


# *Helicobacter pylori* attenuates lipopolysaccharide-induced nitric oxide production by murine macrophages

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## Abstract

Intra-gastric growth of *Helicobacter pylori* and non-*Helicobacter* microorganisms is thought to be associated with elevated levels of pro-inflammatory cytokines and the production of NO these effects can lead to chronic inflammation. Microorganisms can activate the expression of iNOS and the production of NO by macrophages through stimulation with bacterial LPS. *Helicobacter pylori* can evade these vigorous immune responses, but the underlying mechanism remains unknown. In this study, we used a murine model of macrophage infection to demonstrate that *H. pylori* inhibits LPS-induced expression of iNOS and production of NO by macrophages. Suppression of LPS-induced NO production by macrophages led to elevated survival of *H. pylori* in a trans-well system. This effect was abrogated in macrophages from iNOS<sup>-/-</sup> mice. Analysis of iNOS mRNA and protein levels revealed that *H. pylori* inhibits iNOS expression at both transcriptional and post-transcriptional levels, and that these effects occurred with live bacteria. Furthermore, the effect of *H. pylori* involved down-regulation of the mitogen-activated protein kinase pathway and the translocation of active NF-κB into the nucleus. Taken together, our results reveal a new mechanism by which *H. pylori* modulates the innate immune responses of the host and maintains a persistent infection within the stomach.

## Keywords

Lipopolysaccharide, nitric oxide, *Helicobacter pylori*, macrophage, NF-κB

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## Introduction

*Helicobacter pylori* is the most common causative agent of gastrointestinal disease in humans. Infection with this pathogen usually occurs in childhood and the bacteria can persist in the stomach for the lifetime of an individual.<sup>1,2</sup> Persistent infection with *H. pylori* in the gastric mucosa induces the expression of NF-κB and the secretion of pro-inflammatory cytokines, including IL-1β, IL-6, IL-8, and TNF-α.<sup>3,4</sup> Other inflammatory mediators, such as NO, a bactericidal agent generated by inducible nitric oxide synthase (iNOS) during the conversion of L-arginine to L-citrulline, are activated by *H. pylori* infection in both macrophages<sup>5</sup> and the gastric epithelium.<sup>6</sup> These findings indicate that *H. pylori* is an important factor for the induction of pro-inflammatory cytokines and NO in the host stomach.

Nitric oxide is derived from iNOS in LPS-activated macrophages during inflammatory responses. Following

treatment of macrophages with LPS, the NF-κB heterodimer rapidly translocates to the nucleus where it

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activates the transcription of target genes, including iNOS and several pro-inflammatory cytokines.<sup>7</sup> In addition, p38 MAPK, protein kinase C and ERK are also involved in the activation of NF- $\kappa$ B and the expression of iNOS in response to LPS.<sup>8,9</sup> Several microorganisms disrupt the activation of MAPKs or the NF- $\kappa$ B signalling pathway in macrophages to evade immune attack.<sup>10–14</sup> The effect of *H. pylori* on the modulation of LPS-activated molecules in macrophages remains unknown.

In addition to *H. pylori*, non-*Helicobacter* microorganisms are found in the gastric environment.<sup>15,16</sup> One study reported that *H. pylori*-associated gastritis was associated with the presence of several other microbes in the stomach, including *Enterococcus*, *Pseudomonas*, *Streptococcus*, *Staphylococcus* and *Stomacoccus*.<sup>17</sup> A more recent study identified 128 phylotypes in 23 gastric biopsy samples; however, the presence of *H. pylori* did not affect the composition of microbiota in the gastric microbial community.<sup>18</sup> These findings indicate that *H. pylori* and non-*Helicobacter* microorganisms are present in the microbiota of the human stomach, and these microbes can elicit pro-inflammatory mediators and induce vigorous immune responses.<sup>19</sup> These findings also raise the question of how *H. pylori* persists in the microbial ecosystem under the harsh environment of the stomach.

The aim of the present study was to address the question of how *H. pylori* evades the vigorous antimicrobial activities of macrophages. We established an *in vitro* murine model system and an *ex vivo* murine model system to examine whether this bacterium could suppress LPS-induced NO production through the MAPK or the NF- $\kappa$ B signalling pathways. We showed that *H. pylori* inhibits iNOS expression and NO production by murine macrophages stimulated with a high dose of LPS. We further demonstrated that *H. pylori* down-regulates the LPS-induced activation of phosphorylated p38, ERK1/2 and NF- $\kappa$ B, and it subsequently suppresses LPS-induced macrophage responses. Thus, our study revealed that *H. pylori* attenuates LPS-induced NO production in macrophages and consequently evades early host immune responses.

## Materials and methods

### Antibodies and reagents

Polyclonal rabbit anti-iNOS, anti-phosphorylated c-Jun-N-terminal kinase (p-JNK), and anti- $\alpha$ -tubulin Abs were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Mouse mAbs specific for p38 MAPK, JNK, and p44/42 (ERK1/2) were purchased from Cell Signaling (Beverly, MA, USA). Mouse monoclonal anti-phosphorylated p38 MAPK,

and the anti-phosphorylated ERK1/2 (Thr185/Tyr187) Abs were purchased from Upstate (Billerica, MA, USA). LPS (*Escherichia coli* O55: B5) and amino-guanidine hemisulfate (AG) were purchased from Sigma-Aldrich (St Louis, MO, USA). SB203580 (p38 inhibitor), PD98059 (ERK inhibitor) and SP600125 (JNK inhibitor) were purchased from Calbiochem (San Diego, CA, USA). The activator protein (AP)-1-Luc and NF- $\kappa$ B-Luc plasmids were purchased from Stratagene (San Diego, CA, USA). The iNOS promoter construct (piNOSLuc) was a kind gift from Dr E. A. Ratovitski (Johns Hopkins University, Baltimore, MD, USA). The pSV- $\beta$ -galactosidase vector and the luciferase assay kit were purchased from Promega (Madison, WI, USA). All other reagents were obtained from Sigma-Aldrich.

### Bacterial strains, cell culture and mice

*Helicobacter pylori* 26695 (ATCC 700392) was used as a reference strain. The *cagA* or *vacA* isogenic mutants derived from *H. pylori* 26695 were constructed as described.<sup>20</sup> *Helicobacter pylori* strains were recovered from frozen stocks on Brucella agar plates (Becton Dickinson, Franklin Lakes, NJ, USA) containing 10% sheep blood. *Helicobacter pylori* strains were stored and cultivated as described,<sup>21</sup> and *H. pylori* extracts were prepared as described.<sup>22</sup> Heat-killed *H. pylori* was obtained by boiling  $1 \times 10^9$ /ml of bacteria suspended in PBS for 30 min. Crude *H. pylori* extracts were prepared by sonicating  $1 \times 10^9$ /ml of bacteria suspended in PBS for 5 min on ice. Crude extracts were then centrifuged at 16,000 g for 5 min at 4°C. The supernatant was filtered through a 0.22  $\mu$ m filter and used for further analysis.

RAW 264.7 cells (ATCC TIB-71) were cultured in Roswell Park Memorial Institute (RPMI) 1640 medium (Invitrogen, Carlsbad, CA, USA). De-complement fetal bovine serum (10%; HyClone, Logan, UT, USA) was added to the culture medium. For bacterial infection experiments, the cell culture medium was not supplemented with antibiotics.

Male wild-type C3H/HeN and TLR4-deficient C3H/HeJ mice aged 6–8 wk were kindly provided by Dr Ai-Li Shiau (Departments of Microbiology and Immunology, National Cheng Kung University Medical College, Taiwan). C57BL/6 iNOS knockout (C57BL/6-Nos<sup>tm1Lau</sup>) (iNOS<sup>-/-</sup>) and wild-type mice aged 6–8 wk were kindly provided by Dr Ming-Chei Maa (Graduate Institute of Basic Medical Science, China Medical University, Taiwan). Mice were maintained at the animal centre of China Medical University (Taichung, Taiwan). All procedures were performed according to the 'Guide for the Care and Use of Laboratory Animals' (National Research Council, USA) and were approved by the animal experiment committee of China Medical University.

### Preparation of murine peritoneal exudate macrophages

C57BL/6iNOS knockout (C57BL/6-Nos<sup>tm1Lau</sup>) (iNOS<sup>-/-</sup>) and wild-type mice of the same age and gender were used to assess the role of iNOS in *H. pylori*-induced suppression of LPS-induced NO production by macrophages. Murine peritoneal exudate macrophages (PEMs) were obtained after euthanasia by lavaging each mouse with 10 ml of cold PBS three days after intraperitoneal injection of 2 ml of 3% thioglycolate in PBS. Two h after seeding the cells in culture plates, the non-adherent cells were removed by washing with PBS and the adherent cells were used for further experiments.

### Mouse inoculations

C3H/HeN (n = 6) and C3H/HeJ (n = 6) mice aged 6–8 wk were intragastrically inoculated with *H. pylori*. All mice were maintained under fasting for 24 h before inoculation. The protocol of administration of mouse with LPS was performed as described with slight modifications.<sup>23</sup> Each mouse was administered  $1 \times 10^9$  CFU/ml of *H. pylori* and purified LPS (75 µg, phenol extracted from *E. coli* O55: B5, Sigma-Aldrich) by intragastric gavage for three consecutive days. Six h after the final inoculation with *H. pylori*, the mice were fed with standard food and water and housed for 1 wk. On the seventh day after infection, six mice in each group were sacrificed and the number of *H. pylori* in their stomachs was determined by plating on *Brucella* blood agar plates and expressed as CFU/g tissue.

### Immunoblotting

*H. pylori*-infected cells were washed three times with PBS and boiled in SDS-PAGE sample buffer for 10 min. The samples were then resolved by 10% SDS-PAGE and transferred onto polyvinylidene difluoride (PVDF) membranes (Millipore, Billerica, MA, USA). The membranes were incubated with primary Abs and then with horseradish peroxidase-conjugated secondary Abs (Invitrogen). The proteins of interest were visualized with ECL<sup>TM</sup> Western blotting reagents (GE Healthcare, Buckinghamshire, UK) and were detected by exposure to X-ray film (Kodak, Boca Raton, FL, USA).

### Reverse transcription and quantitative real time-PCR

Total RNA was extracted from PEMs using TRIzol reagent (Invitrogen), and 1 µg of total RNA was reverse transcribed into cDNA using the oligo(dT) primer. Quantitative real-time PCR using SYBR Green I Master Mix and a model 7900 Sequence Detector System was conducted according to the manufacturer's instructions (Applied Biosystems, Foster City, CA,

USA). After pre-incubation at 50 °C for 2 min and 95 °C for 10 min, PCR was performed with 40 cycles of 95 °C for 10 s and 60 °C for 1 min. The threshold was set above the non-template control background and within the linear phase of target gene amplification in order to calculate the cycle number at which the transcript was detected (denoted as C<sub>T</sub>). The oligonucleotide primers were: iNOS, forward, 5'-CCCAGAGTTCCA-GCTTCTGG-3', and reverse, 5'-CCAAGCCCCTCA-CCATTATCT-3'; and GAPDH, forward, 5'-CTCA-ACTACATGGTCTACATGTTCCA-3', and reverse, 5'-CTTCCCATTCTCAGCCTTGACT-3'.

### Bacterial survival assay

Bacterial survival was assessed in cultures of *H. pylori*-exposed, LPS-treated macrophages using a trans-well system, as described<sup>22</sup> with slight modification. Briefly, murine PEMs were cultured in the bottom layer of trans-well plates (Corning, Corning, NY, USA). After 48 h,  $1 \times 10^6$  *H. pylori* were added to the insert membrane (0.1 µm pore size) and co-incubated for an additional 6 h in culture. The bacteria on the insert membrane were then resuspended and cultured by serial dilution onto *Brucella* blood agar plates. Colonies were counted after 4 to 5 d of incubation. The CFUs were used to determine anti-bacterial effects.

### Determination of NO production and cell viability assay

The NO production was estimated from the accumulation of nitrite (NO<sub>2</sub><sup>-</sup>), a stable end product of NO metabolism, in the culture medium, using the Griess reagent (Sigma-Aldrich).<sup>24</sup> The MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] assay was used to measure the effects of LPS and *H. pylori* on macrophage viability.<sup>25</sup> RAW 264.7 cells or PEMs were infected with various multiplicities of infection (MOI) of *H. pylori* for 24 or 48 h, respectively. Cell viability was then measured by examining the ability of viable cells to chemically reduce MTT to formazan, which was quantified by measurement of optical density at 570 nm.

### Transfection and reporter gene assay

RAW 264.7 cells were grown to 90% confluency in a 12-well plate and transfected with NF-κB-Luc, AP-1-Luc, or iNOS-Luc reporter plasmid using Lipofectamine 2000 (Invitrogen).<sup>26,27</sup> After 24 h, cells were incubated without or with LPS and then infected with *H. pylori* during an additional 24 h culture. To prepare cell lysates, 100 µl of reporter lysis buffer (Promega) was added to each well and cells were scraped from dishes. An equal volume of luciferase substrate was added to all samples and luminescence was measured

using a microplate luminometer (Biotek, Winooski, VT, USA). Luciferase activity was normalized to the transfection efficiency as determined by co-transfection of the  $\beta$ -galactosidase expression vector (Promega).<sup>28</sup>

### Immunofluorescence labeling of phosphorylated p65

To visualize *H. pylori*-induced inhibition of the translocation of phosphorylated p65 into the nucleus of macrophages, RAW 264.7 cells were seeded onto cover-slips and treated without or with LPS for 2 h and then with *H. pylori* for an additional incubation at 37°C for 1 h. Cells were fixed in 3.7% (w/v) paraformaldehyde and permeabilized with 0.5% (v/v) Triton X-100 in PBS for 2 min. For labeling of p65, cells were incubated for 30 min with rabbit polyclonal anti-p65 (H-286; Santa Cruz Biotechnology) and propidium iodide (Calbiochem). Cells were then incubated with a secondary antibody, fluorescein isothiocyanate-conjugated anti-mouse IgG (Chemicon) and they were fixed in paraformaldehyde. Fixed cells were mounted and observed with a confocal laser scanning microscope (Zeiss LSM 510, Carl Zeiss, Göttingen, Germany). The quantification of fluorescence intensity for p65 was analyzed by ZEN software (Carl Zeiss).

### Statistical analysis

The Student's *t*-test was used to calculate statistical significance; a *P* value of <0.05 was considered significant.

## Results

### *Helicobacter pylori* inhibits LPS-induced NO production by macrophages

To assess whether *H. pylori* inhibits LPS-induced NO production by macrophages, mouse macrophage RAW 264.7 cells were cultured with LPS (2  $\mu$ g/ml) and infected with *H. pylori* at a MOI of 0 to 100 for 24 h. NO production, measured by nitrite levels, was not suppressed when LPS-stimulated RAW 264.7 cells were infected with *H. pylori* at a low MOI of 10 (Figure 1A). However, NO production by LPS-stimulated cells was decreased at a MOI of 50–100. When cells were infected with *H. pylori* at a MOI of 100, LPS-induced NO production was reduced by approximately 50%. Macrophage viability, as determined by the MTT assay, did not change after 24 h of incubation with LPS and *H. pylori* (Figure 1B).

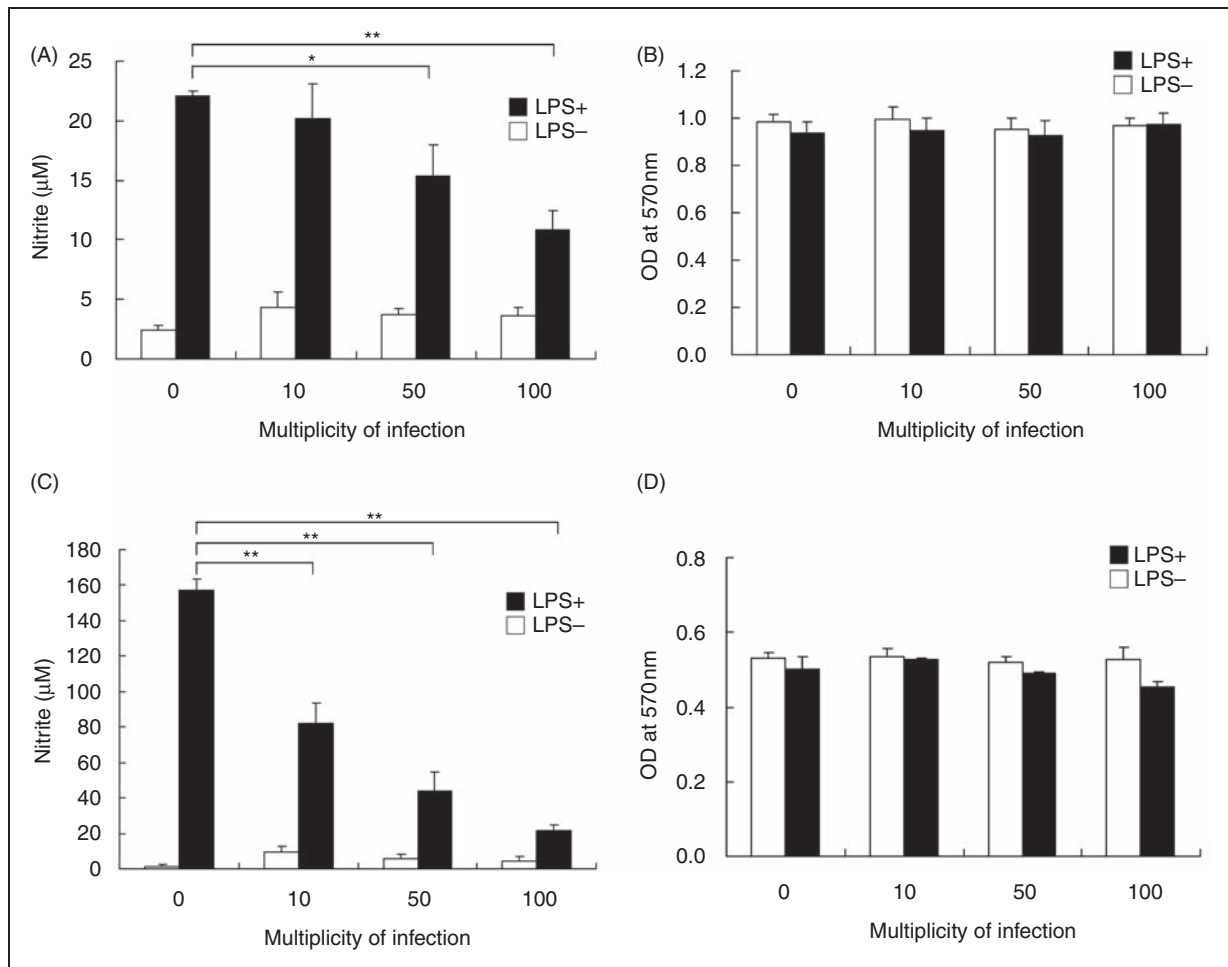
To further delineate the suppressive effects of *H. pylori* on LPS-induced NO production, PEMs were prepared and co-incubated with LPS and *H. pylori* for 48 h. This *ex vivo* approach showed that LPS treatment induced NO production in uninfected

PEMs (MOI of 0; Figure 1C), and LPS-induced NO production was reduced in a MOI-dependent manner when PEMs were infected with *H. pylori* (Figure 1C). The viability of PEMs, as determined using the MTT assay, was hardly influenced by treatment with LPS and *H. pylori* at different MOIs (Figure 1D). Thus, *H. pylori* inhibited, in a MOI-dependent manner, LPS-induced NO production, not only in RAW 264.7 cells, but also in murine primary PEMs, and PEMs were more sensitive than the macrophage cell line to the suppressive effects of *H. pylori*.

### Live *Helicobacter pylori* is essential for the inhibition of LPS-induced NO production

To determine the functional role of *H. pylori* in inhibiting LPS-induced NO production by RAW 264.7 cells, live bacteria, heat-killed bacteria and crude bacterial extracts were tested for their ability to inhibit LPS-induced NO production. As shown in Figure 2A, live *H. pylori* attenuated LPS-induced NO production by RAW 264.7 cells. In contrast to the effects observed with live bacteria, neither heat-killed bacteria nor crude bacterial extracts inhibited LPS-induced NO production. We also used PEMs to study the effects of *H. pylori* on the inhibition of LPS-induced NO production. Consistent with the results for RAW 264.7 cells, LPS-induced NO production by PEMs was inhibited by live *H. pylori* but not by heat-killed bacteria or crude bacterial extracts (Figure 2B). We further investigated the *H. pylori*-derived components that are responsible for these effects. Virulence factor isogenic mutants of *H. pylori*, *ΔcagA* and *ΔvacA*, were tested for their ability to inhibit LPS-induced NO production by RAW 264.7 cells. Both isogenic mutants suppressed LPS-induced NO production, similar to that observed with wild-type *H. pylori* (Figure 3).

To mimic the environment of the bacterial infection, we further analyzed the ability of *H. pylori* to inhibit LPS-induced NO production and to enhance the survival of bacteria adjacent to infected macrophages *in vitro*. Using a trans-well culture system to assay bacterial survival of LPS-stimulated murine PEMs, we found that *H. pylori* at a MOI of 100 increased bacterial viability (Figure 4A). To test the hypothesis that *H. pylori* triggers iNOS activity, which is necessary for NO production, we tested the effects of *H. pylori* in C57BL/6 iNOS<sup>-/-</sup> mice. *Helicobacter pylori* viability was similar with or without LPS and with *H. pylori* at a MOI of 0 or 100 in PEMs from iNOS<sup>-/-</sup> mice, unlike the effects observed in wild-type mice (Figure 4B). These results revealed that the ability of *H. pylori* to inhibit LPS-induced NO production resulted from a reduction in the antimicrobial activity of macrophages with subsequent enhancement of *H. pylori* survival.



**Figure 1.** *Helicobacter pylori*-mediated inhibition of LPS-induced NO production in the RAW 264.7 cell line (A) and in murine primary PEMs; (C). Cells were treated without or with LPS (2 μg/ml) and infected with *H. pylori* at various MOIs from 0 to 100. After 24 h (RAW 264.7 cells) or 48 h (PEMs) incubation, the culture supernatants were collected for determination of nitrite levels using the Griess reagent. The MTT assay showed that there was no loss of cell viability in RAW 264.7 cells (B) or murine PEMs (D) during the incubation period. The data represent the mean ± standard deviation derived from three independent experiments. Statistical significance was determined using the Student's *t*-test (\**P* < 0.05; \*\**P* < 0.01).

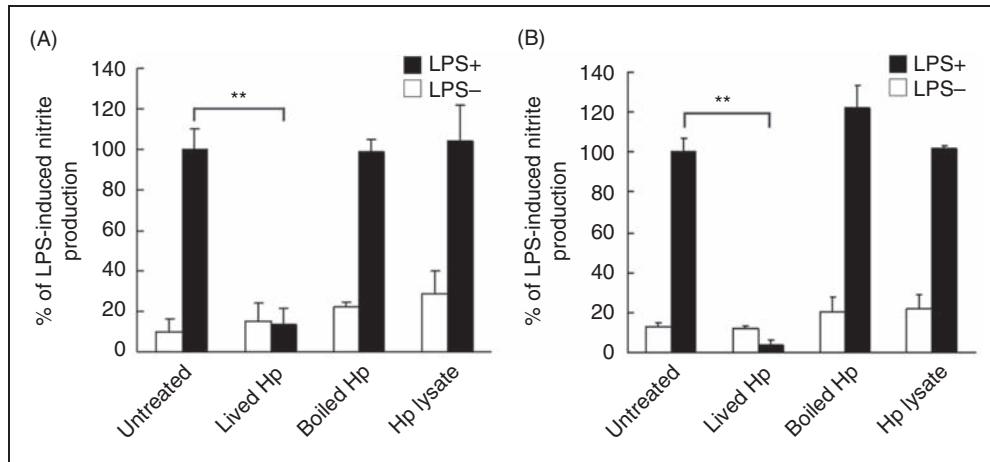
### *Helicobacter pylori* inhibits LPS-induced iNOS expression at transcriptional and post-transcriptional levels

To investigate the effects of *H. pylori* on the regulation of iNOS expression, PEMs were treated (or left untreated) with LPS and exposed to *H. pylori* at various MOI for 48 h. Subsequently, iNOS protein levels were analyzed by Western blotting. *Helicobacter pylori* infection led to a significant decrease, in a MOI-dependent manner, in LPS-induced iNOS level (Figure 5A). When cells were infected with *H. pylori* at a MOI of 100, LPS-induced iNOS expression decreased by approximately 80%. We then measured expression of iNOS mRNA using reverse transcription and quantitative real-time PCR in PEMs that were co-cultured without or with LPS and with *H. pylori* at various MOI for 6 h. The expression of

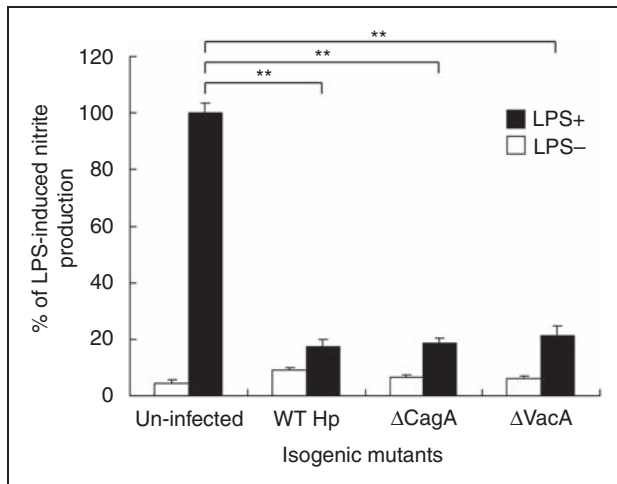
iNOS mRNA was decreased by *H. pylori* infection at a MOI of 50 to 100 (Figure 5B). Taken together, the results indicated that *H. pylori* inhibited iNOS transcription, which subsequently influenced the translation of iNOS mRNA and NO production by macrophages.

### *Helicobacter pylori* attenuates LPS-induced NO production by macrophages through the p38 and ERK 1/2 signalling pathways

The LPS-induced NO production by macrophages involves several signalling pathways, including p38, ERK 1/2 (p42 and p44), and JNK. We used Western blotting to analyze the signal transduction pathways involved in the inhibitory effects of *H. pylori* on NO production by LPS-stimulated PEMs for 60 min of culture. The data showed that without treatment with LPS



**Figure 2.** Live *H. pylori* are essential for inhibition of LPS-induced NO production by macrophages. Cells were treated without or with LPS and were either infected or un-infected with live *H. pylori* (live Hp), heat-killed *H. pylori* (boiled Hp) or crude extracts prepared from *H. pylori* (Hp lysate). Live and heat-killed *H. pylori* were used at a multiplicity of infection (MOI) of 100. After RAW 264.7 cells were cultured for 24 h (A) or murine PEMs were cultured for 48 h (B), culture supernatants were collected to measure nitrite levels. Nitrite release from control cells treated with LPS alone was set as 100%. Results are expressed as the mean  $\pm$  standard deviation derived from three independent experiments. Statistical significance was determined using the Student's *t*-test (\*\* $P < 0.01$ ). Hp, *H. pylori*.



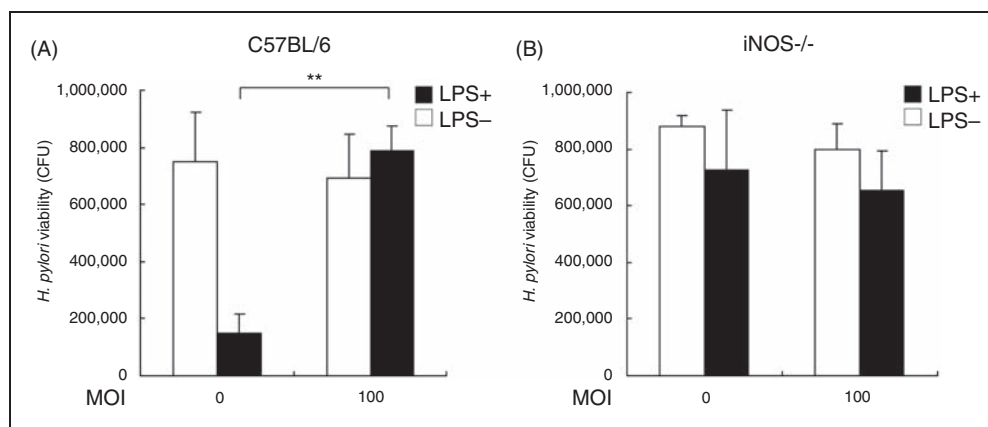
**Figure 3.** Wild-type and isogenic mutants of *H. pylori* produce the same inhibitory effect on LPS-induced NO production by macrophages. RAW 264.7 cells were treated without or with LPS and infected with wild-type (WT) or isogenic mutants of *H. pylori* at a MOI of 100. After incubation for 24 h, culture supernatants were collected to determine levels of nitrite. The data represent the mean  $\pm$  standard deviation derived from three independent experiments. Statistical significance was determined using the Student's *t*-test (\*\* $P < 0.05$ ). Hp, *H. pylori*.

and *H. pylori*, the phosphorylated molecules involved in the MAPK signalling pathway were expressed at a basal level (Figure 6A). *Helicobacter. pylori* infection

led to a decrease in phosphorylated p38 in LPS-stimulated macrophages within 5 to 10 min of culture (Figure 6B). Phosphorylation of ERK1/2 was also inhibited by *H. pylori* infection from 5 to 60 min of culture. In contrast to the results obtained for p38 and ERK1/2, phosphorylation of JNK1/2 in LPS-stimulated PEMs was not affected by *H. pylori* infection. The suppressive effect of *H. pylori* on LPS-induced iNOS expression was augmented by SB203580 and PD98059, specific inhibitors of p38 and ERK1/2, respectively, but not by SP600125, a specific JNK inhibitor (Figure 6D). Taken together, these results suggested that *H. pylori* inhibited LPS-stimulated NO production and iNOS expression in macrophages through the p38 and ERK1/2 signalling pathways.

### Suppression of LPS-induced NF- $\kappa$ B activation in macrophages by *Helicobacter pylori*

We next investigated the effects of *H. pylori* on the levels of two transcription factors, NF- $\kappa$ B and AP-1, which play essential roles in the regulation of iNOS expression.<sup>29</sup> We first examined the effects of *Helicobacter pylori* on NF- $\kappa$ B expression using the luciferase assay in RAW 264.7 cells transfected with the NF- $\kappa$ B-luciferase reporter. *Helicobacter pylori* inhibited LPS-stimulated activation of the NF- $\kappa$ B promoter (Figure 7A), but only slightly inhibited LPS-stimulated AP-1 activity; however, the change was not statistically significant (Figure 7B). We next examined p65 localization and observed that it was primarily



**Figure 4.** Bactericidal effects of NO on *H. pylori*. Murine PEMs from C57BL/6 wild-type (A) or iNOS<sup>-/-</sup> (B) mice were grown on the bottom layer of trans-well culture plates and infected with *H. pylori* at a MOI of 0 or 100 for 48 h. The PEMs were then co-incubated with *H. pylori* in the trans-well insert membrane (0.1  $\mu$ m) for another 6 h and CFU were counted. Bactericidal activity is expressed as the mean  $\pm$  standard deviation derived from three independent experiments. Statistical significance was determined using the Student's t-test (\*\* $P < 0.01$ ).

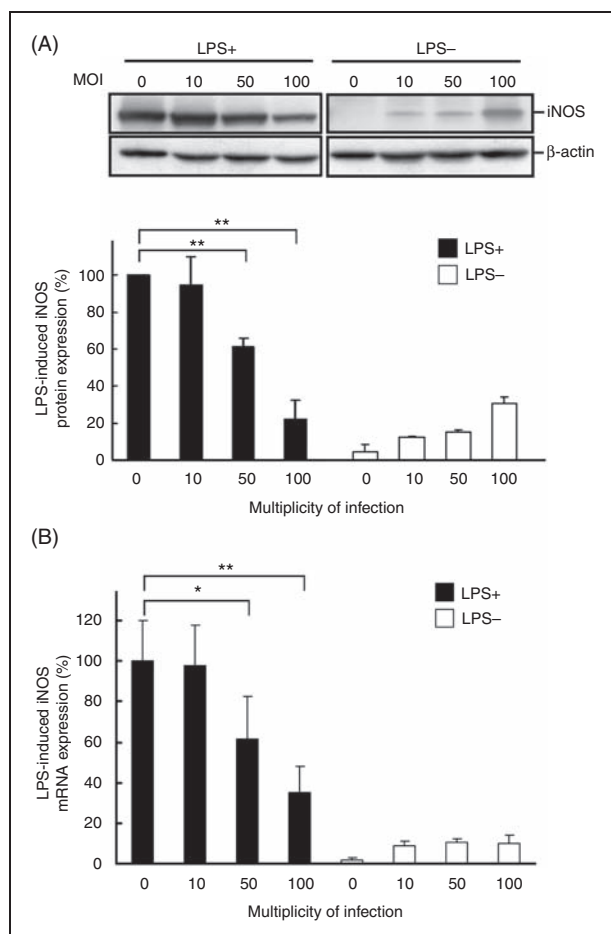
located in the cytosol before LPS treatment. After 1 h of stimulation with LPS, p65 translocated into the nucleus of RAW 264.7 cells. When cells were co-cultured with LPS and *H. pylori*, however, p65 largely remained in the cytosol (Figure 7C). The quantitative data showed that the inhibition of p65 translocation into the nucleus in LPS-treated macrophages upon *H. pylori* infection (Figure 7D). These data suggested that *H. pylori* alone did not alter the distribution of p65 but rather prevented LPS-induced translocation of p65 into the nucleus.

## Discussion

Inducible iNOS and NO are two well known factors that play important roles in the antimicrobial response of macrophages.<sup>30</sup> Our results showed that infection of *H. pylori* elicited a small amount of NO production by macrophages (Figure 1). Live *H. pylori* and also heat-killed *H. pylori* and bacterial lysate slightly stimulated NO synthesis in the absence of bacterial LPS (Figure 2). In addition, without LPS treatment, *H. pylori* stimulated, in a MOI-dependent manner, the expression of iNOS mRNA and protein (Figure 5). Our results confirm the results of a previous study that showed that *H. pylori* can stimulate iNOS expression and activity in murine macrophages.<sup>5</sup> The release of NO appears to be stimulated by *H. pylori*-derived components because even heat-killed *H. pylori* can stimulate NO production.<sup>31</sup> Lipopolysaccharide derived from *H. pylori* produces only low biological activity as a stimulator of NO,<sup>32</sup> with estimates that it is 2000–30,000-fold less potent than LPS derived from *E. coli*.<sup>33</sup> Wilson *et al.* suggested that *H. pylori* induces iNOS expression through both LPS-dependent and -independent mechanisms.<sup>5</sup> The expression of iNOS and the accumulation of NO

have been linked to *H. pylori*-associated gastritis.<sup>6,34</sup> These results demonstrate that NO production plays an important role in gastric inflammatory responses elicited by *H. pylori*.

Apart from infection with *H. pylori*, the stomach is colonized with non-*Helicobacter* microorganisms. The most common non-*Helicobacter* bacteria of the gastric microflora are *Streptococcus*, *Staphylococcus*, *Neisseria*, *Pseudomonas* and *Enterobacteriaceae*.<sup>15</sup> In addition, endoscope-transmitted infections have been reported, including *Salmonella* spp., *E. coli*, *Klebsiella pneumoniae* and *Pseudomonas aeruginosa*.<sup>35</sup> Long-term infection with *H. pylori*<sup>15</sup> or acid-suppressive therapy<sup>16</sup> enables non-*Helicobacter* bacteria to colonize the host stomach. A diverse community of 128 phylotypes was identified using large-scale 16S rRNA sequencing in 23 gastric endoscopic biopsy samples, suggesting that the human stomach may be host to a distinct microbial ecosystem.<sup>18</sup> Both LPS and peptidoglycan (PGN), which is found in the cell wall of Gram-negative and Gram-positive bacteria, have been implicated in the production of NO and pro-inflammatory cytokines by macrophages.<sup>36–39</sup> Patients with both *H. pylori* and non-*Helicobacter* bacteria in their gastric mucosa have higher levels of pro-inflammatory cytokines than patients without bacterial infection.<sup>19</sup> The issue that these findings raise, therefore, is how *H. pylori* can survive in a hostile gastric environment surrounded with profound inflammatory responses provoked by other microorganisms. To mimic the hostile gastric environment with activated macrophages and co-infection of *H. pylori* and non-*Helicobacter* bacteria, purified LPS or PGN were added to the macrophage culture system to determine the effects on NO production. In a preliminary study, we found that treatment with PGN (10  $\mu$ g/ml) for 24 h resulted in an approximately



**Figure 5.** *Helicobacter pylori*-mediated inhibition of iNOS expression in LPS-treated macrophages. Murine PEMs were treated without or with LPS and infected with *H. pylori* at different MOIs. (A) Cell lysates were prepared after 48 h of incubation to measure iNOS protein expression by Western blotting. Protein expression levels were quantified with densitometric analysis and normalized to  $\beta$ -actin (B) iNOS mRNA expression was measured by quantitative real-time PCR after 6 h of incubation, and GAPDH was used as an internal control. The data are presented as the mean  $\pm$  standard deviation of three independent experiments. Statistical significance was determined using the Student's *t*-test (\* $P < 0.05$ ; \*\* $P < 0.01$ ).

four-fold increase in nitrite (8  $\mu$ M) over basal levels in RAW 264.7 cells (Supplementary Figure 1). After 24 h of treatment with LPS (2  $\mu$ g/ml), there was an approximate 8- (22  $\mu$ M) and 150-fold (158  $\mu$ M) increase in nitrite over basal levels in RAW 264.7 cells and PEMs respectively (Figure 1). Murine macrophages were also more sensitive to LPS than PGN. Because of the results of our preliminary studies, we chose LPS for use in the current study.

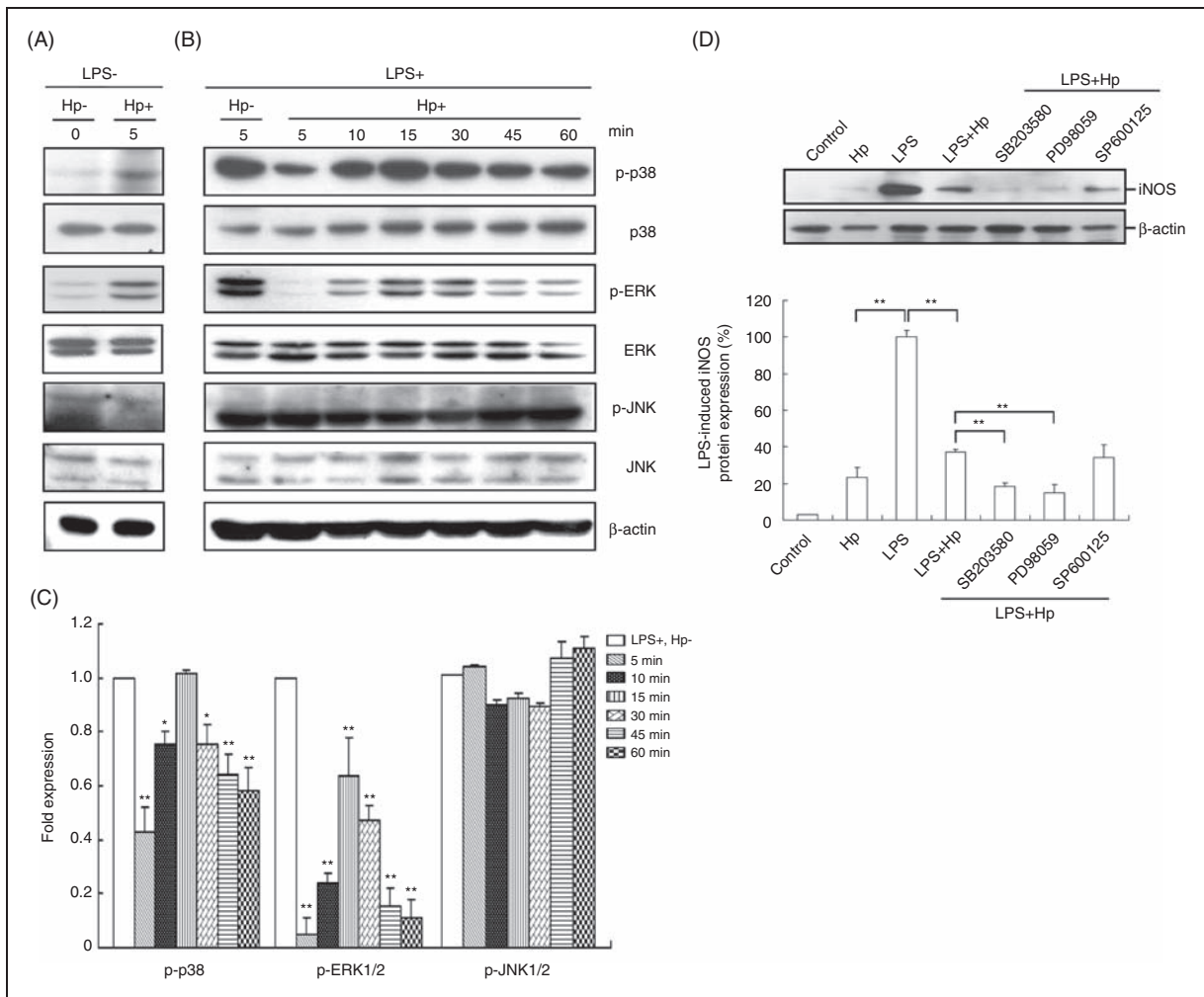
In the present study, we used *in vitro* and *ex vivo* murine model systems to reveal how *H. pylori* evades LPS-dependent killing by macrophages. To demonstrate that the rational design of our murine model

systems mimic a real-life *in vivo* situation, we used wild-type C3H/HeN and TLR4-deficient C3H/HeJ mice to study the role of LPS in killing gastric *H. pylori* *in vivo*. As shown in Supplementary Figure 2, *H. pylori* was more significantly eradicated from the stomachs of LPS-administered C3H/HeN mice than from the stomachs of LPS-administered C3H/HeJ mice. The response of LPS-administered C3H/HeN mice was approximately 11-fold greater than that of LPS-administered C3H/HeJ mice. This result confirmed that LPS enhanced *in vivo* anti-*H. pylori* activity in LPS-responsive mice (C3H/HeN). In addition, the results of this experiment proved our assumption from the *in vitro* and *ex vivo* murine models as the responses were mimicked in the *in vivo* setting.

In this study, we demonstrated that LPS-induced NO production by macrophages was suppressed by *H. pylori* and that this effect was dependent on the presence of live bacteria (Figure 2). Neither heat-killed bacteria nor crude extracts of *H. pylori* suppressed LPS-induced NO production. Likewise, the work of von Bothmer *et al.* demonstrated that *H. pylori* water extract and whole-bacterial suspension produced an L-arginine-sensitive inhibition of NO synthesis.<sup>40</sup> These data raise the issue as to why the bacteria need to be alive. One possibility is that the effect is mediated through the *cag*-pathogenicity island (*cag*-PAI) or VacA. Our results using isogenic mutants of *ΔcagA* or *ΔvacA* indicated that mutant *H. pylori* strains also inhibited LPS-induced NO production in a manner similar to the effects observed with wild-type *H. pylori* (Figure 3). Despite our results suggesting that the suppressive effect is mediated through direct interaction of *H. pylori* with macrophages, the extent to which virulence factors are associated with the inhibition of NO remains unknown. Future studies also are needed on genetic analysis of the virulence factors in different *H. pylori* strains.

Several reports have shown that NO can kill *H. pylori* in cell culture systems.<sup>22,41</sup> Our data indicated that *H. pylori* could survive when *H. pylori* was co-cultured with LPS-activated macrophages in a trans-well system (Figure 4). This effect may be as a result of the suppressive effects of *H. pylori* on LPS-induced NO production by macrophages. This possibility is consistent with the results of previous studies that demonstrated that arginase produced by *H. pylori*<sup>22</sup> and arginase II released from macrophages<sup>42</sup> suppress NO production and lead to immune evasion by the bacteria. Another explanation for the mechanisms through which *H. pylori* inhibits NO production is a decrease in the concentration of L-arginine in culture medium, which interferes with the L-arginine/NO pathway.<sup>40</sup> A reduction in L-arginine availability inhibits iNOS expression and attenuates NO-dependent bactericidal activity.<sup>43</sup> Apart from competition or inhibition



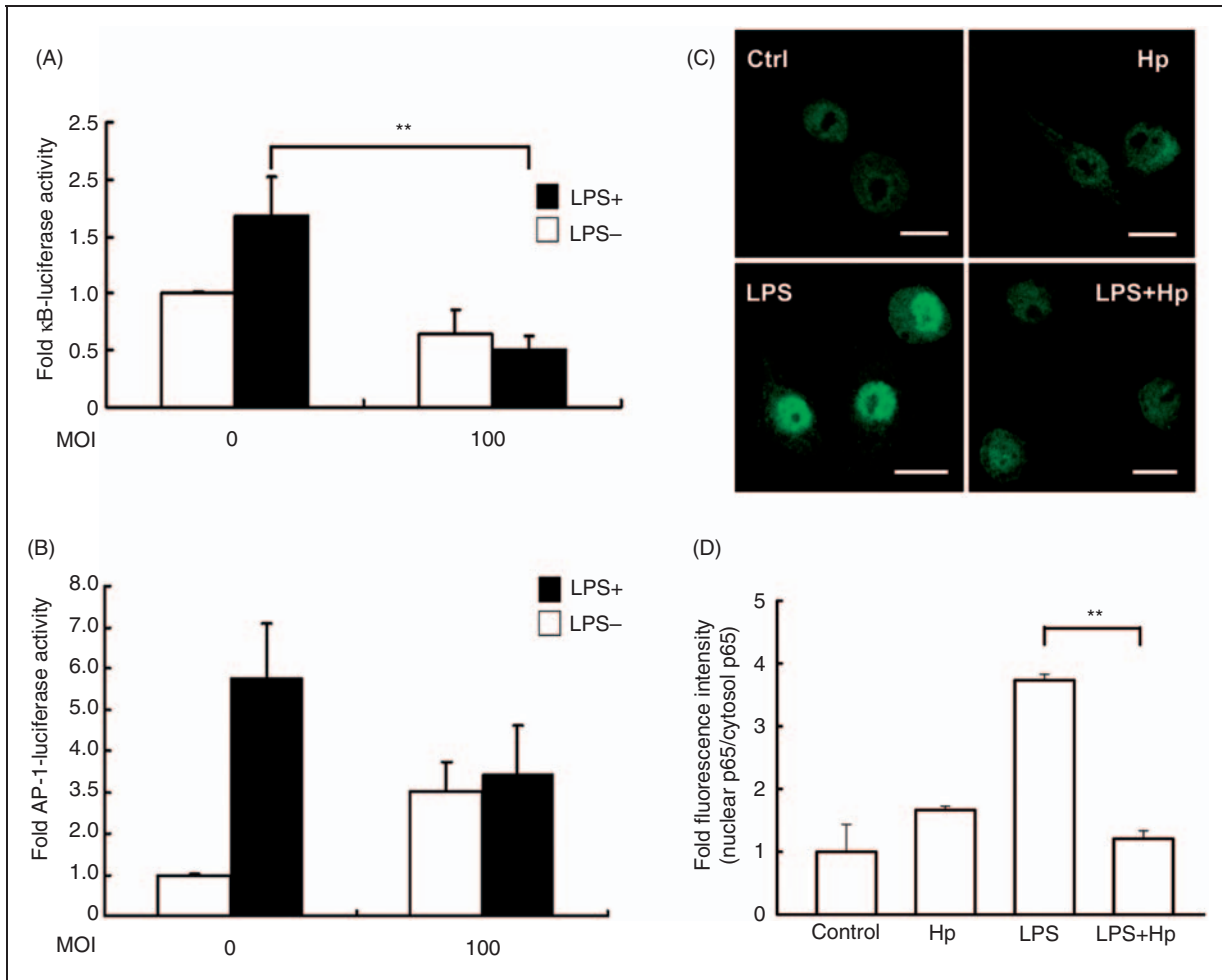


**Figure 6.** Signalling pathways involved in the inhibition of LPS-induced macrophage activation by *H. pylori*. Murine PEMs were incubated without LPS (A) or with LPS (B) and infected (or not infected) with *H. pylori* at a MOI of 100 for the indicated times. Expression levels of phosphorylated p38 (p-p38), anti-phosphorylated extracellular signal-regulated kinases (p-ERK), and anti-phosphorylated c-Jun-N-terminal kinase (p-JNK) were determined by Western blotting. Representative Western blot results from one of three independent experiments are shown. (C) Protein expression levels were quantified with densitometric analysis, normalized to  $\beta$ -actin, and presented as the mean  $\pm$  standard deviation derived from three independent experiments. Statistical significance was determined using the Student's *t* test [ $*P < 0.05$ ;  $**P < 0.01$  as compared with LPS-stimulated cells in the absence of *H. pylori* infection (LPS+, Hp-), lower panel]. (D) Murine PEMs were pretreated for 30 min with SB203580 (p38 inhibitor), PD98059 (ERK inhibitor), or SP600125 (JNK inhibitor), followed by incubation with LPS and infection with *H. pylori* at a MOI of 100 for 48 h. Representative Western blot results from one of three independent experiments are shown. The data are expressed as the mean  $\pm$  standard deviation determined from three independent experiments. Statistical significance was determined using the Student's *t*-test ( $**P < 0.01$ ). Hp, *H. pylori*.

of NO production by macrophages, previous studies showed that *H. pylori* induces apoptosis in both macrophages<sup>44</sup> and T lymphocytes.<sup>45</sup> In the present study, we did not add L-arginine to cultures; we used a high dose of LPS to induce iNOS expression, and the effect of LPS was independent of the concentration of L-arginine, as previously demonstrated.<sup>40</sup> Cell viability was not influenced when cells were incubated with *H. pylori* at a high MOI of 100 and in the presence of LPS (Figure 1). *H. pylori* appears to have intricate mechanisms to regulate the activity of macrophages

and to maintain bacterial survival under various infectious conditions.

Lipopolysaccharide stimulates iNOS gene expression and NO production. Stimulation is positively regulated by NF- $\kappa$ B, which is normally bound to its inhibitor, I $\kappa$ B, in the cytoplasm. Phosphorylation of I $\kappa$ B by I $\kappa$ B kinase results in the degradation of I $\kappa$ B, which dissociates NF- $\kappa$ B and leads to the nuclear translocation of NF- $\kappa$ B and the up-regulation of downstream gene expression. In addition to NF- $\kappa$ B, LPS can activate MAPK pathways in macrophages, including p38,



**Figure 7.** *Helicobacter pylori*-mediated attenuation of LPS-induced NF- $\kappa$ B activation. RAW 264.7 cells transfected with a reporter gene for NF- $\kappa$ B (A), AP-1 (B), or  $\beta$ -Gal-lacZ (1  $\mu$ g each) were treated without or with LPS and infected (or not infected) with *H. pylori* at a MOI of 100. Luciferase activity was normalized to the expression level of  $\beta$ -Gal-lacZ. The data are expressed as the mean  $\pm$  standard deviation derived from three independent experiments. Statistical significance was determined using the Student's *t*-test (\*\* $P$ <0.05). (C) The nuclear translocation of p65 after 1 h treatment with or without LPS and infection of *H. pylori* at a MOI of 100. Cells were washed and treated with anti-p65 followed by fluorescein isothiocyanate-conjugated anti-mouse IgG (green). Cells were co-stained with propidium iodide to visualize the nucleus (red). Cells were analyzed by confocal fluorescence microscopy. Regions of p65 nuclear localization appear in yellow in the overlay. Representative images from one of three independent experiments are shown. Scale bar, 10  $\mu$ m. Ctrl, control; Hp, *H. pylori*.

ERK-1/2, and JNK-1/2. A common strategy for pathogens to overcome host defense is interference with the activity of NF- $\kappa$ B or MAPKs.<sup>10,12,13,46</sup> In the present study, *H. pylori* may have suppressed LPS-induced NO production by macrophages by inhibiting LPS-stimulated NF- $\kappa$ B activation. Our data demonstrated that the phosphorylation of p38 and ERK1/2 was attenuated by *H. pylori* infection at an early stage of infection, whereas JNK-1/2 was not affected. These results suggest that *H. pylori* targets p38 and ERK 1/2, but not JNK1/2. Our data are consistent with previous findings that pathogens can exploit NF- $\kappa$ B to manipulate cellular responses.<sup>10,13,14,47</sup> Together, these findings support our hypothesis that *H. pylori* modulates host signalling to evade the host immune system.

In addition, our results provide insight into the molecular mechanisms through which indigenous *H. pylori* survive commensurately in the stomach with non-*Helicobacter* microorganisms that induce potent immune responses.

In conclusion, we demonstrate that infection with live *H. pylori* attenuates LPS-induced iNOS gene transcription and NO production in a mouse macrophage model. We further find that *H. pylori* inhibits LPS-induced MAPK signalling and NF- $\kappa$ B activation in macrophages. Collectively, this study reveals a new mechanism through which *H. pylori* modulates host cell signalling to protect itself from inflammatory responses and to survive in the harsh environment of the stomach.

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## Conflicts of interest

None declared.

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