Activation of macrophage mediated host defense against Salmonella typhimurium by Morus alba L.

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Abstract

Background: The innate immune system plays a crucial role in the initiation and subsequent direction of adaptive immune responses, as well as in the removal of pathogens that have been targeted by an adaptive immune response.

Objective: Morus alba L. was reported to have immunostimulatory properties that might protect against infectious diseases. However, this possibility has not yet been explored. The present study investigated the protective and immune-enhancing ability of M. alba L. against infectious disease and the mechanisms involved.

Design: To investigate the immune-enhancing effects of M. alba L., we used a bacterial infection model.

Results and discussions: The lifespan of mice infected with a lethal dose of Salmonella typhimurium (1 × 10⁷ colony forming units – CFU) was significantly extended when they were administered M. alba L. Furthermore, M. alba L. activated macrophages, monocytes, and neutrophils and induced Th1 cytokines (IL-12, IFN-γ, TNF-α) in mice infected with a sublethal dose (1 × 10⁶ CFU) of S. typhimurium. M. alba L. significantly stimulated the uptake of bacteria into peritoneal macrophages as indicated by increased phagocytosis. Peritoneal macrophages derived from C3H/HeJ mice significantly inhibited M. alba L. induced NO production and TNF-α secretion compared with peritoneal macrophages derived from C3H/HeN mice.

Conclusions: These results suggest that the innate immune activity of M. alba L. against bacterial infection in mice occurs through activation of the TLR4 signaling pathway.

Keywords: Morus alba L; TLR4; Salmonella; immune defense; macrophage

Bacterial infectious agents are responsible for high morbidity and mortality in humans (1). Antibiotic drugs are used to combat these pathogens; however, the rise in antibiotic-resistant pathogens has led to the development of new therapeutic agents that are effective against these bacteria (2, 3).

The immune system is our natural defense system against pathogens such as viruses, bacteria, and other agents (4). It is composed of the innate immune system and the adaptive immune system. The innate immune system is a primary defense mechanism against invading organisms, while the adaptive immune system acts as a second line of defense. Both aspects of the immune system have cellular and humoral components that mediate their protective functions. Macrophages are the most abundant cells in granulomas and have been shown to play a key role throughout the course of infection in all infected hosts, including humans and non-human primates. Once these bacteria enter a host, Toll-like receptors (TLRs) recognize a variety of microbial products, including bacterial cell wall components and endocytosed nucleic acids, thereby triggering innate immune responses (5–7). Most laboratory studies have focused on the modulation of antimicrobial peptides and innate immune mediators by microbial components (8, 9). However, the effect of herbal plant extracts on immune responses to infectious diseases is not well established. An attempt has been made to identify a potent immunostimulator from plant extracts selected for their immune pharmacological properties. Our previous study showed that Morus alba L. had immunostimulatory effects in macrophages (10). In addition, M. alba L. might have an indirect anticancer effect by enhancing immune responses through TLR4 signaling (11). However, the protective effects of M. alba L. against pathogens have not been examined. In the present study, we investigated the protective effect and macrophage-mediated immune...
responses of *M. alba* L. in mice challenged with pathogenic *Salmonella typhimurium*.

**Materials and methods**

**Chemical and reagents**
Roswell Park Memorial Institute medium 1640 (RPMI) and fetal bovine serum (FBS) were obtained from GIBCO (Grand Island, NY, USA). A nitric oxide (NO) detection kit was obtained from INTRON Biotechnology (Sunnam, Korea). Trizol was obtained from Invitrogen (Carlsbad, CA, USA). Interleukin (IL) 6, IL-12, interferon gamma (IFN-γ) and tumor necrosis factor alpha (TNF-α) were obtained from R&D Systems (Minneapolis, MN, USA). Penicillin, streptomycin, neutral red, 3-(4,5-dimethylthiazolol-2-yl)-2,5-diphenyltetrazolium bromide, lipopolysaccharide (LPS), and all other chemicals were obtained from Sigma-Aldrich (St. Louis, MO, USA).

**Cell culture and animal care**
Peritoneal macrophages were prepared from BALB/c, C3H/HeJ (wild type) and C3H/HeN (TLR4 mutant) mice as described previously (12). RAW264.7 cells were routinely cultured in Dulbecco’s Modified Eagle Medium (DMEM; Sigma-Aldrich) supplemented with 10% heat-inactivated FBS, 100 U/mL penicillin, and 100 μg/mL of streptomycin. *Salmonella typhimurium* was grown on Luria-Bertani (LB) agar or in LB broth, where appropriate, at 37°C under aerobic conditions. Mice were housed in specific pathogen-free (SPF) conditions at 21–24°C and between 40 and 60% relative humidity with a 12 h light-dark cycle. All animals were acclimatized for at least 1 week prior to the start of experiments. All studies were performed in accordance with the guide for animal experimentation by Wonkwang University and approved by the university’s institutional animal care and use committee (Approval No. WKU11-21).

**Preparation of standardized *M. alba* L. extract**
Dried fruits of *M. alba* were purchased from a local herbal market in Jeonbuk, Korea. The fruits were pulverized into powder and extracted twice with hot water (80°C) for 5 h. The solvent was removed under reduced pressure in a RV10 rotary evaporator (IKI, Staufen, Germany) to yield *M. alba* fruit hot water extract (42.8%, w/v). The extract was dried to a powder and kept in a closed container until use. To avoid variations in activity for different preparations, sufficient extract was obtained in one batch for use throughout the study. The content of the marker chlorogenic acid in *M. alba* L. was quantitated using high performance liquid chromatography. Results indicated that *M. alba* L. possessed 1.18 mg of chlorogenic acid per 1.0 g of extract (Fig. 1).

**Induction of *S. typhimurium* infectious model**
SPF male ICR (Institute for Cancer Research) mice (5 weeks old and 25–27 g body weight) were purchased from Hanil Laboratory Animals (Iksan, Korea). The mice were randomly divided into two sets (Fig. 2). The first set contained three groups: a normal group (*n* = 10), *S. typhimurium*-infected group (*N* = 10), and 500 mg/kg/day *M. alba* L. + *S. typhimurium*-infected group (*n* = 10). The second set contained five groups: a normal group (*n* = 7), *S. typhimurium*-infected group (*N* = 7); 100, 300, and 500 mg/kg/day *M. alba* L. + *S. typhimurium*-infected group (*n* = 7, respectively). The normal and *S. typhimurium*-infected groups in these experiments were administered the same volume of distilled water. *M. alba* L. was dissolved in sterile double-distilled water and administered for 5 days consecutively. Twenty-four hours after the last drug administration, the mice of the first set were intraperitoneally injected with a lethal dose of bacterial suspension (10\(^7\) colony forming units (CFU)/mouse) to induce peritonitis. The mice in the second set were intraperitoneally injected with a sublethal dose of bacterial suspension (10\(^6\) CFU/mouse) to induce peritonitis. The challenged mice were fed a standard diet and water for 8 days and their survival and health were monitored twice per day throughout the experimental period. Any animal that exhibited severe clinical abnormalities, became moribund, or lost 20% of its initial body weight was sacrificed by ethyl ether narcosis and exsanguination. At the end of the experiment, the mice were subjected to ether anesthesia. The blood samples were collected from the central vein.

**Measurement of cytokines**
Blood and cell culture media were collected and then TNF-α, IL-6, IL-12, and IFN-γ production was measured by the sandwich ELISA method according to the manufacturer’s instructions (R&D Systems (TNF-α; DY410, IL-6; DY406, IL-12; DY419), Minneapolis, MN, USA and BD PharMingen (IFN-γ; RUO – 558258), CA, USA).

**Hematology test**
Blood samples were collected from the vena cava of all animals under ethyl ether anesthesia at necropsy for hematology. Blood was transferred to a container (EDTA K2; BD Biosciences, South Korea) for WBC (white blood cell), LYMPH (lymphocyte), MON (monocyte), and NEUTRO (neutrophil) counts using an HORIBA (Edison, NJ, USA).
Host defense activity of *Morus alba* L.

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confirmed in the final volume of 200 μL. After incubation at 37°C for 2 h, phagocytic cells were washed 3 times with phosphate-buffered saline (PBS), which was aspirated to remove non-ingested cell particles. Extracellular fluorescence was quenched by the addition of 100 μL of trypan blue. After 1 min, the FITC bacteria that had not been ingested washed away and the macrophages were rinsed twice with PBS. Next, the macrophages were lysed with lysis buffer (10 mM Tris-HCl, pH 7.5, 130 mM NaCl, 1% Triton X-100, 10 mM Na₂HPO₄, 10 mM Na₄P₂O₇). The fluorescence intensity (relative fluorescence unit) of bacteria inside the cells was determined at excitation

**Assay for peritoneal macrophage phagocytosis**

Fig. 1. The structure and high-performance liquid chromatography (HPLC) chromatographic profile of chlorogenic acid in *M. alba* L. extract at 340 nm.

**Fig. 2.** Time schedule of the experimental procedures.
and emission wavelengths of 480 and 520 nm by using a spectrofluorometer (Tecan Infinite F200, Männedorf, Switzerland). The relative phagocytic activity was calculated as the percentage of fluorescence intensity in a sample supplemented with FITC bacteria compared with that with no supplementation (control).

**Intracellular killing assay**

The intracellular killing assay was conducted as described by Wu et al. (14). RAW264.7 cells were seeded at a density of 5 × 10^4 cells/well in a 96-well plate containing 200 μL complete RAW medium and incubated at 37°C for 24 h in an atmosphere of 5% CO_2. Three colonies of each S. typhimurium strain were inoculated into 10 mL LB broth with appropriate antibiotics and incubated for 16 h at 37°C, with shaking at 200 rpm. Cultures were then centrifuged at 3,000 g for 10 min and resuspended in complete medium to 5 × 10^7 CFU/mL. The culture medium was removed from the RAW264.7 cells and replaced with 200 μL of this bacterial inoculum per well, to give a MOI of 10:1. This MOI was used as the frequency of infection and was high enough to enable observation of infection events at the single-cell level in subsequent imaging experiments but minimize Salmonella-induced macrophage death. The cells were then incubated at 37°C for 30 min in an atmosphere of 5% CO_2, after which the supernatant was removed from each well and 200 μL complete RAW medium containing 50 μg/mL gentamicin was added; plates were incubated for either 15 min (T0) or 2 h (T2) at 37°C for 30 min in an atmosphere of 5% CO_2. The cells were then washed twice with 200 μL DMEM and lysed in 1% saponin, and the number of intracellular bacteria (CFU) was determined by serial dilution and plating on plate count agar. Intracellular killing percentage was calculated as: [(T0 – T2)/T0] × 100.

**NO assay**

Peritoneal macrophages (1 × 10^5 cells/well) in a 96-well plate were incubated in the presence of three concentrations of M. alba L. (10, 30, and 100 μg/mL) for 24 h. NO was measured by determining the concentration of its stable oxidative metabolite nitrite using a microplate assay according to a described method (10). Supernatants (100 μL) were collected and mixed with an equal volume of Griess reagent (1% sulfanilamide and 0.1% N-(1-naphthyl)ethylenediamine dihydrochloride in 5% phosphoric acid) at room temperature for 15 min. The absorbance was read at 570 nm using a microplate reader. NaNO_2 was used as a standard.

**Statistical analysis**

Data are expressed as the mean ± SD and were examined for their statistical significance of difference by analysis of variance and Student’s t-test. A p-value < 0.05 was considered statistically significant.

## Results

**Effect of M. alba L. on pathogenic S. typhimurium in mice**

To determine the therapeutic effects of M. alba L. on life expectancy, mice were infected with a lethal dose (1 × 10^7 CFU) of S. typhimurium intraperitoneally and assessed for mortality. Figure 3 shows that the mortality rates in the normal and S. typhimurium groups were 100 and 0% on Day 8. By contrast, the groups treated with M. alba L. showed 60% had survived on Day 8. These observations demonstrate the potential of M. alba L. to protect mice against the lethal effects of S. typhimurium.

**Effects of M. alba L. on body and organ weight**

Body weight was monitored during the experiment to reflect the health status of mice. All the infected groups had significantly greater weight loss compared with normal mice. At Day 5, mice infected with S. typhimurium had lost significantly more weight compared with the M. alba L. group (Fig. 4). After sacrificing the animals, the organs were removed and weighed. No significant changes in the weights of livers were seen in normal, M. alba L. + S. typhimurium, or S. typhimurium mice (Table 1). However, spleen weights were significantly increased in animals infected with S. typhimurium compared with the normal group. The M. alba L. + S. typhimurium group had a significantly higher spleen weight than the S. typhimurium group (p < 0.05).

**Effects of M. alba L. on hematological parameters**

Table 2 shows the effect of M. alba L. on the hematological parameters of mice. WBC count was significantly increased in the M. alba L. + S. typhimurium group (p < 0.05) compared with the S. typhimurium group. Lymphocyte count was decreased in the M. alba L. + S. typhimurium group and infected control group. Monocyte and neutrophil counts were significantly increased in the M. alba L. + S. typhimurium group and infected control group compared with the uninfected control group (p < 0.05 for all).

**Effect of M. alba L. on cytokine production in S. typhimurium-infected mice**

To examine the immunomodulatory effect of M. alba L. on pathogenic S. typhimurium infection, the mice were administered M. alba L. (100, 300, and 500 mg/kg) for 5 days, or were not pretreated and were then challenged with 1 × 10^8 CFU/mouse (sublethal dose) of S. typhimurium. After 8 days, the levels of key immune molecules, IFN-γ, TNF-α, IL-12, and IL-6 were measured in serum. The mice fed with M. alba L. (300 or 500 mg/kg) had higher IFN-γ, IL-12, and TNF-α production compared with the infected group (Fig. 5a, c, d). However, IL-6 production was not significantly different in the serum of mice in all groups (Fig. 5b).
Fig. 3. Effects of *M. alba* L. on survival rates in mice challenged with pathogenic *S. typhimurium*. *Morus alba* L. (500 mg/kg) was orally administered for 5 days to mice, which were then intraperitoneally injected with a lethal dose of a bacterial suspension (10⁷ CFU/mouse) to induce peritonitis. The challenged mice were kept for 8 days on a standard diet with water and their survival was monitored twice a day throughout the experimental period.

Fig. 4. Effects of *M. alba* L. on change in body weight in mice challenged with pathogenic *S. typhimurium*. *Morus alba* L. (100, 300, or 500 mg/kg) was orally administrated for 5 days to mice, which were then intraperitoneally injected with a sublethal dose of a bacterial suspension (10⁵ CFU/mouse) to induce peritonitis. The body weight was recorded daily. The values are the means ± S.D. *p < 0.05, **p < 0.01, ***p < 0.001 compared with the *M. alba* L. group.
Table 1. Effects of *M. alba* L. on organ weight in mice challenged with pathogenic *S. typhimurium*. The values are the means ± S.D. *p < 0.05, compared with the *M. alba* L. group

| Group                        | Dose (mg/kg) | Body weight (g) | Relative liver weight (%) | Relative Spleen weight (%) |
|------------------------------|--------------|-----------------|---------------------------|----------------------------|
| Normal mice                  |              | 36.0 ± 1.41     | 5.46 ± 0.08               | 0.35 ± 0.08                |
| *Mice infected with S. typhimurium (1 × 10⁵ CFU/mouse)* |              |                 |                           |                            |
| Saline                       | -            | 25.8 ± 1.18     | 6.88 ± 0.20               | 1.16 ± 0.08                |
| *Morus alba* L. 100          | 27.6 ± 0.47  | 7.40 ± 0.64     | 1.41 ± 0.09 *             |                            |
| *Morus alba* L. 300          | 26.4 ± 2.46  | 7.91 ± 0.11     | 1.55 ± 0.11 *             |                            |
| *Morus alba* L. 500          | 27.0 ± 3.57  | 7.40 ± 0.93     | 1.60 ± 0.04 *             |                            |

WBC, white blood cells; LYM, lymphocytes; MON, monocytes; NEU, neutrophils.

**Table 2.** Effects of *M. alba* L. on hematological parameters in mice challenged with pathogenic *S. typhimurium*. The values are the means ± S.D. *p < 0.05, compared with the *M. alba* L. group

| Group                        | Dose (mg/kg) | WBC (10³/ml) | LYM (%) | MON (%) | NEU (%) |
|------------------------------|--------------|--------------|---------|---------|---------|
| Normal mice                  |              |              |         |         |         |
| Saline                       | -            | 4.32 ± 0.19  | 76.3 ± 10.28 | 18.4 ± 2.32 | 6.15 ± 3.27 |
| *Mice infected with S. typhimurium (1 × 10⁵ CFU/mouse)* |              |              |         |         |         |
| Saline                       | -            | 1.78 ± 0.24  | 61.3 ± 7.25  | 23.8 ± 1.93 | 16.1 ± 1.27 |
| *Morus alba* L. 100          | 100          | 1.93 ± 0.27  | 62.7 ± 8.93  | 25.4 ± 1.42 | 13.6 ± 1.95 |
| *Morus alba* L. 300          | 300          | 4.22 ± 0.43 * | 55.4 ± 11.75 | 29.5 ± 1.28 * | 22.7 ± 2.95 |
| *Morus alba* L. 500          | 500          | 5.91 ± 1.73 * | 53.4 ± 7.32  | 28.7 ± 1.15 * | 25.1 ± 1.87 * |

**Fig. 5.** Effect of *M. alba* L. on the production of cytokines in mice challenged with pathogenic *S. typhimurium*. *Morus alba* L. (100, 300, or 500 mg/kg) was orally administrated for 5 days to mice, which were then intraperitoneally injected with a sublethal dose of a bacterial suspension (10⁵ CFU/mouse) to induce peritonitis. At the end of the experiment, the mice were subjected to ether anesthesia. Serum was collected from the blood of infected mice pre-administered with *M. alba* L. to evaluate the serum levels of (a) IFN-γ, (b) IL-6, (c) IL-12, and (d) TNF-α. The data are the means ± SD, *p < 0.05* compared with the *S. typhimurium* group.
Discussion

Morus alba L., white mulberry, is a deciduous tree that belongs to the Moraceae family, which is widely distributed in Asia. Morus alba L. has been used worldwide in traditional medicine for the treatment of various diseases from ancient times to the present (12, 15). Previous studies have indicated that Morus alba L. possesses various pharmacological properties, including antimicrobial (16), antioxidant (17), antitumor (18), anti-obesity (19), antihypotensive (20), neuroprotective (21, 22), and antidiabetic (23, 24) actions. In addition, our previous studies indicated that the extracts and components of Morus alba L. fruit water extract were safe (25), immunostimulatory (10, 12, 26), and had indirect anticancer activity by enhancing immune responses mediated by TLR4 signaling (12). However, the effects and mechanisms involved in pathogen defense actions have remained elusive. Therefore, in this study, we examined the pathogen defense method commonly used to test intracellular bactericidal activity with live Salmonella as the target. The intracellular killing of macrophages was increased by Morus alba L. treatment in a dose-dependent manner (Fig. 6b), which demonstrated that Morus alba L. enhanced macrophage intracellular killing.

Effects of Morus alba L. on peritoneal macrophages from TLR4-deficient mice

To investigate the involvement of TLR4 signaling in mediating the beneficial effects of Morus alba L., we compared NO and TNF-α production in peritoneal macrophages from wild-type (C3H/HeN) or TLR4-deficient mice (C3H/HeJ). Significantly increased NO and TNF-α was detected in peritoneal macrophages from C3H/HeN but not C3H/HeJ mice after Morus alba L. treatment (Fig. 7). Therefore, Morus alba L. induces the production of proinflammatory cytokines via TLR4.

Fig. 6. Effects of Morus alba L. on phagocytic activity and intracellular killing. Peritoneal macrophages derived from BALB/c mice or RAW 264.7 cells were treated with 10, 30, or 100 μg/mL of Morus alba L. for 24 h, or with LPS (1 μg/mL) as a positive control. (a) After the addition of FITC-labeled E. coli, the cells were incubated for 2 h. The supernatants containing unphagocytosed bacteria were removed and then lysed with Triton X-100 containing lysis buffer. Fluorescence intensities of the lysed cells were measured by using a fluorescence microplate reader. (b) After the addition of S. typhimurium, the cells were incubated for 2 h. Unphagocytosed bacteria were removed and the cells were lysed with 1% saponin. The number of intracellular bacteria (CFU) was determined by serial dilution and plating on plate count agar (PCA). The values are the mean ± S.D. **p < 0.01, ***p < 0.001 compared with the untreated group.

Fig. 7. Effects of Morus alba L. on peritoneal macrophages derived from C3H/HeN and C3H/HeJ mice. Peritoneal macrophages from C3H/HeN and C3H/HeJ mice were stimulated with Morus alba L. or LPS (1 μg/mL) as a positive control for 24 h. (a) NO and (b) TNF-α levels were determined by ELISA. The values are the means ± S.D. **p < 0.01, ***p < 0.001, compared with C3H/HeN mice.
properties of *M. alba* in mice challenged with pathogenic *S. typhimurium*.

Several studies have suggested that host cells exposed to different groups of pathogens respond with common transcriptional activation programs, referred to as the *core response to infection* (27). Macrophages, monocytes, and granulocytes are involved in the non-specific defense mechanism against pathogens. The common response of macrophages to bacterial infections involves the upregulation of genes involved in M1 polarization (28).

We report that *M. alba* L. enhanced immune activity and resistance to experimental challenge with *S. typhimurium* infection. However, *M. alba* L. did not directly kill *S. typhimurium*. Therefore, the pathogen defense effect of *M. alba* L. is likely to be mediated by potentiation of the host’s defense system, such as macrophage activation, rather than by the indirect inhibition of bacterial growth.

We observed hepatomegaly and splenomegaly in *S. typhimurium* infected mice. The present results showed that the killing activity of monocytes and neutrophils was increased in *M. alba* L. treated mice infected with *S. typhimurium*, indicating its potential to enhance non-specific immune responses. Systemic *Salmonella* infection commonly induces splenomegaly in murine or human hosts. The capacity of *Salmonella* to evade killing and replicate within tissue phagocytes means that infection can quickly overwhelm local host defenses at the site of infection. Thus, inflammatory signals produced by cells in the infected tissue rapidly recruit neutrophils and inflammatory monocytes to engulf replicating bacteria. Splenomegaly can cause splenic structure change; splenic tissue structure is correlated with the recruitment and expansion of T and B cells and macrophages (29). The present study provides evidence of the induction of the macrophage phagocytic effect by *M. alba* L.

Furthermore, cytokines such as IL-12, IFN-γ, and TNF-α have a vital role in both innate and adaptive immunity and are responsible for Th1 polarization (30, 31). Because IL-12, IFN-γ, and TNF-α were upregulated in *M. alba* L. treated groups, this suggests it has important immunoregulatory activity by activating macrophages. However, Th2 cytokine production (IL-6) was unaffected. IL-12 is produced primarily by monocytes, macrophages, and other antigen-presenting cells and is essential for fighting infectious diseases and cancer. IL-12 synergizes with TNF-α and other proinflammatory cytokines to stimulate IFN-γ production, as well NK and CD8 T cell cytotoxicity (32). TNF-α is another proinflammatory cytokine and its levels in plasma are directly correlated with the ability of phagocytes to generate superoxide and the activity of iNOS and, thus, NO levels (31). One of the key requirements for the accuracy of the in vitro bactericidal assay is to eliminate the bacteria remaining extracellularly after ingestion by macrophages. For this purpose, gentamicin at a concentration of c. 50 mg/L has been widely used, as this antibiotic is believed to be incapable of penetrating the macrophage membrane. However, there are reports that such a high concentration of gentamicin penetrates HeLa cells, reaching c. 90% of the extracellular concentration after incubation for 72 h. The M1, or proinflammatory, macrophage phenotype is characterized by high levels of proinflammatory cytokines and reactive nitrogen and oxygen intermediates, the promotion of a Th1 response, and strong microbicidal and tumoricidal activity. Phagocytosis is the first step in the macrophage response to invading microorganisms and activation of phagocytosis enhances innate immune responses (33). Phagocytosis by macrophages was increased by *M. alba* L. treatment in a dose-dependent manner. The activation of intracellular killing mechanisms is critical to the antimicrobial activity of phagocytes. The pathogen internalization is followed by the fusion of phagosomes and lysosomes, which exposes the internalized pathogen to antimicrobial proteins and reactive molecules. Results indicate that *M. alba* L. is armed with such a killing capacity in a dose-dependent manner.

The first step in the modulation of cellular events is binding to receptors. TLRs play a central role in macrophage activation and the control of pathogen infections (5).

TLR4 is expressed on macrophages, dendritic cells, B cells, T cells, and endothelial cells. The role of TLR as a *M. alba* L. receptor was confirmed in our previous study (10, 12). TLR4 binds with *M. alba* L. activated signaling pathways, including MAPKs and NF-κB. Activation of MAPKs is required for the induction of NO as it controls the activation of NF-κB. In this study, the role of TLR4 as an *M. alba* L. receptor was confirmed in macrophages. To investigate this membrane receptor, we examined the effect of *M. alba* L. on primary macrophages isolated from wild-type C3H/HeN and C3H/HeJ mice that have mutant TLR4. *Morus alba* L. induced NO production and TNF-α in macrophages from C3H/HeN, but not from TLR4 mutated C3H/HeJ mice, which suggests that TLR4 is the membrane receptor for *M. alba* L.

In summary, our data suggest that *M. alba* L. activated macrophages via TLR4 to provide protection against *Salmonella* infection, indicating that *M. alba* L. modulates the effector functions of immunocompetent cells.

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**Authors’ contributions**

All authors read and approved the final version of the manuscript. S.Y.K. conceived and designed the experiments. B.Y.C. and H.C.L. performed the experiments. B.Y.C., J.S.O., and S.Y.K. analyzed the
data. S.Y.K. and B.Y.C. drafted the manuscript. B.Y.C., B.S.K., H.C.L., J.S.O., and S.Y.K. revised the manuscript.

Conflicts of interest and funding

The authors declare that they have no competing interests.

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