

Salmonella Outer Protein B Suppresses Colitis Development via Protecting Cell From Necroptosis

Gui-Qiu Hu^{\dagger}, Yong-Jun Yang^{\dagger}, Xiao-Xia Qin, Shuai Qi, Jie Zhang, Shui-Xing Yu, Chong-Tao Du and Wei Chen^{*}

Key Laboratory of Zoonosis Research, Ministry of Education, College of Veterinary Medicine, Jilin University, Changchun, China

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> *Correspondence: Wei Chen chw_cc@jlu.edu.cn

[†]These authors have contributed equally to this work

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Hu G-Q, Yang Y-J, Qin X-X, Qi S, Zhang J, Yu S-X, Du C-T and Chen W (2019) Salmonella Outer Protein B Suppresses Colitis Development via Protecting Cell From Necroptosis. Front. Cell. Infect. Microbiol. 9:87. doi: 10.3389/fcimb.2019.00087 Salmonella effectors translocated into epithelial cells contribute to the pathogenesis of infection. They mediate epithelial cell invasion and subsequent intracellular replication. However, their functions in vivo have not been well-identified. In this study, we uncovered a role for Salmonella outer protein B (SopB) in modulating necroptosis to facilitate bacteria escape epithelial cell and spread to systemic sites through a Salmonellainduced colitis model. Mice infected with SopB deleted strain $\Delta sopB$ displayed increased severity to colitis, reduced mucin expression and increased bacterial translocation. In vitro study, we found there was an increased goblet cell necroptosis following $\Delta sopB$ infection. Consistently, mice infected with $\Delta sopB$ had a strong upregulation of mixed lineage kinase domain-like (MLKL) phosphorylation. Deletion of MLKL rescued severity of tissue inflammatory, improved mucin2 expression and abolished the increased bacterial translocation in mice infected with $\Delta sopB$. Intriguingly, the expression of sopB in LS174T cells was downregulated. The temporally regulated SopB expression potentially switched the role from epithelial cell invasion to bacterial transmission. Collectively, these results indicated a role for SopB in modulating the onset of necroptosis to increased bacteria pathogenesis and translocated to systemic sites.

Keywords: Salmonella, SopB, colitis, MLKL, necroptosis

INTRODUCTION

Salmonella enterica is a leading cause of enteric disease in human and animals that is capable of causing a wide range of illnesses ranging from a localized gastroenteritis and lymphadenitis resulting in diarrhea to life-threatening systemic infections (Kirk et al., 2015). Salmonella has evolved many mechanisms to evade host immune response to survive in host cell via virulence effectors (Tobar et al., 2006; McGhie et al., 2009; Raymond et al., 2013). Salmonella has two major virulence determinants called Salmonella pathogenicity island (SPI)-1 and SPI-2 (Hensel, 2004). SPI-1 is essential for invasion of non-phagocytic cells and responsible for inflammation in Salmonella induced colitis, whereas SPI-2 is required for intracellular survival and proliferation in phagocytes, and is important in systemic infection (Barthel et al., 2003; Abrahams and Hensel, 2006). Both SPI-1 and SPI-2 encode separated type III secretion systems (T3SSs) that direct the translocation of virulence proteins into host environment (Galan, 2001; Schmidt and Hensel, 2004). These effector proteins modulate host cell immunity and are important for bacterial pathogenesis.

Salmonella outer protein B (SopB) is SPI-1 encoded protein. Its synthesis is tightly regulated. The long half-life in cells allows SopB to exert multiple roles during Salmonella infection (Giacomodonato et al., 2011). SopB is responsible for membrane ruffle formation and subsequent invasion (Piscatelli et al., 2016). In addition to its role in invasion, numerous biological roles have been explored attributing to its inositol phosphatase activity. For instance, SopB participates in modulation of chloride secretion (Bertelsen et al., 2004) and Akt phosphorylation (Cooper et al., 2011). Studies have shown that SopB protects epithelial cell from apoptosis (Knodler et al., 2005; Ruan et al., 2016). Additionally, SopB mutation mediated increased macrophage death results in enhanced inflammasome activation in inositol phosphatase activity dependent manner (Hu et al., 2017). The role of SopB in protecting cells from death represents a bacterial strategy to reduce host response.

Cells death mediated by Salmonella infection is considered to be an important pathological process and a major cause of tissue damage during infection. Initial studies showed that Salmonella induce cell apoptosis (Knodler et al., 2005), however, recent studies demonstrate that cell death can be mediated through many different ways such as necroptosis and pyroptosis (Broz et al., 2010; Hefele et al., 2018). Apoptosis is a non-lytic programmed cell death which is usually immunological silent, while necroptosis and pyroptosis are lytic cell death and usually result in inflammatory response. Many effector proteins have involved in regulating epithelial cell death which is considered a mechanism used by Salmonella to escape epithelial cell and spread to systemic sites. Previous studies showed SopB protects cell from death in mechanisms of apoptosis and/or pyroptosis (Knodler et al., 2005; Hu et al., 2017), however, the mechanisms of SopB in vivo has not been well-studied.

In this study, we used a *Salmonella* induced colitis model to evaluate the role of SopB in bacterial pathogenesis and employed SopB mutant strain $\Delta sopB$ as well as mixed lineage kinase domain-like (MLKL) deficient mice to elucidate the contribution of SopB in manipulating host immune response during bacterial invasion and dissemination *in vivo*.

MATERIALS AND METHODS

Animals

C57BL/6J (WT) and MLKL deficient (MLKL^{-/-}) mice, on a C57BL/6J background, were bred in house. Mice were maintained under conditions of a 12 h light/dark cycle at 23°C with food and water *ad libitum* in the animal care facility. All animal experiments were performed in accordance with the relevant guidelines and regulations that were approved by the Committee on Animal Care and Use of Jilin University, China.

In vivo Infection

Six- to eight- weeks old and sex-matched mice were used in this study. The *Salmonella*-induced colitis model was established as previously described (Stecher et al., 2005). Briefly, following the administration of 20 mg streptomycin per mouse for 24 h, mice were orally challenged with 5×10^7 colony-forming unit

(CFU) of *Salmonella* strain SL1344 or SopB deleted strain $\triangle sopB$. Bacterial burdens and histopathology of cecum were determined at indicated time point post infection (p.i.).

Cell Culture

LS174T human goblet-like cells were cultured in DMEM/F12 (Gibco, Waltham, MA, USA) containing 10% heat inactivated fetal bovine serum. Cells were seeded in 24-well dish at 0. 3 \times 10⁶ cells per well and cultured overnight to yield monolayers of 80-90% confluence. Cells were pretreated with or without the MLKL inhibitor NSA (Toronto Research Chemical Inc., Toronto, Ontario, Canada) at a final concentration of 2 nM or the receptor-interacting protein kinase (RIP1) inhibitor Nec-1(Tocris, Minneapolis, MN, USA) at a final concentration of 50 µM for 1 h. Salmonella strains were then inoculated to LS174T cells at a multiplicity of infection = 100 for 1 h, then the extracellular bacteria were removed by washing with DMEM/F12. Cells were then maintained in DMEM/F12 containing 200 U/mL streptomycin and 100 µg/mL gentamicin for another 3 h, cell viability was evaluated. For mucin-2 examination, cells were seeded in 6-well dish at 1.5×10^6 cells per well and cultured overnight, then infected with Salmonella strains for 14 and 24 h respectively and lysed in lysis buffer.

Dot Blot Assay

Add 20 ug total protein to NC membranes $(0.45 \,\mu\text{m})$, Merck Millipore, Darmstadt, Germany) and let the membrane dry at room temperature (RT). Blocking non-specific sites by soaking in 5% BSA in Tris buffered saline containing 0.1% (v/v) Tween 20 (Sigma-Aldrich, St. Louis, MO, USA) (TBST) for 2 h at RT and then incubating the membranes with primary antibody anti-Mucin2 (1:500, Santa Cruz, CA, USA) for 1 h at RT. Wash the membrane with TBST for 3 times, each time for 10 min, and then incubate with secondary antibody for 1 h at RT.

Histology and Immunohistochemisty (IHC)

On day 2 p.i., the cecum was harvested and fixed in 4% paraformaldehyde and then embedded in paraffin; $5 \mu m$ sections were used for hematoxylin-eosin (H&E) staining and IHC staining. The assessment of tissue pathology was scored as previously described (Stecher et al., 2005). For IHC staining, sections were dewaxed and rehydrated, and antigen was unmasked in a citrate-containing buffer, and then immersed the sections in 3% hydrogen peroxide for 10 min, blocked the non-specific antigen with 5% goat serum for 1 h at RT, then incubated with the primary antibody anti-Mucin2 (1:100, Santa Cruz, CA, USA) and anti-p-MLKL (1:100, AbCam, Cambridge, MA, USA) at 4°C overnight respectively, then cover sections with detection antibody.

PAS Staining

After IHC staining, sections were oxidized in periodic acid solution (Solarbio, Beijing, China) for 5 min and then rinsed in distilled water. Subsequently, sections were immersed in Schiff reagent (Solarbio, Beijing, China) for 15 min, and then rinsed in distilled water.

Total RNA Extraction and Quantitative Real-Time PCR

Total RNA was extracted from LS174T cells and cecal tissue samples at indicated time point using TRI-Reagent (Sigma-Aldrich, St. Louis, MO, USA) according to the manufacturer's guidelines. Bacterial RNA was isolated following manufacturer's instructions (Promega, Madison, WI, USA). cDNA was synthesized using Rtase M-MLV (TaKaRa, Kyoto, Japan) with oligo (dT). Quantitative real-time PCR was performed on an ABI 7500 Fast Real-Time PCR System (Applied Biosystems) using SYBR Green PCR Master Mix (Roche, South San Francisco, CA, USA). GAPDH was used as the reference gene. The relative changes in gene expression were analyzed by the $2^{-\Delta\Delta ct}$ method. The relative primers are listed in **Table 1**.

Cytotoxicity Assays

Cell viability was evaluated by the measurement of lactate dehydrogenase (LDH) leakage from damage or destroyed cells with a CytoTox 96 Non-Radioactive Cytotoxicity Assay Kit under the manufacturer's instruction (Promega, Madison, WI, USA).

Bacterial Counts

Cecum of mice were flushed free of feces with PBS. Tissues including liver, spleen, MLN and cecum of mice were weighted and then homogenized in cold PBS at a ratio of 1:6 (g/mL). The homogenates were serial diluted and 50 ul was plated on LB plates containing streptomycin. After 16 h, colonies were counted.

Cytokines Measurement

Cecum of mice were harvested and flushed free of feces with PBS, and then weighted and homogenized in cold PBS at a ratio of 1:6 (g/mL). Homogenates were centrifuged at 13000 rpm/min for 30 min and supernatants were collected and used for cytokines measurement. Cytokines were detected with ELISA assay following R&D systems instruction.

TABLE 1	Primers used for gRT-PCR.

Gene	Sequence
Mouse Mucin2	F:5'-GCTGACGAGTGGTTGGTGAATG-3'
	R:5'-GATGAGGTGGCAGACAGGAGAC-3'
Mouse GAPDH	F:5'-AGGTCGGTGTGAACGGATTTG-3'
	R:5'-GGGTCGTTGATGGCAACA-3'
Human Mucin2	F:5'-GAGGGTGGAAGTGGCAT-3'
	R:5'-TGTCGGCAGGGTTGA-3'
Human GAPDH	F:5'-ACATCATCCCTGCCTCTACTG-3'
	R:5'-ACCACCTGGTGCTCAGTGTA-3'
SopB	F:5'-GGAATTGTAAAAGCGGCAAA-3'
	R:5'-TTTTCTGTCCACCGCTATCC-3'
SopE ₂	F:5'-GGAGAGGTTATGCCGCCTTT-3'
	R:5'-CGGAGTGATCCTCAAGGCAA-3'
Eubacteria (Universal)	F:5'-ACTCCTACGGGAGGCAGCAGT-3'
	R:5'-ATTACCGCGGCTGCTGGC-3'

Statistical Analysis

All experiments were independently performed three times in triplicate. All values were expressed as means \pm standard deviations. Differences between mean values were assessed by ANOVA test. The analysis of survival was determined via logrank test. Data analysis was performed with GraphPad Prism software version 6.0 (GraphPad, La Jolla, CA, USA). *P* value was < 0.05 was considered to be significant different. **P* < 0.05, ***P* < 0.01, ****P* < 0.001, #*P* < 0.05, ##*P* < 0.01.

RESULTS

Mice Display Increased Susceptibility to $\Delta sopB$ Induced Colitis

Although SopB mediates sustained activation of the pro-survival kinase Akt in infected epithelial cells (Knodler et al., 2005), and moreover, Akt2 deficiency mice display increased susceptibility to Salmonella infection (Kum et al., 2011), the role of SopB in vivo is not clear to now. We hypothesized that SopB deleted SL1344 strain $\triangle sopB$ would affect the host's defense against Salmonella infection, leading to increased severity to disease. Therefore, we initially examined the susceptibility to colitis following the oral administration of 5 \times 10⁷ colony-forming units (CFU) of Salmonella strain SL1344 or SopB mutant strain $\Delta sopB$. Compared with SL1344 infection, $\Delta sopB$ infected mice displayed significant higher mortality (Figure 1A) and increased body weight loss (Figure 1B). Mice infected with $\triangle sopB$ displayed a significant increased splenomegaly (Figure 1C) and decreased cecum weight (Figure 1D) when compared with SL1344 infection. Consistently, H&E staining showed $\Delta sopB$ infected mice had increased intestinal damage and histopathology score characterized by greater submucosa edema, elevated polymorphonuclear (PMN) leukocyte infiltration, severe mucosa ulceration and increased goblet cells depletion (Figures 1E,F). Collectively, these data indicated that SopB plays a redundant role in the induction of intestinal inflammation during Salmonella infection, which deletion increased severity to colitis.

∆sopB Infection Leads to Reduced Mucin Expression

The major difference of histopathology in cecum between SL1344 and $\Delta sopB$ infection was especially in goblet cell depletion (**Figure 1F**). Goblet cells exert numerous biological roles to protect the gut, such as mucin secretion and antigens delivering (Pelaseyed et al., 2014; Birchenough et al., 2016). Mucus serves as the first physical barrier encountered by *Salmonella*. Therefore, mucins expression was further investigated. When compared with SL1344 infection, there was a significant reduction in mucin expression (**Figure 2A**) and numbers of mucin containing goblet cells (**Figure 2B**) in cecum following $\Delta sopB$ infection. Mucin2 is a major mucin secreted by goblet cells and has an important function in protecting against enteric pathogens (Bergstrom et al., 2010). The expression of mucin2 in cecum was subsequently examined. In concert with mucins expression, we found there was a significant



P-value for survival was determined via log-rank test. *P < 0.05, **P < 0.01, **P < 0.001.

reduction of mucin2 mRNA and protein in cecum following $\Delta sopB$ infection for 2 days (**Figures 2C,D**). Consist with the *in vivo* study, we also found there was a reduction in mucin2 expression in LS174T cells infected with $\Delta sopB$ for 14 and 24h respectively, when compared with SL1344 infection (**Figure 2E**). These results indicated that $\Delta sopB$ infection results in significant reduction in goblet cells number and mucin expression.

∆sopB Infection Promotes Goblet Cells Necroptosis

The reduced mucin expression attributes to the disturbed goblet cell function. Thus, we subsequently examined whether there was an increased goblet cells death following $\Delta sopB$ infection. We found that $\Delta sopB$ infection induced increased LS174T cell death when compared with SL1344 infection (**Figure 3A**), however, whether the goblet cells went on death in a manner of necroptosis has not been investigated. To elucidate this question, LS174T cells were pretreated with

a necroptosis inhibitor necrostatin-1 (nec-1) or NSA before bacterial infection. Surprisingly, treatment with NSA, an inhibitor of MLKL, significant reduced LS174T cell death following $\Delta sopB$ infection, however, had no obvious effect on SL1344 induced cell death (Figure 3B). These results indicated that $\triangle sopB$ infection can induce goblet cells death in a manner of necroptosis. The molecular nec-1 is a RIPK1 inhibitor which is upstream of MLKL. We found nec-1 treatment did not significantly reduce LS174T cell death following $\triangle sopB$ infection (Figure 3B). According to these results, we concluded that $\Delta sopB$ infection increased LS174T cell necroptosis via upregulating MLKL phosphorylation. To further confirm the observed phenomenon in vitro, we examined the level of MLKL phosphorylation in cecum following bacterial infection for 2 days. Consistent with the in vitro results, we found there was a significant increased expression of MLKL phosphorylation in goblet cells as well as in the other epithelial cells (Figure 3C). Collectively, these results indicated that $\triangle sopB$ infection promotes cell necroptosis via upregulating MLKL phosphorylation.



treated with SL1344 or $\triangle sopB$ for 14 and 24 h respectively. *P < 0.05, **P < 0.01.

MLKL Deletion Abolishes the Severe Inflammation in Cecum Following $\Delta sopB$ Infection

To further identify the contribution of necroptosis in $\Delta sopB$ mediated increased severity to colitis, we employed MLKL deficient mice. Intriguingly, MLKL deletion rescued severity to $\Delta sopB$ induced colitis. Following $\Delta sopB$ infection for 2 days, there was a significant decrease in the weight of cecum, however, MLKL deletion improved the loss of cecum weight (**Figure 4A**). Consistently, MLKL deletion also ameliorated the inflammatory cytokines production in cecum following $\Delta sopB$ infection (**Figure 4B**). There was a significant increased inflammatory cytokine production including TNF- α , IL-6, IFN- γ , KC, CCL2 and CXCL10, however, no significant difference was found in IL-12 production in mice infected with $\Delta sopB$ when compared with mice infected with SL1344 (**Figure 4B**). Moreover, the cytokines production including IFN- γ , CCL2, and CXCL10 were reduced in the cecum of MLKL^{-/-} mice infected with $\Delta sopB$ when compared with WT mice (**Figure 4B**). These results indicated that MLKL deletion improved the severity to $\Delta sopB$ induced colitis. Additionally, we found mucin2 expression in the cecum of MLKL^{-/-} mice infected with $\Delta sopB$ was increased when compared to WT mice (**Figure 4C**). Collectively, these data indicated that MLKL mediated necroptosis contributes to the increased severity to colitis in mice infected with $\Delta sopB$.

MLKL Deletion Reduces Bacterial Translocation Following *△sopB* Infection

Salmonella disturbs intestinal barrier and then transmits to systemic sites to establish infection. We subsequently evaluated the effect of $\Delta sopB$ mediated necroptosis on Salmonella translocation. Compared to Salmonella infection, $\Delta sopB$ infected mice had an increased bacterial colonization in cecum (**Figure 5A**) and significant increased bacterial burdens in mesenteric lymph node (MLN) (**Figure 5B**), liver (**Figure 5C**), and spleen (**Figure 5D**) in WT mice indicating that SopB deletion enhanced Salmonella pathogenesis characterized by



increased bacterial translocation. However, MLKL deletion abolished the increased bacterial translocation. Following $\Delta sopB$ infection, MLKL^{-/-} mice had decreased bacterial burdens in liver (**Figure 5C**) and spleen (**Figure 5D**) when compared to WT mice. These results indicated that $\Delta sopB$ infection induced increased necroptosis contributing to the increased bacterial translocation.

The Level of SopB Expression Is Downregulated During Salmonella Infection

Salmonella is an intracellular bacterial pathogen and has evolved many mechanisms to evade host immune response to facilitate bacteria translocation, such as regulating effector proteins expression. We proved that SopB deletion results in increased cell necroptosis and bacterial translocation. Thus, we hypothesized that *Salmonella* might adoptively downregulate SopB expression to enhance its translocation. Indeed, results showed that the expression level of *sopB* was downregulated in LS174T cells during *Salmonella* infection (**Figures 6A,B**). The downregulated *sopB* expression would enhance bacteria pathogenesis and increase cell necroptosis which facilitates bacterial translocation. Additionally, we also examined the expression level of $SopE_2$ which has a similar function as SopB. In contrast to *sopB*, the expression level of *sopE*₂ was upregulated in LS174T cells infected with *Salmonella* for 2 h (**Figures 6A,C**). These results indicated effector proteins were differently regulated during *Salmonella* infection and the downregulated *sopB* expression might represent a strategy used by *Salmonella* to spread to systemic sites and establish infection.

DISCUSSION

SopB is an important SPI-1 secreted virulence effector and exerts many roles during *Salmonella* infection. However, the mechanisms of SopB in *Salmonella* induced colitis have not been well-studied. To address this, we investigated the severity to colitis following SL1344 or $\Delta sopB$ infection. $\Delta sopB$ infection





increased goblet cell necroptosis associated with enhanced intestinal inflammation and increased bacterial translocation. The increased goblet cell necroptosis was caused by upregulating MLKL phosphorylation. MLKL deletion rescued the intestinal inflammation and bacterial loads in mice infected with $\Delta sopB$. Interestingly, SopB expression was downregulated during *Salmonella* infection. The down regulated SopB would increase cell necroptosis similar to SopB deletion which enable bacteria escape from the epithelial cell to infect neighbor cells and

consequently translocated to systemic sites. Collectively, this study indicated a functional role for SopB in preventing cell necroptosis and the downregulation of SopB expression represents a mechanism used by *Salmonella* to manipulate onset of epithelial cell death to establish infection.

Salmonella infection induced cell death is considered to be a major course of pathogenesis of gut inflammation and is tightly controlled. Many effectors translocated into epithelial cells involve in modulating cell death. It has been suggested that the







effector protein SopB protects cell from apoptosis and pyroptosis (Knodler et al., 2005; Hu et al., 2017). In our study, SopB was additionally shown to prevent goblet cell necroptosis. We found $\Delta sopB$ infection increased goblet cell necroptosis in cecum via upregulating MLKL phosphorylation. Consistently, $\Delta sopB$ infection increased LS174T cell death, whereas NSA treatment rescued the increased LS174T cell death induced by $\Delta sopB$. These results indicated that SopB play a role in preventing cell necroptosis. However, NSA treatment did not absolutely abolish

the increased cell death mediated by $\Delta sopB$. This might cause by that SopB can regulate cell death in a manner different from necroptosis. Consistent with a recent study that $\Delta sopB$ infection induced increased numbers of caspase-3 and caspase-8 positive cells, while a similar phenotype was not observed after $\Delta sopBE_2$ infection indicated that SopB counteracts the proapoptotic effect mediated by SopE₂ in an undefined mechanism (Zhang et al., 2018). Intriguingly, we found the expression of SopE₂ in LS174T cell were increased after $\Delta sopB$ infection for 2 h which might promote cell apoptosis and contribute to the increased numbers of caspase3 and caspase-8 positive cells induced by $\Delta sopB$ infection.

Necroptosis is generally considered to be pro-inflammatory and has a role in pathogenesis of bacterial infection (Kitur et al., 2015; Pasparakis and Vandenabeele, 2015; Parker and Prince, 2016). The hall mark of necroptosis is that MLKL forms pores or actives ion channels in the membrane of cells resulting in membrane rupture and the leakage of intracellular contents such as inflammatory mediators (Jorgensen et al., 2017). Due to overt inflammatory response caused by the leakage of immunostimulatory contents, necroptosis is considered to be an inflammatory death (Newton and Manning, 2016). Inflammation is required for the control of bacterial infection. However, the overt and/or prolonged inflammation is one of the reasons for local tissue pathogenesis contributing to the excessive bacterial translocation and the development of pathogenesis of inflammatory disease. Additionally, epithelial cell necroptosis facilitates bacteria escape host cell and translocate to systemic sites. Numerous studies have demonstrated that necroptosis has a crucial role in enhancing infection (Robinson et al., 2012; Kitur et al., 2015). Studies have shown that Salmonella exploits type I IFN signaling to induce macrophage necroptosis (Robinson et al., 2012; Hu and Zhao, 2013). Ifnar^{-/-} mice challenged with Salmonella showed an improved survival as well as lower bacterial burdens in liver and spleen. Inhibition of necroptosis induced by Salmonella infection reduced inflammation and improve bacterial clearance (Robinson et al., 2012). Consistently, in present study we found $\triangle sopB$ infection induced an increased level of MLKL phosphorylation in cells of cecum associated with increased inflammation and bacterial translocation. Deletion the final executor of necroptosis MLKL abolished the increased bacterial translocation and severity to inflammation following $\triangle sopB$ infection. All these results indicated necroptosis represents a pro-inflammatory response. The increased epithelial cell necroptosis mediated by $\triangle sopB$ infection is a mechanism used by Salmonella to breakdown epithelial barrier and spread to systemic sites. However, we did not consider the potential effect of MLKL deficiency on intestinal microbiota composition, which may affect intestinal disease symptoms. Recently, a newly published study found MLKL deficiency has a little effect on intestinal microbiota composition which did not affect severity to DSS induced colitis (Zhang et al., 2019). Whether the altered intestinal microbiota contributes to Salmonella induced disease need to be further investigated.

To establish an infection, *Salmonella* first enters into epithelial cell and replicates within *Salmonella*-containing vacuole, and then escapes from epithelial cell in the help of effectors (Fabrega and Vila, 2013). Once the epithelial barrier is disrupted,

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Abrahams, G. L., and Hensel, M. (2006). Manipulating cellular transport and immune responses: dynamic interactions between intracellular *Salmonella enterica* and its host cells. *Cell. Microbiol.* 8, 728–737. doi: 10.1111/j.1462-5822.2006.00706.x Salmonella is internalized by macrophage and subsequently transmits to systemic sites. All these phases are connected with cell death. In the initial stage of infection, cell death must be prevented which provides a favorite environment for Salmonella replication. During the last phage of infection, cell death is required to promote Salmonella escape epithelial cells. To realize this role switching, effector proteins are temporally regulated. In present study, we found SopB expression was downregulated. The downregulated SopB would induce cell necroptosis which potentially switches the role of SopB from epithelial cell invasion to escape host cells. Therefore, the temporally regulated SopB secretion represents a virulence strategy used by Salmonella to establish infection. Epithelial cells have important roles in protecting enteric pathogens infection (Broz et al., 2012). SopB cooperates with other effectors to mediate bacterial invasion and cell death. Though SopB is required for invasion, deletion of SopB fail to alter the ability of invasion and has no pathogenesis defect. Additionally, we found SopB not only mediated the upregulation of MLKL phosphorylation in epithelial cell, but also in immune cells. However, we focus the function of SopB in goblet cells due to the mainly difference in pathogenesis as well as the crucial role of goblet cell in defending against enteric pathogens infection.

In addition to the role in delaying apoptosis and pyroptosis, this study indicated a different role for SopB in modulating cell necroptosis. We found SopB deletion upregulated MLKL phosphorylation which increased cell necroptosis and consequently increased severity to colitis and promotes bacterial translocation. This study further identified a different biological mechanism that the downregulated SopB expression increased host cell necroptosis represents a devised strategy used by *Salmonella* to establish infection.

AUTHOR CONTRIBUTIONS

G-QH and Y-JY performed most of the experiments, collected and analyzed the results, and wrote the manuscript. WC conceived and designed the study. X-XQ, SQ, and S-XY performed the animal study. JZ performed the IHC experiments. C-TD performed the H&E study. All authors have read the manuscript and approved the final manuscript.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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