



p40^{phox}-Deficient Mice Exhibit Impaired Bacterial Clearance and Enhanced Pro-inflammatory Responses during Salmonella enterica serovar Typhimurium Infection

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Li Y, Lv M, Su C, Long S, Zhang W, Conway KL, Li W, Xavier RJ and Shi HN (2017) p40^{phox}-Deficient Mice Exhibit Impaired Bacterial Clearance and Enhanced Pro-inflammatory Responses during Salmonella enterica serovar Typhimurium Infection. Front. Immunol. 8:1270. doi: 10.3389/fimmu.2017.01270 Salmonella enterica serovar Typhimurium (S. Typhimurium) is a major cause of acute gastroenteritis in humans. During infection, reactive oxygen species (ROS), generated from NADPH oxidase (a multisubunit enzyme complex), are required for pathogen killing upon phagocytosis and for regulating pro-inflammatory signaling in phagocytic cells. Mutations in subunits forming the NADPH complex may lead to enhanced susceptibility to infection and inflammatory disease. Compared to other NADPH oxidase subunits, the function of p40^{ohox} is relatively understudied, particularly in the context of intestinal bacterial infection. In this study, we utilized genetically engineered mice to determine the role of $p40^{phox}$ in the response to S. Typhimurium infection. We show that mice lacking p40^{phox} are more susceptible to oral infection with S. Typhimurium, as demonstrated by significantly enhanced bacterial dissemination to spleen and liver, and development of exacerbated bacterial colitis. Moreover, we demonstrate that the increased infection and disease severity are correlated with markedly increased F4/80⁺ macrophage and Ly6G⁺ neutrophil infiltration in the infected tissues, coincident with significantly elevated pro-inflammatory cytokines (IL-1 β and TNF- α) and chemoattractant molecules in the infected tissues. Functional analysis of macrophages and neutrophils further shows that p40^{ohox} deficiency impairs bacteriaor PMA-induced intracellular ROS production as well as intracellular killing of Salmonella. These observations indicate that the p40^{ohox} subunit of NADPH oxidase plays an essential role in suppressing intracellular multiplication of Salmonella in macrophages and in the regulation of both systemic and mucosal inflammatory responses to bacterial infection.

Keywords: NADPH oxidase, bacterial killing, mucosa, inflammation, Salmonella infection

INTRODUCTION

Salmonella enterica serovar Typhimurium (*S.* Typhimurium) is a Gram-negative food-borne pathogen that is frequently associated with disease in various host species, including humans, livestock, domestic fowl, and rodents (1). In human, *S.* Tyhimurium is a major cause of acute gastroenteritis. There are approximately 40,000 cases of acute gastroenteritis annually in the United States (1).

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Although mice usually do not develop signs of the intestinal inflammation observed in humans following oral infection by S. Typhimurium, the bacteria penetrate the intestinal epithelium and move into the mesenteric lymph nodes (MLN), then spread to the circulatory system, causing systemic disease (2, 3). The Salmonella-induced intestinal inflammation model has also been established, which involves pretreatment of mice with streptomycin and then oral inoculation with a strain of S. Typhimurium that is naturally resistant to the antibiotic (4, 5). This bacterial colitis model provides an experimental system that is well suited to the analysis of host mucosal immune and inflammatory responses. In this model, replication of S. Typhimurium in lamina propria monocytic phagocytes/macrophages plays an important role in the development of colitis (6). However, it is less clear whether/how host mucosal defenses are affected by NADPH oxidase. In the infected intestinal mucosa, a mixture of different phagocytes has been observed, including polymorphonuclear neutrophils (PMN), dendritic cells (7), and macrophages, which play an essential role in host defense against invading bacterial pathogens. To effectively control the infection, molecules that can enhance the microbicidal capacity of phagocytes and the recruitment of various cell populations, such as neutrophils and monocytes, are required (2, 8, 9).

Phagocyte NADPH oxidase, a membrane-bound enzyme complex that is composed of six subunits, the catalytic subunit gp91^{phox} and p22^{phox}, and the cytosolic regulatory components (p47^{phox}, p67^{phox}, p40^{phox}, and Rac), plays an important role in microbial killing through the generation of reactive oxygen species (ROS) (10-12). ROS generation is one of the central mechanisms used by phagocytes to defend against bacterial infection (13-15). The importance of ROS production in host defense is demonstrated by the enhanced susceptibility to infection of individuals who have inherited deficiencies of NADPH oxidase and develop chronic granulomatous disease (CGD) (16). gp91^{phox} is the catalytic core of the NADPH oxidase and can produce ROS in the absence of other cytosolic subunits. Mutations in gp91^{phox} account for the majority of CGD patients in the world (65% of all cases are the X-linked gp91^{phox}-deficient form) (17). Evidence also shows that there is an excessive inflammation in CGD even in the absence of infectious agents (7). Furthermore, mice deficient in NADPH oxidase (Cybb^{-/-}, encoding gp91^{phox}) failed to control infection with a normally avirulent bacterial infection (6). Mutations in the NCF4 gene encoding $p40^{phox}$, a less well understood subunit of the NADPH oxidase complex, were first described in CGD with abnormal neutrophil function in 2009, which disproved the view that p40^{phox} was not essential for NADPH oxidase activity (18). Evidence is also available to suggest a role for p40^{phox}, in host resistance against Staphylococcus aureus infection and in the regulation of neutrophil recruitment and function (19, 20). In genetic studies the NCF4 gene ($p40^{phox}$) locus has been found to be associated with ileal and perianal Crohn's disease (21-23), supporting a role for NADPH oxidase dysfunction in the intestinal inflammation of IBD. It has been reported that p40^{phox-/-} mice showed increased susceptibility to both DSS- and anti-CD40-induced colitis (19). However, the role of p40^{phox} in the regulation of intestinal immune response during infectious colitis and the importance of p40^{phox} expression

by phagocytes in controlling and regulating intestinal mucosal bacterial infection is less well understood.

Macrophages and neutrophils contribute significantly to the effector phase of the immune response, i.e., elimination of bacteria, and are also thought to be critical mediators of many chronic inflammatory diseases. These phagocytic cells have evolved a repertoire of antimicrobial mechanisms based on the formation of toxic radicals, including NADPH phagocytic oxidase and inducible nitric oxide synthase. In line with many other investigations, our previous studies showed that macrophages can kill infected bacterial pathogens, such as Salmonella and Citrobacter rodentium via autophagy (24), a catabolic process that several lines of evidence have suggested can be regulated by ROS and reactive nitrogen species (RNS) via undefined molecular mechanisms (25). The generation of ROS has also been shown to induce the activation of inflammasome (26), a cytosolic protein complex that senses microbial stimuli and regulates the maturation of inflammatory cytokine IL-1β production (27). However, mononuclear phagocytes from CGD patients (deficient in ROS production) have been shown to have an increased IL-1 β secretion (28). In this study, we utilized genetically engineered mice to determine the role of p40^{phox} in host innate defense against both systemic and mucosal infection of S. Typhimurium, as well as the mechanism of action.

MATERIALS AND METHODS

Mice

Six- to eight-week-old female C57BL/6 [wild-type (WT)] mice were obtained from the Jackson Laboratory (Bar Harbor, ME, USA). p40^{phox} knockout mice, which have been backcrossed to the B6 background for 10 generations, were kindly provided by Dr. Ramnik Xavier (Massachusetts General Hospital, Boston, MA, USA) (19). All mice were fed autoclaved food and water and maintained in a specific-pathogen-free facility at Massachusetts General Hospital.

S. Typhimurium Infection

Mice were infected orally with 3×10^8 CFU of the streptomycin-resistant SL1344 strain of S. Typhimurium. After 72 h of infection, mice were euthanized by CO2 inhalation. Spleens and livers were harvested to determine bacterial translocation and systemic inflammatory response. To study whether p40^{phox} deficiency affects the early mucosal innate immune and inflammatory response to Salmonella infection, the streptomycinpretreatment mouse model was used as described previously (4). Briefly, mice were each given 20 mg of streptomycin, followed 24 h later by oral infection with 10⁸ CFU of S. Typhimurium. Ceca were collected for histopathology 24 h after infection. To determine the impact of p40^{phox} deficiency on the host during Salmonella infection, WT mice and mice lacking p40^{phox} were infected orally with 3×10^8 CFU of S. Typhimurium. Body weight loss and survival of mice were monitored daily. Mice were humanely euthanized if they exhibited greater than 15% weight loss.

Determination of Salmonella Translocation

Spleens and livers collected from *Salmonella*-infected mice (WT and $p40^{phox-/-}$ mice) (with or without streptomycin treatment), were excised using sterile procedure, weighed, homogenized, and plated on Luria–Bertani plates containing 50 µg/ml streptomycin (3). The CFU were quantified. To examine the appearance and distribution of *Salmonella* in cecum and spleen, cryostat sections of the intestinal tissues and spleens were incubated with a rabbit antibody against *Salmonella* (Thermo Fisher Scientific), followed by fluorescein isothiocyanate (FITC) or Cy3-conjugated goat anti-rabbit IgG antibody (Biosource Cat. #554020), and analyzed by immunofluorescence microscopy.

Histopathology

At necropsy, tissue samples of spleen, liver, and cecum were collected, frozen in Tissue-Tek OCT compound (Miles Inc., Elkhart, IN, USA) and then stored at -80°C. Then, 5-µm sections were cut and stained with hematoxylin and eosin (H&E). Pathology was scored using a modified histological scoring systems previously published in the literature (5, 29, 30). The scores were assessed by determination of infiltration of inflammatory cells, with scores ranging from 0 to 4 (0, normal cell pattern; 1, scattered inflammatory cells in the lamina propria; 2, increased numbers of inflammatory cells in the lamina propria; 3, confluence of inflammatory cells extending into the submucosa; and 4, transmural extension of the infiltrative inflammatory cells), together with the evaluation of cecal tissue damage (score range, 0-4; 0, normal tissue pattern; 1, minimal inflammation and crypt hyperplasia; 2, mild crypt hyperplasia with or without focal invasion of epithelium; 3, obvious crypt hyperplasia, invasion of epithelium, and goblet cell depletion; and 4, extensive mucosal damage and extension through deeper structures of the bowel wall). Stained sections were analyzed without prior knowledge of the type of treatment.

Immunofluorescence Microscopy

Tissue cryosections were fixed in ice-cold acetone, washed, and then blocked with avidin/biotin agent (Vector Laboratories, Burlingame, CA, USA). To analyze the location and abundance of macrophages and neutrophils, cecal slides were stained with FITC-labeled anti-mouse F4/80 (eBioscience Cat. #11-4801-81) and Cy5-labeled anti-mouse Ly6G (Biolegend Cat. #108410). DNA was stained and mounted using the 4', 6-diamidino-2-phenylindole (DAPI) (Vector Laboratories). To examine the bacterial killing capacity, peritoneal macrophages were collected from uninfected WT and p40^{phox} KO mice, grown on cover slips, and infected with S. Typhimurium for 1 h (cell/bacteria, 1:10). Cells were then stained with phycoerythrin (PE)-conjugated anti-F4/80 at both 2 and 6 h after infection (24). Internalization and elimination of bacteria by macrophages were visualized by detecting Salmonella with a rabbit antibody against Salmonella, followed by FITC-labeled anti-rabbit IgG antibody.

Quantitative Real-time PCR Analysis

Total RNA was isolated from spleen, liver, cecum, and peritoneal macrophages using TRIzol reagent (Invitrogen Life Technologies,

Carlsbad, CA, USA) or RNeasy Kit (Qiagen, Valencia, CA, USA) following the manufacturer's instruction. All RNA samples were reverse transcribed into cDNA using the Superscript First-Strand Synthesis System (Invitrogen Life Technologies). The cDNA samples were then tested for the expression of TNF- α , IL-1 β , KC, MCP1, and MIP2 by real-time RT-PCR performed as previously described (29). Results were normalized to GAPDH expression and relative quantification was calculated using the $2^{-\Delta\Delta CT}$ method.

Measurement of Cytokine Production by ELISA

Liver and spleen homogenate was prepared for cytokines measurement through ELISA. Protein concentrations were determined using a bicinchoninic acid protein assay. IL-1 β was measured by DuoSet ELISA kit from R&D Systems (Cat. #DY401) according to the manufacturer's instruction. ELISA capture antibodies and biotinylated secondary antibodies for IFN- γ (BD Cat. #554410; 551216), TNF- α (BD Cat. #554416; 557432), and IL-17A (BD Cat. #555068; 555067) were purchased from DB Bioscience. Standard curves were obtained using recombinant murine IFN- γ (BD Cat. #554587), TNF- α (BD Cat. #554589), and IL-17A (eBioscience Cat. #14-8171-80) from BD Bioscience and eBioscience.

Flow Cytometry Analysis

Splenic single-cell suspensions were prepared from uninfected mice and mice infected orally with 3×10^8 CFU S. Typhimurium at 24 h after bacterial infection. Cells were surface stained with fluorescent-conjugated antibodies against CD11b (eBioscience, Cat. #11-0112-85; BD Cat. #553311), Ly6G (BioLegend, Cat. #108410), and F4/80 (eBioscience, Cat. #12-4801-80) before being subjected to flow cytometry analysis with an Accuri C6 FACS machine. Gates set on forward and side angle light scatter were used to exclude dead cells and debris.

Lucigenin-Dependent Chemiluminescence Assay and 2',7'-Dichlorofluorescin Diacetate (DCFDA) Intracellular ROS Detection Assay

Reactive oxygen species generation was measured using lucigenin-dependent chemiluminescence. Peritoneal macrophages were collected by washing the peritoneal cavity of WT or p40^{phox-/-}mice with ice-cold PBS. Neutrophils were purified using EasySep[™] Neutrophil Enrichment Kit (Stemcell Technologies, Catalog #19762) according to the manufacturer's instructions. Briefly, mouse bone marrow cells were flushed from femurs and tibias, centrifuged and, after hypotonic lysis of erythrocytes, mixed with biotinylated antibodies and magnetic particles. Neutrophils were then separated by immunomagnetic negative selection. Macrophages and neutrophils were then resuspended in 100 μ M lucigenin (5 \times 10⁷/ml) and incubated for 15 min on ice. Subsequently, cells were seeded in a black 96-well plate (4 \times 10⁶/well) and stimulated with 200 nM PMA (Sigma-Aldrich). Macrophages without PMA treatment were used as a control. Light emission was measured on a TopCount NXT scintillation and luminescence reader (PerkinElmer, Waltham, MA,

USA) every 5 min for a period of 1 h at 37°C. Data were expressed as relative light units (RLU) normalized to cell number and corrected for a sample blank. Each experiment was performed in triplicate.

In addition, intracellular ROS generation was measured using DCFDA Cellular Reactive Oxygen Species Detection Assay Kit (Abcam, Ab113851) in accordance with the manufacturer's instructions and published methods (31). DCFDA is a fluorogenic dye that measures ROS activity within the cell. For these experiments, both intestinal and peritoneal macrophages were isolated from C57BL/6 and p40^{phox-/-} mice. The macrophages from intestine were isolated using published methods (32). Briefly, small and large intestine were collected and Pever's patches were removed. The intestinal tissues were cut open longitudinally (to 1.5 cm pieces) and washed in PBS. The epithelial layer was removed by shaking the tissues in EDTAcontaining HBSS/FBS buffer (250 rpm for 20 min at 37°C). The intestinal tissue was then digested with CollagenaseVIII/DNAse I solution and filtered through a 100 µm cell strainer. After washing, the cell pellets were collected. Intestinal macrophages were enriched using magnetic beads (anti-F4/80 and CD11b). The cells were then stained in the diluted DCFDA solution at 37°C for 30 min in the dark, then washed. Cells were seeded in a dark side, clear bottom 96-well microplate (2×10^5 stained cells/ well). The cells were then stimulated with 20 nM PMA (Sigma-Aldrich) or infected with S. Typhimurium (cell/bacteria, 1:100). Macrophages without PMA treatment or Salmonella infection were used as a control. The microplates were read using a fluorescent plate reader (FluoSTAR OPTIMA-BMG Labtech) with the excitation wavelength set at 485 nm and emission wavelength at 535 nm. Results are expressed as fluorescent intensity of samples minus blank (DCFDA-unstained cells). Each experiment was performed in triplicate.

Gentamicin Protection Assay

The resident peritoneal macrophages were collected from uninfected WT and $p40^{phox}$ knockout mice. After 2 h of incubation in complete DMEM, non-adherent cells were removed by washing. The adherent cells were cultured at 37°C overnight and then exposed to 10⁷ S. Typhimurium (cell/bacteria, 1:10) for 1 h in antibiotic-free medium. After completion of the infection period, cells were washed with PBS and then incubated with gentamicin-containing medium (100 µg/ml) for 2 h to eliminate extracellular bacteria. Cells were then lysed immediately in sterile 1% Triton X-100 in water or, after a further 4 h, in medium containing 10 µg/ml gentamicin. Serial dilutions of the cell lysates were plated on LB agar containing streptomycin overnight. Colonies were counted after incubation and bacterial numbers present inside the cells at each time point were calculated (24, 33).

Statistical Analysis

Results are expressed as the mean \pm SEM. Survival curves of infected mice (WT and p40^{phox-/-}) were compared using Kaplan–Meier analysis followed by log-rank test. Statistical differences were determined using one-way ANOVA with Tukey's *post hoc* test, the two-tailed Student's *t*-test or paired *t*-test (body weight

loss) with GraphPad Prism (GraphPad Software, San Diego, CA, USA). Significance was defined as a *P*-value <0.05.

RESULTS

p40^{ohox} Deficiency Increases Morbidity of Mice during *Salmonella* Infection

NADPH oxidase is critical in host defense against invading pathogens (34). To investigate the role of p40^{phox}, one of the subunits of the NADPH oxidase, in host protective and inflammatory responses against Salmonella infection and in the regulation of the severity of bacterial-induced disease, p40phox-/- mice and C57BL/6 WT control mice were infected orally with 3×10^8 CFU of S. Typhimurium (SL1344 strain). As expected, mice infected with S. Typhimurium exhibited significant body weight loss and high mortality (100%) by day 5 of infection in WT mice. By contrast, p40^{phox-/-} mice developed more severe disease, resulting in a more significant body weight loss and early mortality reaching 100% mortality by 4 days after inoculation (Figures 1A,B). To examine whether the increased susceptibility of p40^{phox}-deficient mice to Salmonella infection was related to poor control of bacterial replication, bacterial translocation to spleens and livers was determined 72 h after infection. Significantly, greater bacterial numbers were present in the spleens (Figure 1D) and livers (Figure 1C) of p40^{phox-/-} mice compared with C57BL/6 controls, suggesting an impaired mucosal barrier function and inability to effectively clear microbes in p40^{phox}-deficient mice. This is further supported by the results from our immunofluorescence microscopic analysis revealing enhanced bacterial dissemination in the spleen tissues of p40^{phox}-deficient mice (Figure 1E).

The increased tissue bacterial loads were accompanied by exacerbated inflammation and tissue damage, which was evidenced by the observations from analysis of histopathology and pro-inflammatory cytokine profiles of these organs from WT and p40^{phox-/-} mice with and without S. Typhimurium infection. Histological analysis of infected mice revealed extensive pathological changes in the spleen, evidenced by a neutrophilic infiltrate and congestion within the red pulp in Salmonellainfected WT mice (Figure 2A). In contrast to WT mice, p40^{phox} knockout mice developed more severe tissue damage, characterized by red pulp congestion, thrombosis, and splenic architecture disruption (Figure 2A). Histological analysis of the liver sections from Salmonella-infected mice showed massive necrosis accompanied by a mixed inflammatory cell infiltration in both WT and p40^{phox}-deficient mice. However, more pronounced and extensive hepatic lesions were noted in Salmonella-infected p40^{phox-/-} mice compared to WT mice (Figure 2B). These results suggest that Salmonella infection of p40^{phox} KO mice results in an exacerbated inflammation and tissue injury in the spleen and livers due to the higher numbers of bacteria in the tissues.

Consistent with more severe inflammatory changes in the spleen and liver, we found there was a significant upregulation of gene expression of inflammatory cytokine IL-1 β (**Figure 2C**) and TNF- α (**Figures 2D,E**) in infected p40^{*phox*} KO mice. Furthermore, ELISA analysis of cytokine concentration in spleen and liver homogenates confirmed that *Salmonella* infection induced a



were measured 72 h post infection. Significance was determined using a paired *t*-test for body weight (n = 5/group; *p < 0.05; **p < 0.01; ***p < 0.01). Bacterial translocation to spleens (**C**) and livers (**D**) was determined 72 h after infection using a Student's *t*-test. The number of CFUs per gram of tissue is shown (n = 9-12/group; ***p < 0.001). (**E**) Immunofluorescence microscopy data show the distribution of *Salmonella* (in Green) in spleen sections (magnification 40x). Neutrophils were identified by positive staining with anti-Ly6G. Data shown are generated from three independent experiments.

marked increase in production of TNF- α in spleen from mice lacking p40^{*phox*} (**Figure 2H**). Moreover, a trend of increase was found for IL-1 β production in spleen from p40^{*phox-/-*} mice (**Figure 2I**) while there was no statistical significance. However, the levels of IFN- γ (**Figure 2F**) and IL-17A (**Figure 2G**) present in the livers and spleens of infected $p40^{\textit{phox}-\!/-}$ mice were comparable to WT levels. These findings suggest that $p40^{\textit{phox}}$ deficiency results in increased pro-inflammatory cytokine production (including TNF- α and IL-1 β) during bacterial infection, which may be responsible for the observed enhanced tissue damage.





Collectively, the data demonstrate that p40^{phox} is critical for host resistance and survival during *Salmonella* infection.

p40^{phox} Deficiency in S. Typhimurium-Infected Mice Enhances the Recruitment of Macrophages

To further study the influence of p40^{phox} deficiency on host innate immunity against Salmonella infection, we next analyzed the frequency of macrophages and neutrophils in the splenic tissue of p40^{phox} knockout mice with Salmonella infection. Our FACS analysis of spleen cells revealed that the frequency of macrophages expressing CD11b⁺ F4/80⁺ was significantly enhanced in infected p40^{phox-/-} mice compared to WT mice (Figures 3A,C). A significant increase in splenic Ly6G⁺CD11b⁺ neutrophil numbers was also observed between WT mice and p40^{phox}-deficient mice with Salmonella infection (Figures 3B,D). These results are further supported by the results from our immunofluorescence microscopic analysis of spleen revealing enhanced macrophages and neutrophil in the spleen tissues of p40^{phox}-deficient mice (Figure 3E). These data suggest that loss of p40^{phox} results in increased transmigration of macrophages and neutrophils to the site of infection. We next determined the role for p40^{phox} in the regulation of bacteria-induced chemoattractant expression, which regulates macrophage and neutrophil recruitment. Accordingly, we assessed the expression of the three major chemoattractants: KC (CXCL1), MCP1 (CCL2), and MIP2 (CXCL2) in the spleen tissues. Our RT-PCR results showed that Salmonella infection induced an upregulation of KC, MCP1, and MIP2 gene expression in the spleens. This bacteria-induced upregulation of chemokines expression was more pronounced in the spleen of p40^{phox}-deficient mice than that detected in WT mice (Figures 3F-H). These observations show that p40^{phox} deficiency in mice may enhance the recruitment of phagocytes to the infection site by regulating the expression of KC, MCP1, and MIP2 during Salmonella infection and suggest that the excessive accumulation of granulocytes in the spleens of infected p40^{phox} KO mice may thereby enhance the inflammatory response.

p40^{ohox} Deficiency Exacerbates Salmonella-Induced Mucosal Injury

Our data mentioned above suggest that Salmonella induced more severe systemic infection (i.e., spleen and liver infection) in p40phox-deficient mice. To study whether and how p40phox deficiency affects the early mucosal innate immune and inflammatory responses, we utilized the streptomycin-pretreatment model of Salmonella infection, which has been widely used in the field to analyze Salmonella-induced intestinal inflammation (2, 4, 5, 29). In brief, mice were treated with streptomycin prior to infection with S. Typhimurium (108 CFU) and sacrificed 24 h post infection. Our results showed that mice infected with Salmonella developed acute disease, as indicated by a hunched posture, disturbed body hair, soft stool, and body weight loss soon after bacterial inoculation. A more severe disease status was observed in p40^{phox-/-} mice after Salmonella infection (data not shown). Histological examination of cecum showed extensive pathological changes that were typical in this model, including pronounced submucosal edema, dramatic disruption of epithelial architecture, and marked inflammatory cell infiltrates (Figure 4A). The ceca of p40^{phox-/-} mice showed a more pronounced edema of the intestinal tissue and increased inflammatory cellular infiltration (Figure 4A) than that detected in WT mice. These cecal inflammatory features were absent from uninfected mice (Figure 4A). Using the pathological scoring system (29), we found that the cecal inflammation was notably more severe in p40^{phox} KO mice than in WT mice (Figure 4B). Immunofluorescence microscopic analysis of infected intestine (cecum) revealed increased bacterial loads in the tissue of the p40^{phox} KO mice (Figure 4D) compared to WT B6 mice (Figure 4C). Further analysis of tissue bacterial loads showed that the enhanced severity of intestinal injury in the p40^{phox} KO mice was associated with significantly higher bacterial numbers in livers (Figure 4E) and spleens (Figure 4F) compared to WT mice, again an indication of more severely impaired bacterial killing activity in p40phox-deficient mice. Moreover, and consistent with the severe intestinal injury, the mRNA expression levels of IL-1 β (Figure 4G) and TNF- α (Figure 4H) were upregulated markedly in the cecum of infected mice compared with uninfected mice at 24 h post exposure to Salmonella.

Next, we examined whether p40^{phox} deficiency affected the infiltrating cell population in cecal lamina propria of mice in the streptomycin-pretreated Salmonella-colitis model using the immunofluorescence approach. Ceca sections were stained with anti-F4/80 (35) for macrophages and anti-Ly6G for neutrophils (29). As shown in Figures 5C,D, the immunofluorescence examination revealed a clear infiltration of F4/80⁺ macrophages and Ly6G⁺ neutrophils in the cecal submucosa in the infected mice, but was absent in the ceca of the uninfected mice in both groups (Figures 5A,B). A significantly higher frequency of both F4/80⁺ macrophages and Ly6G⁺ neutrophils was noted in the cecal lamina propria of p40^{phox} knockout mice with Salmonella infection (Figures 5E,F) compared to WT mice. The observed increased macrophage and neutrophil infiltration in the intestinal tissue was found to be correlated with upregulation of KC (Figure 5G) and MCP1 (Figure 5H) in the cecum of infected mice compared with uninfected mice at 24 h post exposure to Salmonella. Notably, a much higher expression level of KC was observed in the inflamed cecum of $p40^{phox-/-}$ mice than in WT mice (Figure 5G). Taken together, these results suggest that p40^{phox-/-} mice develop exacerbated Salmonella-induced enterocolitis, characterized by severe tissue injury and enhanced macrophages and neutrophil infiltration in the intestinal tissue.

p40^{ohox-/-} Macrophages Exhibit Normal Bacterial Phagocytosis and Reduced Bactericidal Activity

Despite the observed increase in macrophages and neutrophils, our results also showed that $p40^{phox-/-}$ mice have elevated numbers of bacteria present in infected tissue, suggesting that these phagocytes are unable to effectively control bacterial multiplication in $p40^{phox-/-}$ mice. Previous study showed that the NADPH oxidase deficiency of $p40^{phox-/-}$ neutrophils were severely deficient in bacterial killing (20). In this study, we evaluated the bacterial killing capacity of $p40^{phox-/-}$ macrophages, as macrophages are







FIGURE 4 | $p40^{phore-r/-}$ mice show more severe *Salmonella*-induced mucosal injury and enterocolitis. Mice lacking $p40^{phore}$ and wild-type (WT) mice were treated with streptomycin prior to infection with *Salmonella enterica* serovar Typhimurium (10⁸ CFU) and sacrificed 24 h post infection. (**A**) Representative hematoxylin and eosin-stained cecum sections from each group of mice (original magnification 40x). (**B**) Histopathological score of cecal inflammation in mice with or without infection of *Salmonella*. The scores were assessed by determination of infiltration of inflammatory cells (score range, 0–4), together with the evaluation of cecal tissue damage (score range, 0–4). (**C**,**D**) Representative immunofluorescence-stained cecum sections from each group of mice showing increased intestinal bacterial loads in the cecum of $p40^{ohorx}$ deficient mice compared to WT mice. Numbers of bacteria recovered from liver (**E**) and spleen (**F**) homogenates of *Salmonella*-infected mice were determined using a Student's *t*-test. The number of CFUs per gram of tissue is shown (n = 5-6/group; *p < 0.05). Cecal levels of IL-1 β (**G**) and TNF- α (**H**) were determined by quantitative PCR (n = 5/group; *p < 0.05; **p < 0.01). Data are representative of two independent experiments.

the major population of tissue-resident mononuclear phagocytes, contributing significantly to the elimination of bacteria, and are the major source of critical mediators of inflammation. We isolated peritoneal macrophages from WT and $p40^{phax-/-}$ mice,

infected the cells with *Salmonella in vitro*, and then used gentamicin protection assay as a measure of macrophage microbicidal function (24). Our results showed that the macrophages isolated from $p40^{phox-/-}$ mice internalized similar numbers of bacteria at



FIGURE 5 | $p40^{phax}$ deficiency enhances the recruitment of macrophages and neutrophils in cecal lamina propria. Mice deficient in $p40^{phax}$ and normal mice were treated with streptomycin prior to Salmonella enterica serovar Typhimurium infection and sacrificed 24 h post infection. Ceca from uninfected B6 **(A)** and $p40^{phax}$ KO **(B)** mice and infected B6 **(C)** and $p40^{phax}$ KO **(D)** mice were stained with anti-F4/80 for macrophages and anti-Ly6G (Red) for neutrophils (magnification 100x). The mean number of F4/80⁺ cells **(E)** and Ly6G⁺ cells **(F)** detected in each high power field was determined by counting four fields from each sample (samples from three mice per group were counted) (*p < 0.05; **p < 0.01). Cecal levels of KC **(G)** and MCP1 **(H)** were determined by quantitative PCR (n = 5/group; *p < 0.05). Data are representative of two independent experiments.

2 h after infection compared with cells isolated from WT mice (**Figure 6A**). However, at 6 h after initial bacterial exposure, the number of viable bacteria recovered from $p40^{phox}$ -deficient macrophages was significantly higher than that in macrophages isolated from WT mice (**Figure 6A**), indicating an impaired bacterial killing capacity of $p40^{phox-/-}$ macrophages.

To further determine the functional alterations of p40^{phox-/-} macrophages, peritoneal macrophages from WT and p40^{phox-/-} mice were collected, grown on cover slips, and infected with Salmonella. After incubation in gentamicin-containing medium, we visualized the influence of p40^{phox} deficiency on the uptake and subsequent elimination of Salmonella by macrophages in vitro using immunofluorescent staining approach at 2 and 6 h after bacterial infection. Our results showed that at the early time point (2 h), F4/80⁺ macrophages isolated from WT and p40^{phox-/-} mice internalized bacteria equally well, evidenced by the detection of similar frequency and intensity of Salmonella in macrophages (Figure 6B). At the 6 h time point, the number of cells that contained bacteria was markedly reduced in WT macrophages and compared to that detected in p40^{phox-/-} macrophages (Figure 6B), confirming that macrophages from p40^{phox-/-} mice are impaired in killing internalized bacteria.

Our cytokine production analysis of *Salmonella*-infected tissues revealed a more pronounced increase in TNF- α and IL-1 β response in p40^{phox}-deficient mice (**Figures 2** and 4). To define the impact of p40^{phox}-deficiency on bacterial antigen-induced macrophage cytokine response, we collected peritoneal macrophages from WT and p40^{phox-/-} mice and exposed the cells to *Salmonella* LPS *in vitro* (36). Our real-time PCR analysis revealed that LPS treatment significantly upregulated the gene expression of TNF- α and IL-1 β in both WT and p40^{phox-/-} cells, whereas the expression levels of IL-1 β and TNF- α in p40^{phox-/-} macrophages were significantly higher than those isolated from normal mice (**Figure 6C**). These results suggest that p40^{phox} deficiency induces dysregulation of pro-inflammatory responses of macrophages.

p40^{phox-/-} Macrophages Exhibit Reduced ROS Generation

Using a lucigenin-dependent chemiluminescence assay, we measured PMA-induced ROS generation by peritoneal macrophages as well as neutrophils isolated from WT and p40^{phox-/-} mice and confirmed that the production of ROS was markedly reduced in p40^{phox-/-} cells (Figures 7A,B). However, PMA-induced ROS production was not completely abolished in p40^{phox-/-} macrophages and neutrophils. Upon phagocytosis of the pathogens, the ROS are produced. In the next set of experiments, using the DCFDA Cellular ROS Detection Assay Kit, we determined the impact of p40^{phox} deficiency on the intracellular ROS production of both intestinal (Figures 7C,D) and peritoneal (Figures 7E,F) macrophages in response to S. Typhimurium infection. Intestinal and peritoneal macrophages were isolated from both WT and p40^{phox}-deficient mice and stained in the diluted DCFDA solution before being infected with S. Typhimurium or treated with PMA. Our results show that Salmonella-induced macrophage ROS production was significantly higher in cells from WT mice compared to p40^{phox-/-} mice (Figures 7C,E). Moreover, PMA-induced ROS production of macrophages was found to be markedly lower in p40^{*phox-/-*} macrophages compared to WT cells (**Figures 7D,F**), confirming the results from lucigenin-dependent chemiluminescence assay. These results demonstrate that intracellular ROS generation by macrophages from p40^{*phox-/-*} mice was significantly impaired.

DISCUSSION

Salmonella infection is a major cause of human food-borne gastroenteritis worldwide. To effectively control the infection, molecules that can enhance the microbicidal capacity of phagocytes, such as phagocyte NADPH oxidase, are required. Individuals who have inherited deficiencies of NADPH oxidase show enhanced susceptibility to infection and develop CGD (16). Mice deficient for *Cybb* gene, encoding gp91^{phox}, the core component of the NADPH oxidase complex, failed to control infection with a normally avirulent bacterial infection (6). In this study, we determined the functional role of p40^{phox} in the effective control of both systemic and mucosal infection of Salmonella. Our results showed that defects in p40^{phox} in mice resulted in significantly increased morbidity and mortality during Salmonella infection. This is evidenced by the detection of higher tissue bacterial loads in the liver and spleen, supporting the role for p40^{phox} in restricting the penetration and growth of invading bacteria in intestinal mucosa, and enhanced level of bacteria-induced pro-inflammatory cytokine response. Moreover, we observed that p40^{phox} deficiency led to markedly increased recruitment of F4/80⁺ macrophages and Ly6G⁺ neutrophils in the infected tissues. Functional analysis of macrophages revealed that, in addition to the anticipated reduced ROS generation, the bacterial killing capacity of macrophages was significantly impaired in p40^{phox}-deficient mice and that p40^{phox}-deficient macrophages produced significantly more IL-1 β and TNF- α . These findings indicate that the p40^{phox} of NADPH oxidase plays an essential role in suppressing intracellular multiplication of Salmonella in macrophages and in the regulation of macrophage inflammatory response to bacterial infection.

Our results showed that p40^{phox} deficiency in mice significantly increased the bacteria-induced infiltration of macrophages and neutrophils into spleen and ceca mucosa (Figures 3 and 5), which correlated with an upregulation of expression of KC and MIP2 in the infected tissues, and yet, higher bacterial loads were detected in these tissues. In line with previous report showing that p40^{phox} deficiency in mice resulted in impaired neutrophil function (20), our functional analysis of macrophages from p40phox-deficient mice demonstrated functional defect of macrophages. This is evidenced by the results showing the elimination of Salmonella is significantly compromised in these cells. The uncontrolled bacterial infection and elevated bacterial loads may contribute to the observed enhanced pro-inflammatory cytokine production and exacerbated Salmonella-induced mucosal inflammation. This observation is in line with the observations in CGD patients, who have impaired ROS production and increased bacterial and fungal infections (16).

It was also reported that in the TNBS colitis model CGD mice $(Nox2^{-/-})$ developed an accentuated colitis (37), with exaggerated



and p40^{prox-/-}rmice. (A) Peritoneal macrophages were exposed to Salmonella enterica serovar Typhimurium (10 bacteria/cell) for 1 h. At different time points after infection (2, 6 h), the number of intracellular bacteria was determined by plating cell lysates onto LB plates supplemented with streptomycin. Results are expressed as mean \pm SEM (n = 5/group; NS = Not Significant; *p < 0.05). The data shown are representative of three experiments with similar results. (B) Immunofluorescence microscopy data show the number of internalized Salmonella in peritoneal macrophages (stained with Cy3 anti-F4/80) at 2 and 6 h after gentamicin treatment. Salmonella bacteria were detected with a rabbit antibody against Salmonella, followed by fluorescein isothiocyanate (FITC)-labeled anti-rabbit IgG antibody. (C) Real-time PCR analysis reveals that Salmonella LPS treatment (100 ng/ml) significantly upregulated the gene expression of IL-1 β and TNF- α in both WT and p40^{shox-/-} cells (n = 3-5/group; *p < 0.05; **p < 0.01; ***p < 0.001). The results are displayed as mean \pm SEM and are representative of three independent experiments.



neutrophil and macrophage infiltration and bacterial dissemination deep into the crypts (37). The observation of increased neutrophils and macrophages in Nox2^{-/-} mice has been suggested due to a functional failure of macrophages in clearance of neutrophils (38), as neutrophil numbers decrease during resolution of TNBS colitis as a result of macrophage clearance (39). New evidence indicated that the infiltration of neutrophils in TNBS colitis consumed sufficient O₂ to stabilize intestinal epithelial cell hypoxia-inducible factor (HIF), resulting in inflammation

resolution. In contrast to WT mice, CGD mice do not provoke mucosal hypoxia and exhibit an increased susceptibility to intestinal inflammation (37). It is clear that further investigation is needed for defining the potential role for $p40^{phox}$ in the regulation of kinetic and spatial distribution and migration of macrophage and PMN and mucosal integrity in the context of bacterial infection and inflammation.

After invasion of macrophages, *Salmonella* remain inside a membrane-bound compartment, the *Salmonella*-containing

vacuole (SCV). We have not directly measured ROS levels inside the SCV, a technically challenging task that requires a special *Salmonella* strain expressing a redox sensitive green fluorescent protein (40). However, the methods that we have used measure total cellular ROS and are widely used in the field. The fact that they clearly demonstrate a p40^{phox}-dependent decrease in ROS production that correlates with increased intracellular survival of *Salmonella* indicates their functional relevance. How p40^{phox} deficiency affects ROS levels inside the SCV remains to be determined.

Reactive oxygen species generation via NADPH oxidase by innate immune cells and the epithelium plays important role in host defense against infection. Although the current study focused on the role of p40^{phox} in innate immune cells in host protection against bacterial infection, epithelial cells also produce ROS, which participate in regulating intracellular signals for barrier repair and healing (41). In the GI tract, the main sources of ROS are NADPH oxidase enzymes. There are seven members in the family of ROS-generating NADPH oxidase, including NOX1-5 and two dual oxidases (DUOX1 and DUOX2). NOX2 is referred to as the phagocyte NADPH oxidase because it was first described in neutrophils and macrophages (42) and is a major ROS-producing enzyme with bactericidal activity (43). The main oxidases expressed along the GI tract are NOX1, often called "colon NADPH oxidase" due to the high expression in colon epithelium and DUOX2, which can be detected throughout all of the digestive tract, particularly in cecum and colon epithelium (44). DUOX2 is thought to participate in antimicrobial defense in the host mucosa (45-47). It has been shown that loss of functional DUOX enzymes lead to increased mucosal colonization by Helicobacter felis, enhanced shedding of bacterial antigens and virulence factors, and severe gastritis in Duoxa (Dual oxidase maturation factor) deficient mice (45). These results suggest an antimicrobial role for Duox at the apical surface in the mucus layer of the gastric epithelium controls growth of H. felis (45). Recently, two functionally altered NOX1 and DUOX2 variants in very early onset (VEO) IBD patients have been identified, both of which are associated with severe pancolitis and reduced ROS production in vitro (48, 49), suggesting that reduced epithelial ROS may provoke a pro-inflammatory phenotype similar to that in CGD patients. Interestingly however, a recent investigation using mice with complete or epithelium-restricted deficiency in Cyba, encoding the p22phox component of NADPH oxidase, showed that epithelial deficiency of Cyba resulted in protection from Citrobacter rodentium infection (50). In that study, it was also shown that intestinal ROS altered the gut microbiota composition, which is evidenced by the detection of enrichment of H₂O₂-producing bacterial strains in mice with epithelial Cyba deficiency. This study further showed that C. rodentium virulence can be attenuated by H2O2-mediated suppression of the virulenceassociated LEE pathogenicity island (50). Although a direct role for p40^{phox} in regulating intestinal epithelial cell protection against enteric bacterial infection needs to be investigated, these published data indicate the possibility that reduced intestinal epithelial ROS generation will influence mucosal responses to both commensal and pathogens, the pathogenesis of IBD and intestinal inflammation.

In both systemic and mucosal infection models utilized in the current study, hyperproduction of IL-1ß was detected in the infected tissues in $p40^{phox}$ -deficient mice (Figures 2 and 4). IL-1 β is produced as a biologically inactive precursor and the caspase-1-dependent IL-1ß secretion requires the activation of inflammasome, a cytosolic protein complex that senses microbial and endogenous stimuli, which is implicated in host protection and a variety of inflammatory diseases (51). Evidence also showed that PMN cells isolated from CGD patient had an increased caspase-1 activation and IL-1ß secretion with or without LPS-stimulation and that CGD patients had uncontrolled inflammation (28, 52). Our observations from the current study and those published results, therefore, suggest that normal NADPH oxidase activity not only plays an important role in killing bacteria but also in downregulating TLR signaling-mediated responses and IL-1ß production.

In summary, in the current study, we utilized the approaches that involved infecting mice with *Salmonella* that causes both systemic and mucosal infection, as well as *in vitro* infection of macrophages, to investigate the function of p40^{phox} in macrophages during bacterial infection. Our data demonstrate that p40^{phox} is critical for host resistance and survival during *Salmonella* infection and bacteria-mediated intestinal inflammation. One of the major complications of CGD is colitis, which may be treated with corticosteroid. A better understanding of the complex interaction between NADPH oxidase, immune defense, intestinal commensal, and enteric bacterial pathogens will provide important information for the development of new management strategies for disease conditions, such as IBD, cystic fibrosis, and so on.

ETHICS STATEMENT

All animal studies were carried out in accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals of the National Institutes of Health. The protocol was approved by the Sub-committee on Research Animal Care of Massachusetts General Hospital (animal welfare assurance number A3596-01).

AUTHOR CONTRIBUTIONS

All the authors reviewed and approved the final version of the manuscript and agreed to be accountable for the content of the work. YL and ML performed most experimental work, acquired and analyzed results, and YL wrote the first draft of the manuscript. CS, SL, and WZ performed experimental work and analyzed results. KC and RX generated and provided gene knockout animals, and analyzed and interpreted the data. WL, RX, and HS designed, supervised, and interpreted the experimental data, critically revised the manuscript. HS wrote the final version of the 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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