups. One group (the inoculated mice) were inoculated with 3.75×10^7 ST-7 (B) lysates by oral gavage for 3 days. The other group, the control mice, were given PBS. Both infected mice and control mice were anesthetized and killed for examination. The small intestine, cecum, proximal colon, and distal colon were harvested to investigate the mRNA expression of the proinflammatory cytokines using real-time reverse transcription (RT)-PCR.

Culture of the RAW264.7 mouse macrophage line. The murine macrophage line RAW264.7 was purchased from the American Type Culture Collection (ATCC TIB-71). RAW264.7 stock cultures were maintained in T-75 flasks (Fisher Scientific) in a humidified 37°C incubator with 5% CO₂, and cultures within 5 to 20 passages were used for this study. RAW264.7 cells were cultured in RPMI 1640 medium with L-glutamine (Invitrogen) supplemented with 10% heat-inactivated fetal bovine serum (FBS) (Invitrogen) and 1% penicillin-streptomycin mixture (Invitrogen) (RPMI 1640 complete medium). Cell viability was assessed by trypan blue assay, and cell cultures with more than 95% viability were used for all experiments. Cells were trypsinized with 0.25% trypsin-EDTA (Invitrogen), seeded in either 6-well plates (Nunc) for Western blot experiments or 12-well plates (Nunc) for real-time PCR and enzyme-linked immunosorbent assays (ELISAs), and grown to 80 to 90% confluence.

For MAPK inhibition experiments, RAW264.7 cells were pretreated in RPMI 1640 complete medium containing one of the following inhibitors: PD98059 (Sigma) (100 μ M), SB203580 (Biorbyt) (20 μ M), or SP600125 (Calbiochem) (20 μ M). After 1 h pretreatment, the RAW264.7 cells were stimulated with ST-7 (B) lysates for the indicated times.

Western blot analysis. After the desired period of coincubation at 37°C in 5% CO₂, the stimulation reaction was ended by removing the RPMI 1640 complete medium, and the monolayers were washed twice with PBS. The cells were then harvested with a cell scraper and spun down at 4°C for 3 min. The supernatant was discarded, while the remaining cell pellet was lysed by adding 150 to 250 pg/ml of cell lysis buffer (50 mM Tris-HCl [pH 7.8], 150 mM NaCl, 1% NP-40) adjusted to the pellet size. The protein concentration of whole-cell lysates was quantitated by spectrophotometric analysis using Bradford reagent (Bio-Rad), according to the manufacturer's instructions. Equal amounts of proteins (30 to 60 µg) were then loaded on 10% polyacrylamide gels for sodium dodecyl sulfatepolyacrylamide gel electrophoresis (SDS-PAGE) using the Mini-Protean 3 Electrophoresis System (Bio-Rad Laboratories) and transferred to a 0.2-µm polyvinylidene difluoride (PVDF) membrane (Bio-Rad Laboratories). After being blocked with 5% nonfat milk in 20 mM Tris-HCl, pH 8.0, 154 M NaCl, 0.05% Tween, the membrane was incubated in primary MAPK antibody at 4°C overnight, followed by incubation with Amersham ECL anti-rabbit IgG and horseradish peroxidase-linked species-specific whole antibody (from donkey; GE Healthcare) for 30 min. The primary MAPK antibodies used were phospho-p44/42 MAPK (Erk1/2) (Thr202/Tyr204) rabbit monoclonal antibody (Cell Signaling Technology), phospho-SAPK/JNK (Thr183/Tyr185) rabbit monoclonal antibody (Cell Signaling Technology), and phospho-p38 MAPK (Thr180/Tyr182) rabbit monoclonal antibody (Cell Signaling Technology). These antibodies were raised against human MAP kinases using epitopes conserved in their murine orthologs. The blots were treated for chemiluminescence detection using Supersignal West Dura Extended Duration Substrate (Pierce). Luminescence was detected using Amersham hyperfilm MP enhanced chemiluminescence (ECL) autoradiography film (GE Healthcare) and processed using an X-ray film processor (Kodak).

ELISA and quantitative real-time PCR (qRT-PCR) for cytokines. An ELISA kit (BD Pharmingen) was used to detect the levels of proinflammatory cytokines, IL-6, IL-1 β , and TNF- α , in supernatants of RAW264.7 cultures coincubated with isolate ST-4 (WR-1) or ST-7 (B) parasite lysates. For live-parasite infection, ST-4 (WR-1) and ST-7 (B) log-phase cultures were harvested and washed twice with PBS at 500 \times g for 15 min at room temperature. The pellet was resuspended in PBS and counted with a hemocytometer. Multiplicities of infection (MOIs) of 20 for ST-4 (WR-1) and 10 for ST-7 (B) were used for infection. For real-time PCR, total cellular RNA was isolated from cells using TRIzol reagent (Invitrogen). First-strand cDNA was prepared using the ImProm-II Reverse Transcription System (Promega) following the manufacturer's protocol. Quantitative real-time PCR was performed using Fast-SYBR green master mix (ABI Applied Biosystems), cDNA, and appropriate primers (see below). Real-time PCR was then run in a 7900HT Fast real-time PCR system at 95°C for 30 s, followed by 40 cycles of 5 s at 95°C and 20 s at 60°C. Melting-curve analysis was conducted after each reaction and examined for a single peak to ensure specificity of the amplification reactions. Relative quantitation of mRNA levels was calculated by the $\Delta\Delta C_T$ method, and the amount of the target relative to glyceraldehyde 3-phosphate dehydrogenase (GAPDH) mRNA was expressed as $2^{-\Delta\Delta CT}$. The primers used were IL-6 sense (5' GAT GCT ACC AAA CTG GAT ATA ATC 3') and antisense (5' GGT CCT TAG CCA CTC CTT CTG TG 3'), IL-1ß sense (5' CCT TCA TCT TTG AAG AAG A 3') and antisense (5' GAG GTG CTG ATG TAC CAG TTG 3'), TNF-α sense (5' TCC CAG GTT CTC TTC AAG GGA 3') and antisense (5' GGT GAG GAG CAC GTA GTC GG 3'), and GAPDH sense (5' GAG AAC TTT GGC ATT GTG G 3') and antisense (5' ATG CAG GGA TGA TGT TCT G 3').

RESULTS

Blastocystis induces proinflammatory cytokine expression in the intestine. The intestinal epithelium provides nonspecific defenses against potential pathogens, and enterocytes secrete cytokines, such as IL-1 β , to regulate inflammation (37). To investigate the possible role of *Blastocystis* infection in intestinal inflamma-

tion, we treated intestinal explants isolated from C57/BL6 mice with *Blastocystis* ST-7 (B) lysate to test its proinflammatory potential. We found that treatment with ST-7 (B) resulted in secretion of IL-1 β and TNF- α from the explants (Fig. 1A). To confirm that the inflammatory cytokine expression in intestinal explants was indeed due to *Blastocystis*, we treated the explants with live ST-7 (B) and found that live ST-7 (B) treatment induced the secretion of IL-1 β and IL-6 (see Fig. S2 in the supplemental material). Together, these results suggested that infection with *Blastocystis* could contribute to acute and chronic inflammatory intestinal diseases.

To examine the possible contribution of Blastocystis to intestinal inflammation, we treated C57/BL6 mice for 5 days with 2% DSS, a heparin-like polysaccharide that induces intestinal mucosal damage, exposing the underlying epithelium to the luminal contents, including microbes, and that therefore is commonly used to induce mouse colitis (1), followed or not by treatment with ST-7 (B) lysate. The expression of inflammatory cytokines, including IL-1 β , IL-6, and TNF- α , in the colon was examined. We observed an increase in the average expression of IL-1B, IL-6, and TNF- α in response to ST-7 (B) treatment in each of the three experiments we performed (see Fig. S3 in the supplemental material). However, the differences between treated and nontreated groups were not significant. It is possible that the effect of Blasto*cystis* treatment was overshadowed by the existing high expression of these cytokines caused by DSS, indicating a suitable mouse model for Blastocystis infection is needed.

Blastocystis induces MAP kinase activation in macrophages. It has been shown that infection of mice by *Blastocystis* ST-7 (B) resulted in intensive proinflammatory cell infiltration and inflammation in the colon (20). Macrophages, one of the major components of innate immunity, are the first cells that encounter pathogens that cross the intestinal epithelium (38). To examine macrophage responses to *Blastocystis* infection, we generated mouse bone marrow-derived macrophages (BMDMs) and treated them with *Blastocystis* ST-4 (WR-1) or ST-7 (B) lysates. We found that ST-7 (B) treatment induced strong expression of proinflammatory cytokines, including IL-1 β , IL-6, and TNF- α , at both the mRNA and protein levels in macrophages compared with cells treated with ST-4 (WR-1) (Fig. 1B and C).

To facilitate the study of the role of Blastocystis infection in intestinal inflammation, particularly in proinflammatory cytokine expression by macrophages, we utilized the murine macrophage line most commonly used in medical research, RAW264.7. We treated RAW264.7 macrophages with ST-4 (WR-1) or ST-7 (B) lysate to examine proinflammatory cytokine expression. As shown in Fig. 2A and B, both ST-4 (WR-1) and ST-7 (B) induced the expression of IL-1 β , IL-6, and TNF- α . However, ST-7 (B) induced significantly higher expression of these cytokines in macrophages than ST-4 (WR-1), which is consistent with our findings in BMDMs (Fig. 1B and C). Similar results were observed when RAW264.7 cells were treated with live ST-4 (WR-1) and ST-7 (B) parasites (see Fig. S4 in the supplemental material). Of note, the lysate of S. boulardii, a strain of yeast, does not induce significant expression of IL-1 β , IL-6, and TNF- α from macrophages (Fig. 2C). This result demonstrates that the inflammatory cytokine expression by macrophages in response to ST7 (B) lysate is indeed caused by factors from Blastocystis ST-7 (B).

Signaling pathways such as the MAP kinases (MAPKs) play a critical role in regulating the expression of proinflammatory cyto-

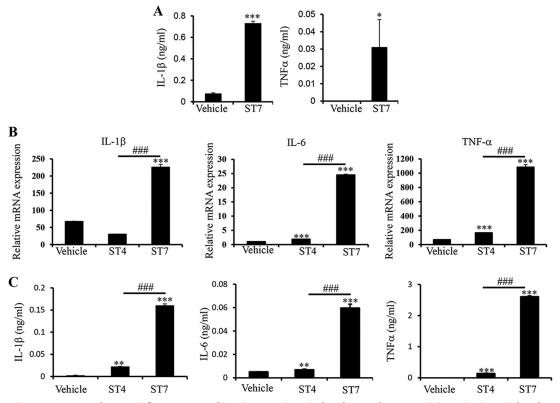


FIG 1 *Blastocystis* treatment upregulates proinflammatory cytokines in mouse intestinal explants and BMDMs. (A) Murine intestinal explants were isolated from C57/BL6 mice and treated with *Blastocystis* ST-7 (B) lysate. The culture supernatants were tested using ELISA for protein concentrations of proinflammatory cytokines, IL-1β and TNF- α . (B) Mouse BMDMs were generated and treated with either *Blastocystis* ST-4 (WR-1) or ST-7 (B) at MOIs of 20 and 10, respectively. The mRNA levels of TNF- α , IL-1β, IL-6, and GAPDH were also measured by qRT-PCR. (C) The concentrations of IL-6, TNF- α , and IL-1β in the culture supernatants of BMDMs were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, *P* < 0.05; **, *P* < 0.005 (representing a statistically significant difference compared to the negative control). ###, *P* < 0.005 (representing a statistically significant difference between *Blastocystis* ST-4 and ST-7).

kines like IL-1 β , IL-6, and TNF- α (39). To investigate the potential role of MAPKs in the expression of proinflammatory cytokines by macrophages in response to Blastocystis, RAW264.7 cells were treated with ST-4 (WR-1) or ST-7 (B) to assess the activation of MAPKs. As shown in Fig. 2D, ST-4 (WR-1) induced ERK activation at 15 min posttreatment, and activation peaked at 30 min. ST-7 (B) induced stronger ERK activation at 15 and 30 min posttreatment, which then decreased at 3 h. Also, ST-7 (B) induced much stronger JNK activation in RAW264.7 that peaked within the first hour and was sustained until 6 h after treatment. Peak phosphorylation of p38 was seen in cells at 30 min when treated with ST-4 (WR-1) or ST-7 (B). Evidently, ST-7 (B) induced much stronger activation of all three MAPKs than ST-4 (WR-1). These results demonstrated the different abilities of these two subtypes to activate the three MAPK pathways, which is correlated with their abilities to induce proinflammatory cytokine expression in macrophages. In the studies described below, we focused on investigating the regulatory function of MAPKs in macrophage proinflammatory cytokine expression induced by ST-7 (B).

Blastocystis-induced proinflammatory cytokine expression is dependent on ERK and JNK. Next, we examined the roles of the three major groups of MAPKs in the expression of proinflammatory cytokines in macrophages in response to *Blastocystis*, using their specific inhibitors. To elucidate the role of ERK in macrophage cytokine expression in response to *Blastocystis*, we treated RAW264.7 cells with PD98059, an MEK1-specific inhibitor. Pretreatment of macrophages with PD98059 for an hour reduced ERK activation upon ST-7 (B) stimulation (Fig. 3A). Treatment with PD98059 did not affect the activation of p38 and JNK in macrophages in response to ST-7 (B) stimulation (Fig. 3A).

Next, we examined the effect of ERK inhibition on the expression of IL-1 β , IL-6, and TNF- α in macrophages in response to *Blastocystis* by qRT-PCR and ELISA. As shown in Fig. 3B, inhibition of ERK significantly impaired the expression of IL-1 β , IL-6, and TNF- α at the mRNA level. Notably, inhibition of ERK more profoundly suppressed the protein production of IL-6 and TNF- α than that of IL-1 β (Fig. 3C). Together, these data indicated that ERK plays a major role in regulating the expression of IL-6 and TNF- α by macrophages in response to *Blastocystis* infection.

To understand the contribution of JNK to the expression of proinflammatory cytokine expression upon *Blastocystis* treatment, RAW264.7 macrophages were pretreated with SP100625, a JNK-specific inhibitor. As shown in Fig. 4A, inhibition of JNK activation in macrophages led to a decrease in activation of both JNK and its downstream target, c-Jun. At the same time, ERK activation was also completely abrogated in response to SP100625 treatment, whereas p38 activation was not affected. A significant reduction of IL-1 β , IL-6, and TNF- α in response to *Blastocystis* stimulation was observed at both the mRNA and protein levels (Fig. 4B and C). Interestingly, SP100625 treatment resulted in

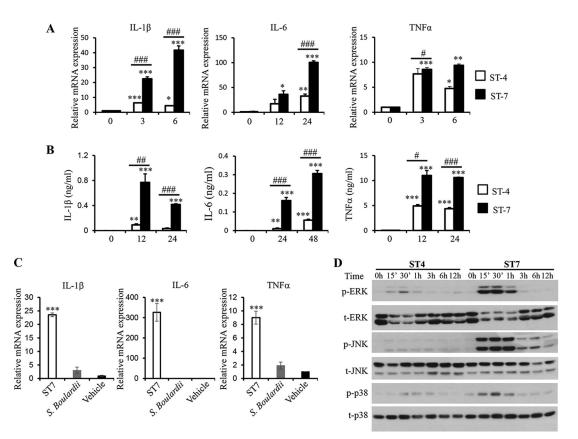


FIG 2 *Blastocystis* ST-7 (B) induces stronger proinflammatory cytokine expression and MAPK activation in murine macrophages than ST-4 (WR1). RAW264.7 cells were incubated with lysates of either *Blastocystis* ST-4 (WR1) or ST-7 (B) at MOIs of 20 and 10, respectively. (A) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH in macrophages were measured by qRT-PCR. The results are normalized to GAPDH and expressed as the mean fold change relative to the 0-h time point, which is set at 1. (B) The concentrations of IL-6, TNF- α , and IL-1 β in the culture supernatants of RAW264.7 cells were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, P < 0.05; **, P < 0.01; ***, P < 0.005 (representing a statistically significant difference compared to the negative control). #, P < 0.05; ##, P < 0.005 (representing a statistically significant difference between *Blastocystis* ST-4 and ST-7). (C) *S. boulardii* lysate does not induce inflammatory cytokine expression in macrophages. RAW264.7 cells were incubated with lysate of either *Blastocystis* ST-7 (B) (MOI, 10) or *S. boulardii* (MOI, 20). The mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH in macrophages were measured by qRT-PCR. (D) Representative Western blots of MAPK activation in RAW264.7 cells in response to ST-4 (WR1) or ST-7 (B) lysate stimulation. Cells were lysed and were processed by SDS-PAGE. Membranes were then blotted with phospho-ERK (p-ERK), total ERK (t-ERK), phospho-JNK, total JNK, phospho-p38, and total p38 antibodies. Each blot is representative of three separate experiments.

reduced expression of IL-1 β at both the mRNA and protein levels. These results showed the important regulatory role of JNK and ERK pathways for these proinflammatory cytokines in response to *Blastocystis* infection.

p38 MAP kinase has been shown to be important for proinflammatory cytokine expression in response to microbial infection (39). To address whether p38 regulates proinflammatory cytokine expression by macrophages in response to Blastocystis, we treated RAW264.7 cells with the p38-specific inhibitor SB203580, which blocks p38 catalytic activity without inhibiting its phosphorylation prior to stimulation with ST-7 (B). As shown in Fig. 5A, treatment with SB203580 resulted in reduced phosphorylation of ATF2, the substrate of p38. Meanwhile, an increase in ERK and JNK phosphorylation was observed after the treatment, suggesting cross talk by p38 with ERK and JNK in macrophages. At the same time, inhibition of p38 did not suppress mRNA expression of IL-1 β , IL-6, and TNF- α in response to *Blastocystis* stimulation or their protein production, except for IL-6 (Fig. 5B and C). This is consistent with the observation that ERK and JNK are required for mRNA expression of these cytokines in response to

Blasotcystis (Fig. 4B and C). Together, these results demonstrated that ERK and possibly JNK activation triggered by *Blastocystis* infection play major roles in regulating the expression of proinflammatory cytokines, including IL-1 β , IL-6, and TNF- α .

Blastocystis-derived serine proteases play a major role in inducing MAPK activation and proinflammatory cytokine expression in macrophages. Proteases produced by parasitic protozoa contribute to their pathogenicity in many ways, ranging from invasion of the host to immune evasion and activation of inflammation (40). Proteases expressed by *Blastocystis* were able to degrade human secretory IgA and induce IL-8 expression in human colonic epithelial cells, indicating their roles in the pathogenesis of Blastocystis (19, 41). We examined the role of Blastocystis proteases in the regulation of proinflammatory cytokine expression in macrophages by incubating ST-7 (B) lysates with serine or cysteine protease inhibitors before macrophage stimulation. ST-7 (B) treated with the serine protease inhibitor 4-(2-aminoethyl)benzenesulfonyl fluoride hydrochloride (AEBSF) induced weaker activation of ERK in RAW264.7 cells than ST-7 (B) treated with vehicle, whereas the activation of JNK and p38 was not reduced

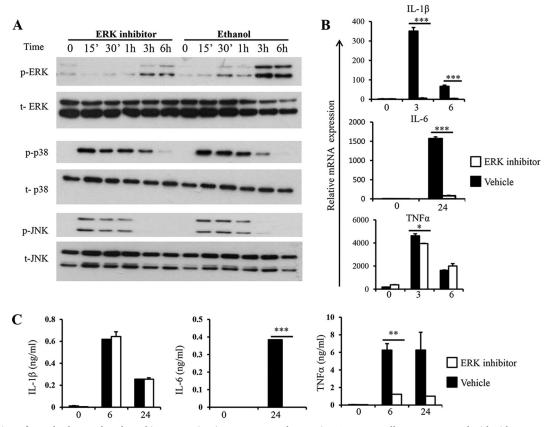


FIG 3 Inhibition of ERK leads to reduced cytokine expression in response to *Blastocystis*. RAW264.7 cells were pretreated with either 100 μ M PD98059 or ethanol vehicle dissolved in RPMI complete medium for 1 hour prior to stimulation with *Blastocysis* ST-7 at an MOI of 10 for the indicated times. (A) Representative Western blots for MAPK activation in RAW264.7 cells. The cells were lysed and processed by SDS-PAGE. Membranes were then blotted with phospho-ERK, ERK, phospho-JNK, JNK, phospho-p38, and p38 antibodies. Each blot is representative of three separate experiments. (B) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH were measured by qRT-PCR. The values are expressed as mean relative transcript levels normalized to GAPDH and are expressed relative to the 0-h time point, which is set at 1. (C) Concentrations of IL-6, TNF- α , and IL-1 β in the culture supernatants were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, *P* < 0.05; **, *P* < 0.01; ***, *P* < 0.005 (representing a statistically significant difference between inhibitor and vehicle treatments).

(Fig. 6A). In correlation with reduced ERK activation, RAW264.7 cells stimulated with AEBSF-treated ST-7 (B) produced significantly smaller amounts of IL-6, TNF- α , and IL-1 β (Fig. 6B). In contrast, treatment of cysteine protease inhibitor with ST-7 (B) induced activation of MAPKs in RAW264.7 cells similar to that induced by the control (Fig. 7A). IL-6 mRNA and protein secretion levels were not affected by the cysteine protease inhibition. Interestingly, inhibiting cysteine proteases in ST-7 (B) resulted in significantly lower levels of IL-1 β and TNF- α mRNA expression (Fig. 7B). However, protein expression of IL-1 β and TNF- α was not reduced (Fig. 7B). Together, these results suggested that serine proteases produced by Blastocystis play a critical role in the induction of MAPK activation and proinflammatory cytokine expression in macrophages and that signaling triggered by cysteine proteases produced by Blastocystis plays a role in the mRNA expression of IL-1 β and TNF- α that is independent of MAPKs.

DISCUSSION

Blastocystis, the most common single-cell parasite in the human large intestine, is associated with various human intestinal disorders, such as acute and chronic diarrhea, irritable bowel syndrome, and possibly inflammatory bowel disease (3, 42, 43). How-

ever, the pathogenic potential of *Blastocystis* remains unclear. Previous studies showed that *Blastocystis* infection led to upregulation of proinflammatory gene expression and intense infiltration of proinflammatory cells into the colon in rats and mice (20, 21). In agreement with these findings, in this study, we found that ST-7 (B) lysate was able to induce inflammatory cytokine secretion in mouse colon explants (Fig. 1A). We further showed that in mice with colitis, treatment with ST-7 (B) lysate resulted in increased inflammatory cytokine gene expression, including the expression of IL-6, TNF- α , and IL-1 β genes (Fig. 1B), suggesting the inflammatory potential of *Blastocystis* infection in the intestine.

Proinflammatory cytokines, such as IL-1, IL-8, gamma interferon (IFN-γ), and TNF-α, secreted by epithelial cells or immune cells, including macrophages, monocytes, neutrophils, and lymphocytes, contribute to mucosal intestinal inflammation and inflammatory disorders (44, 45). Extensive reports have shown that common protozoan pathogens, such as *T. cruzi*, *E. histolytica*, *T. vaginalis*, and *Leishmania*, cause upregulation of proinflammatory cytokines to induce immune responses (46–49). Macrophages, the first cells that encounter pathogens that cross the intestinal epithelium, may be resident in strategic locations, such as the lamina propria, or may be recruited to the inflamed gut by

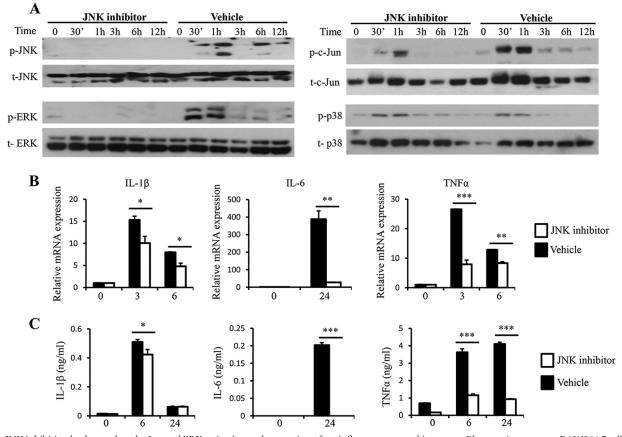


FIG 4 JNK inhibition leads to reduced c-Jun and ERK activation and expression of proinflammatory cytokines upon *Blastocystis* treatment. RAW264.7 cells were pretreated with either 20 μ M SP100625 or DMSO (dimethyl sulfoxide) vehicle dissolved in RPMI complete medium for 1 hour prior to stimulation with *Blastocystis* ST-7 (B) at an MOI of 10. (A) Representative Western blots for activation of c-Jun and MAPKs in RAW264.7 cells. The cells were lysed and processed by SDS-PAGE. Membranes were then blotted with phospho-ERK, ERK, phospho-JNK, JNK, phospho-cJun, c-Jun, phospho-p38, and p38 antibodies. Each blot is representative of three separate experiments. (B) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH were measured by qRT-PCR. The values are expressed as mean relative transcript levels normalized to GAPDH and are expressed relative to the 0-h time point, which is set at 1. (C) Concentrations of IL-6, TNF- α , and IL-1 β in the supernatants of RAW264.7 cells were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, *P* < 0.05; **, *P* < 0.01; ***, *P* < 0.005 (representing a statistically significant difference between inhibitor and vehicle treatments).

chemokine signals (38, 50). Blastocystis has been reported to disrupt intestinal barrier integrity (17, 51). Such tissue damage would presumably allow Blastocystis antigens to cross the epithelium into the lamina propria and contact cells of the immune system. Different types of macrophages with different features and functions exist in the gut, playing important roles in the regulation of the immune response to commensal microorganisms, as well as contributing to the pathogenesis of intestinal inflammatory disorders (50). For instance, the CD14⁺ inflammatory macrophages in the intestine can produce large amounts of proinflammatory cytokines, including IL-23, IL-6, and TNF-α, in response to commensal bacteria to contribute to the intestinal inflammation and pathogenesis of IBD (50, 52). We showed that macrophages treated with Blastocystis ST-4 (WR-1) or ST-7 (B) resulted in upregulation of the proinflammatory cytokines IL-6, IL-1β, and TNF- α (Fig. 1 and 2). In addition, ST-7 (B) evoked a stronger inflammatory response than ST-4 (WR-1), demonstrating the different pathogenic potentials of the two subtypes.

Interestingly, the different abilities of ST7 (B) and ST4 (WR-1) to induce the expression of proinflammatory cytokines in macrophages were correlated with their abilities to induce the activation of MAPKs (Fig. 2). MAPKs are known to play crucial roles in the regulation of cytokine expression in immune responses (28). While the kinetics of MAPK activation induced by the two subtypes were similar, the intensity of phosphorylation of each MAPK activated by ST-7 (B) was distinctly stronger than that by ST-4 (WR-1).

We further examined the contributions of individual MAPK pathways to the expression of these cytokines in macrophages in response to Blastocystis using specific inhibitors. The results suggested that the ERK and possibly JNK pathways play the main roles in regulating the expression of IL-6, IL-1 β , and TNF- α . Interestingly, inhibition of p38 activation resulted in an increase in mRNA expression of IL-1 β , IL-6, and TNF- α , which was correlated with increased JNK and ERK activation (Fig. 5A and B), suggesting cross talk among the three MAPK signaling pathways and supporting the major role of ERK and JNK in the expression of IL-1 β , IL-6, and TNF- α in macrophages in response to *Blasto*cystis. While mechanisms are in place to ensure the specificity of each MAPK pathway, studies have shown that cross talk does occur among different signaling pathways when inhibition of one MAPK pathway augments the phosphorylation of other MAPKs (53, 54).

The pathogenic status of Blastocystis remains controversial due

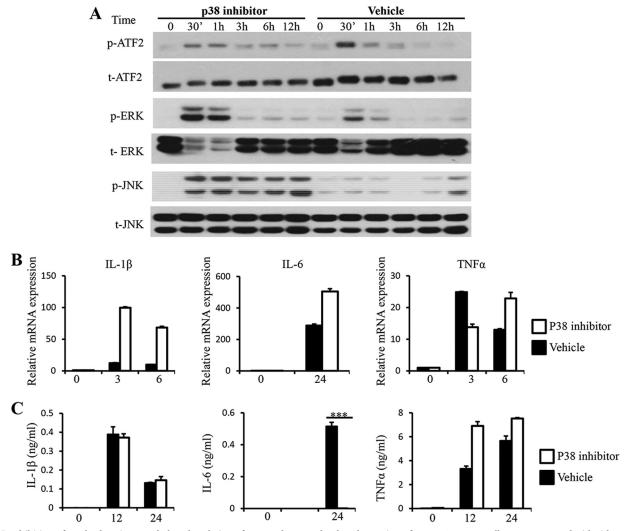


FIG 5 Inhibition of p38 leads to increased phosphorylation of ERK and JNK and reduced secretion of IL-6. RAW264.7 cells were pretreated with either 20 μ M SB203580 or DMSO vehicle dissolved in RPMI complete medium for 1 hour prior to stimulation with *Blastocystis* ST-7 at an MOI of 10. (A) Representative Western blots for activation of ATF2, ERK, and JNK. Cells were lysed and processed by SDS-PAGE. Membranes were then blotted with phospho-ERK, ERK, phospho-JNK, JNK, phospho-ATF2, and ATF2 antibodies. Each blot is representative of three separate experiments. (B) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH were measured by qRT-PCR. The values are expressed as mean relative transcript levels normalized to GAPDH and are expressed relative to the 0-h time point, which is set at 1. (C) Concentrations of IL-6, TNF- α and IL-1 β in the culture supernatants were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. ***, P < 0.005 (representing a statistically significant difference between inhibitor and vehicle treatments).

to lack of knowledge about virulence factors and the existence of intrasubtype variation in pathogenic potential (55). Previous studies suggested that aspartic and cysteine proteases are candidates for *Blastocystis* virulence factors. Proteases derived from parasitic protozoa are major virulence factors that contribute to their pathogenicity by manipulating the immune response in a number of ways, such as tissue invasion, immune evasion, and activation of inflammation (40). Genome sequencing of ST-7 (B) has revealed the major classes of proteolytic enzymes to be serine, aspartic, and cysteine proteases and metalloproteases (56). It has been shown that aspartic and cysteine proteases from *Blastocystis* are able to degrade human secretory IgA, while cysteine proteases could reduce epithelial barrier function and induce IL-8 expression (19, 41, 57). In this study, we found that serine proteases, but not cysteine proteases, play a major role in induction of MAPK

activation and expression of proinflammatory cytokines in macrophages (Fig. 6 and 7). Therefore, serine proteases are possible virulence factors of *Blastocystis*, as well. Serine proteases have been shown to play a role in the viability and pathogenicity of protozoan parasites, such as *Plasmodium falciparum*, *Leishmania amazonensis*, and *T. cruzi*, and have been implicated in gastrointestinal symptoms, such as abdominal aches, muscular contractions, and widespread pain (58–60). Several reports have established a link between *Blastocystis* and IBS after finding a higher prevalence of the intestinal parasite in IBS patients (14, 15, 61). Fecal supernatants collected from IBS patients were found to have significantly higher serine protease activity than those from healthy subjects (62). This elevated serine protease activity in fecal supernatants resulted in an increase in colonic paracellular permeability *in vitro* and evoked visceral hypersensitivity in mice after intracolonic in-

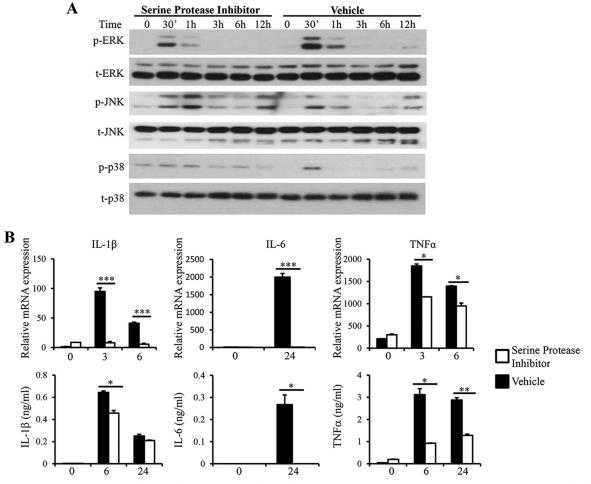


FIG 6 Serine protease inhibition leads to decreased ERK activation and proinflammatory cytokine expression. ST-7 (B) lysates were pretreated with 500 μ M serine protease inhibitor, AEBSF, for 1 hour before coincubation with RAW264.7 cells at 37°C. (A) Representative Western blots for activation of MAPKs. Cells were lysed, and cell extracts were analyzed by Western blotting using phospho-ERK, total ERK, phospho-JNK, total JNK, phospho-p38, and total p38 antibodies. (B) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH were measured by qRT-PCR. The values are expressed as mean relative transcript levels normalized to GAPDH and are expressed relative to the 0-h time point, which is set at 1. (C) Concentrations of IL-6, TNF- α , and IL-1 β in the supernatants were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, P < 0.05; **, P < 0.01; ***, P < 0.005 (representing a statistically significant difference between inhibitor and vehicle treatments).

fusion. *Blastocystis* has been reported to induce contact-independent apoptosis and to disrupt barrier function in rat intestinal epithelial IEC-6 cells (17). It would be interesting to investigate if *Blastocystis* serine protease activity is associated with increased intestinal epithelial permeability in future studies.

The link between *Blastocystis* serine protease, MAPK activation, and inflammation is currently unclear. Macrophages express a group of receptors known as protease-activated receptors (PARs) (63), a family of G-protein-coupled seven-transmembrane receptors (64), on the surface. The activation of PARs is initiated by cleavage of the N terminus of the receptor by a serine protease, resulting in the generation of a new tethered ligand that interacts with the receptor within extracellular loop 2, leading to downstream signaling. PAR2 is one of the four members in this family. PAR2 has been shown to be activated by proteases from enteric bacteria, such as *Serratia marcescens* (65), or proteases from the host in response to bacterial infection, such as infection by *Citrobacter rodentium* (66), playing a major role in the host inflammatory response to enteric bacterial infection (65, 66). PAR2 activation induces MAPK signaling, including ERK1/2 and to a lesser extent p38 and JNK (67). We found that ST-7 (B) lysate-induced ERK activation and inflammatory cytokine expression was reduced by treatment with serine protease inhibitor (Fig. 6). It is possible that serine protease from *Blastocystis* activates PAR2 in macrophages, which triggers MAPK activation and inflammatory cytokine expression. We will further validate this possibility in future investigations.

The different abilities of ST-7 (B) and ST-4 (WR-1) to trigger MAPK activation and proinflammatory cytokine expression could be due to their differential expression of serine proteases or different serine protease activities. Previous studies have shown such intrasubtype variation of virulence potential. For instance, cysteine protease activity from ST-7 (B) exhibited approximately twice the level of cysteine protease activity from ST-4 (WR-1) (23). In addition, ST-7 (B) was found to be resistant to metronidazole, a first-line therapy for *Blastocystis* infections, while ST-4 (WR-1) was susceptible to the drug (68). Despite being more sensitive to NO production by intestinal epithelial cells, ST-7 (B) also exhibited an ability to downregulate iNOS, indicating it possesses

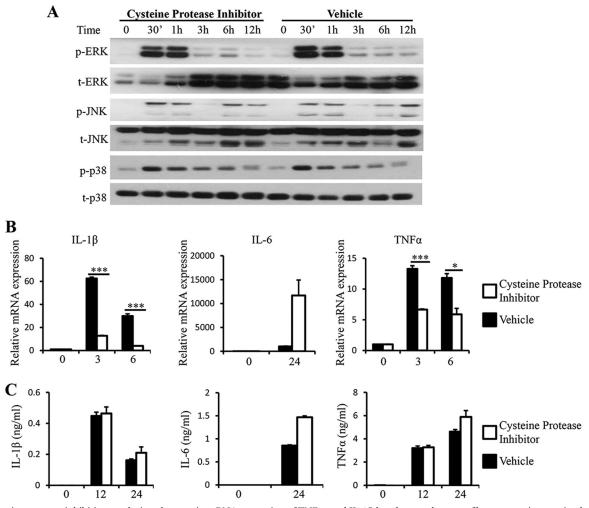


FIG 7 Cysteine protease inhibition results in a decrease in mRNA expression of TNF- α and IL-1 β but does not have an effect on protein secretion levels for all three proinflammatory cytokines examined. ST-7 (B) lysates were pretreated with 100 μ M cysteine protease inhibitor, E-64, for 1 hour before coincubation with RAW264.7 cells at 37°C. (A) Representative Western blots for activation of MAPKs. Cells were lysed, and cell extracts were analyzed by Western blotting using phospho-ERK, total ERK, phospho-JNK, total JNK, phospho-p38, and p38 antibodies. (B) mRNA levels of TNF- α , IL-1 β , IL-6, and GAPDH were measured by qRT-PCR. The values are expressed as mean relative transcript levels normalized to GAPDH and are expressed relative to the 0-h time point, which is set at 1. (C) Concentrations of IL-6, TNF- α , and IL-1 β in the supernatants were measured by ELISA. Each value represents the mean and standard deviation from at least 3 individual experiments. *, *P* < 0.005 (representing a statistically significant difference between inhibitor and vehicle treatments).

unique strategies to evade potential host defenses (24). Together, this evidence suggested that ST-7 (B) is more pathogenic than ST-4 (WR-1). Recent studies have implicated ST-7 (B) as a pathogenic subtype associated with intestinal symptoms, as it was found to be more prevalent in symptomatic patients than in healthy carriers (69). In addition, the prevalence of *Blastocystis* spp. is significant in IBS patients, and inflammation of the mucosa is well recognized in these patients (70). The significantly higher induction of proinflammatory cytokine production by ST-7 (B) found in this study may suggest a contribution by the parasite to the immunopathology of symptomatic patients seen in the clinical setting and supports the pathogenic nature of ST-7 (B) reported in clinical manifestations.

The pathogenic potential of ST-4 (WR-1) also should not be underestimated due to the increasing number of reports associating the subtype with symptomatic infections. Subtype analysis done on 51 clinical isolates has revealed a high prevalence of ST-4 (WR-1) among symptomatic patients from a health district of Valencia, Spain (7). ST-4 (WR-1) was revealed to be predominant in rural communities of Nepal and was also found to be common among *Blastocystis*-positive patients presenting with acute diarrhea in Denmark (71, 72). Oral inoculation of rats with ST-4 (WR-1) resulted in mild goblet cell hyperplasia and significant upregulation of proinflammatory cytokines (21). With these observations showing the pathogenic potential of both ST-4 (WR-1) and ST-7 (B), the results from our study suggest that while ST-4 (WR-1) could induce significant upregulation of proinflammatory cytokines, comparisons between the two subtypes have shown that ST-7 (B) could induce stronger upregulation of proinflammatory cytokines, which suggests that it could be more virulent. Future clinical studies can attempt to determine whether ST-7 (B) is associated with the marked immunopathology seen in symptomatic patients compared to ST-4 (WR-1).

In summary, *Blastocystis* ST-4 (WR-1) and ST-7 (B) exhibit subtype-dependent upregulation of proinflammatory cytokines and MAPK activation in macrophages. This is the first study that demonstrates differential regulation of proinflammatory cytokines by MAPKs in response to *Blastocystis* in macrophages. The present study also demonstrates for the first time that *Blastocystis* ST-7 (B) serine proteases play a major role in MAPK activation and the expression of proinflammatory cytokines, including IL-6, IL-1 β , and TNF- α , supporting serine proteases as virulence factors of *Blastocystis*. While these findings shed new light on the pathogenesis of *Blastocystis*, future studies should focus on the development of suitable animal models to gain further understanding of *Blastocystis*-host interactions.

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