Mechanical ventilation modulates Toll-like receptors 2, 4, and 9 on alveolar macrophages in a ventilator-induced lung injury model

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Objective: To investigate the role of Toll-like receptor 2 (TLR2), TLR4, TLR9 and myeloid differentiation factor 88 (MyD88) on alveolar macrophages in ventilator-induced lung injury (VILI).

Methods: Male, adult pathogen-free Sprague-Dawley rats weighing 300-350 g were used in this study. Animals were tracheotomized and allowed to breathe spontaneously for 4 h or mechanically ventilated for 4 h with low or high tidal volume (7 or 40 mL/kg). TLR2, TLR4, and TLR9, MyD-88 and NF-κB of alveolar macrophages’ expression under the different ventilation conditions were detected. Pulmonary permeability, lung inflammatory, IL-6 and IL-1β were assessed as well.

Results: Rats subjected to high tidal volume showed significantly greater pulmonary permeability and lung inflammatory than the control rats. Alveolar macrophages from rats subjected to high tidal volume also showed significantly higher protein expression of TLR2 (0.59±0.049 vs 0.35±0.036 and 0.36±0.031, both P<0.001), TLR4 (0.845±0.0395 vs 0.401±0.026 and 0.403±0.020, both P<0.001 ), TLR9 (0.727±0.074 vs. 0.383±0.039 and 0.367±0.043, both P<0.001), MyD-88 (1.01±0.060 vs. 0.485±0.045 and 0.507±0.046, both P<0.001) and NF-κB (0.776±0.067 vs. 0.448±0.043 and 0.481±0.047, both P<0.001), as well as significantly higher concentrations of IL-6 (7.32±0.24 vs 2.42±0.13 and 2.44±0.32, both P<0.001) and IL-1β (139.95±9.37 vs 53.63±5.26 and 53.55±6.63, both P<0.001) than the control and low tidal volume group.

Conclusions: The overexpression of TLR2, TLR4, and TLR9 on alveolar macrophages and release of pro-inflammatory cytokines play a role in VILI.

Keywords: Toll like receptor (TLR); myeloid differentiation factor 88 (MyD88); alveolar macrophages; ventilator-induced lung injury (VILI)

Submitted Oct 13, 2014. Accepted for publication Feb 09, 2015.
doi: 10.3978/j.issn.2072-1439.2015.02.10
View this article at: http://dx.doi.org/10.3978/j.issn.2072-1439.2015.02.10

Background

Mechanical ventilation facilitates surgical interventions during general anesthesia and is a life-saving therapy in patients with acute lung injury or with the most severe form of this injury, acute respiratory distress syndrome. At the same time, mechanical ventilation can exacerbate lung injury; the resulting damage has been termed ventilator-induced lung injury (VILI) (1). This type of acute lung injury is characterized by inflammation associated with robust release of proinflammatory mediators and activation of inflammatory signaling pathways (2).

Various inflammatory cytokines are released into the pulmonary distal airspaces during acute lung injury; among these, IL-1β initiates and amplifies inflammation (2,3). This cytokine has also recently been implicated in the pathogenesis of VILI, in which it is a major contributor to alveolar barrier dysfunction (4). In fact, clinical studies have identified IL-1β as one of the best markers of ventilator-induced lung inflammation (5).

Alveolar macrophages account for 5% of peripheral lung cells and reside in the alveolar space, where they account for 90% of all leukocytes, with the remainder being mainly dendritic cells and T cells (6). Alveolar macrophages...
play a crucial role in the maintenance of immunological homeostasis and host defense, and they are the primary producers of proinflammatory cytokines in lungs following exposure to noxious stimuli (2,7,8). These macrophages are rapidly activated by mechanical ventilation, suggesting that they may play a critical role in the pathogenesis of VILI. Indeed, removing alveolar macrophages from rats exposed to mechanical ventilation at high tidal volume attenuates the resulting alveolar barrier dysfunction and inflammatory lung injury (4,8,9).

Toll-like receptors (TLRs) recognize pathogen-associated molecular patterns (PAMPs), which refer to biomolecules produced by invading microbes or released from damaged tissue. These biomolecules, which include lipopolysaccharides, peptidoglycans, lipoteichoic acid and CpG-containing oligonucleotides (CpG ODNs), serve as danger signals that initiate an inflammatory immune response. TLRs trigger inflammation mediated by complement, macrophages, and neutrophils. These cells produce chemokines or cytokines that mediate systemic immune responses and recruit leukocytes to the sites of inflammation. This injury-induced inflammation can lead to systemic inflammatory response syndrome (SIRS) and multiple organ dysfunction syndrome (MODS) (10-12).

TLR2, TLR4 and TLR9 are crucial for PAMP signaling. These receptors work through the adaptor protein myeloid differentiation factor 88 (MyD88), which activates nuclear factor-κB (NF-κB) and ultimately stimulates the production of proinflammatory cytokines (13,14). Rat alveolar macrophages are known to be sensitive to mechanical and CpG ODN (15) stimulation, but we are unaware of studies examining whether they are sensitive to ventilation-induced injury. If so, this would implicate TLR- and MyD88-mediated signaling in alveolar macrophages in the pathogenesis of VILI.

To address this question, we examined the effects of mechanical ventilation on the expression of TLR2, TLR4 and TLR9 genes in rat alveolar macrophages, as well as on activation of MyD88 and NF-κB.

Materials and methods

Animal procedures

Healthy, pathogen-free male Sprague-Dawley rats (300-350 g) were obtained from the Animal Center of Guangxi Medical University (Nanning, China), and kept in an environmentally controlled room (23±2 °C, 55%±10% humidity) with a 12-h light/dark cycle and allowed free access to food and water. Rats were anesthetized initially by intraperitoneal injection of pentobarbital (50 mg narcoren/kg body weight; Merial, Halbergmoos, Germany) and fentanyl (0.05 mg/kg body weight; Janssen-Cilag, Neuss, Germany), and then placed in a supine position on an adjustable warming pad to maintain a temperature of 37±1 °C. Anesthesia was maintained by supplementing with one third of the initial dose of anesthetic agents approximately every 45 min during the experimental period. Temperature was monitored continuously using a rectal probe. Anesthetized animals were randomly allocated into three groups (with 10 rats for each) using table of random number: non-ventilated, ventilated with low tidal volume (7 mL/kg, low V₁), and ventilated with high tidal volume (40 mL/kg, high V₁). The study was performed in accordance with the guiding principles for the care and use of laboratory animals approved by Institutional Animal Care and Use Committee of Tumor Hospital of Guangxi Medical University.

All animals were tracheotomized and then allowed to breathe spontaneously for 4 h or ventilated mechanically for 4 h using a small animal ventilator (TOPO, Kent Scientific, Torrington, USA) set at a positive end-expiratory pressure of 0. The ventilation rate was 80 breaths/min at low tidal volume and 60 breaths/min at high tidal volume. All rats were supplemented with oxygen at approximately 40-50% (16) of the inspired oxygen fraction, as determined by oxygen monitoring; the amount of oxygen was always less than 55% of the inspired oxygen fraction. Fluid-filled polyurethane catheters (inside diameter, 50.58 mm; outside diameter, 50.96 mm; SIMS Portex Ltd., Hythe, UK) were inserted into the right femoral vein for infusion of saline (0.01 mL/g/h body weight) and glycopyrrolate (0.1 μg/g body weight) to maintain intravascular volume, and into the right femoral artery for measurement of blood pressure and blood gas analysis.

After 4 h, all animals were sacrificed by lethal dose of anesthetic agent, and tissue samples were harvested. All animal procedures were performed with special care to minimize activation of an inflammatory response.

Lung histopathology and determination of wet/dry ratio

The middle lobe of the right lung was fixed by injection of 10% formaldehyde solution through the right middle bronchus at a pressure of 20 cmH₂O. After fixation, the tissue was embedded in paraffin and 4-μm sections were...
prepared and stained with hematoxylin and eosin, then examined by light microscopy. To score lung injury we used a modified VILI histopathology scoring system as previously described (17). VILI was scored according to the following four main items: alveolar congestion; haemorrhage; infiltration or aggregation of neutrophils in airspace or vessel wall; and thickness of the alveolar wall/hyaline membrane formation. A score of 0 represented normal lungs; 1 represented mild, less than 25% lung involvement; 2 represented moderate, 25% to 50% lung involvement; 3 represented severe, 50% to 75% lung involvement; and 4 represented very severe, more than 75% lung involvement. An overall score of VILI was obtained based on the summation of all the scores from normal or ventilated lungs (n=10 per group).

Pulmonary wet/dry ratios were measured as an index of pulmonary congestion. Immediately after rats were killed, the right lower lobe was weighed and then dried to a constant weight at 60 °C for 24 h.

Collection and analysis of bronchoalveolar lavage fluid (BALF) and alveolar macrophages

Alveolar macrophages were isolated as described previously (18). In brief, the trachea of killed animals was cannulated and the lungs were flushed once with 5 mL of cold phosphate-buffered saline (Dulbecco’s PBS; Gibco BRL, Grand Island, USA) to collect BALF for mediator analysis (BALF 1 fraction). The lungs were subsequently flushed another eight times with 10 mL PBS to obtain alveolar macrophages (BALF 2 fraction). The two BALF fractions were centrifuged, and the BALF 1 supernatant was stored at −80 °C for enzyme-linked immunosorbent assay (ELISA) and determination of total protein using the bicinchoninic acid (BCA) assay according to the manufacturer’s instructions (Pierce, Rockford, USA).

BALF 1 and 2 pellets were combined, resuspended in Dulbecco’s modified Eagle medium (DMEM, Gibco, USA), counted, and transferred to 24-well culture plates (BD, Franklin Lakes, NJ, USA). After incubation for 60 min at 37 °C in a 5% CO₂ atmosphere, cultures were washed with DMEM to remove nonadherent cells. The adherent cells were counted using a hemocytometer, viability was determined using a 0.2% trypan blue exclusion assay, and cell differentiation and aggregation were examined by counting 500 cells on a Wright-Giemsa-stained slide. These cultures also served as the source of RNA and protein for analyzing mRNA and protein expression in alveolar macrophages.

Cytokine analysis in BALF and plasma

Concentrations of IL-1β, IL-6 and IL-10 were determined in BALF and plasma using a commercial ELISA according to the manufacturer’s recommendations (R&D Systems, Minneapolis, USA). The manufacturer-specified detection limits with this kit were IL-1β, 15.6 pg/mL; IL-6, 0.078 pg/mL; and IL-10, 0.78 pg/mL.

RNA isolation and real-time reverse transcriptase-polymerase chain reaction

Total RNA was extracted from alveolar macrophages using TRIzol (Invitrogen, Carlsberg, USA). RNA quality and quantity were determined by spectrophotometry. RNA samples were reverse-transcribed into cDNA using a reverse transcription kit (Toyobo, Osaka, Japan). The following primers were used to determine mRNA expression levels: TLR2, 5’-GGGATACAGGCGGTCAAGAC-3’ (forward) and 5’-CGAGACAGATGAAATGGTTGT-3’ (reverse); TLR4, 5’-CGCTCTGAGATTCCGATC-3’ (forward) and 5’-CTCCTCAGTTGAAGTTTTC-3’ (reverse); TLR9, 5’-CGCCACACAACTGACATTCA-3’ (forward) and 5’-TAAAGGTCCTCCTCTGCTCCA-3’ (reverse); MyD88, 5’-GAGATCCGCGATATGACG-3’ (forward) and 5’-TGGTTTGATATGACG-3’ (reverse); NF-κB, 5’-GAGGACTTCTGAGTTTG-3’ (forward) and 5’-GAGGACTTCTGAGTTTG-3’ (reverse); and glyceraldehyde phosphate dehydrogenase (GAPDH), 5’-GGCACAGTCAAGGCTGAGAATG-3’ (forward) and 5’-ATGGTGGTGAAGAGAGG-3’ (reverse). Real-time reverse transcriptase PCR was performed as previously described (19). The level of each target gene was normalized relative to that of GAPDH in each sample using the ΔCt method. Relative differences in gene expression among groups of the alveolar macrophages were determined using the comparative Ct (ΔΔCt) method and fold expression was calculated by the formula 2−ΔΔCt, where ΔΔCt represents ΔCt values normalized relative to the mean ΔCt of healthy control samples. All fragments were checked for the specificity by direct sequencing of both strands with an ABI PRISM 310 Genetic Analyzer using Big Dye Terminator kit v 3.1 (Applied Biosystems, CA, USA).

Protein extraction and Western blotting

Proteins were extracted from alveolar macrophages using RIPA buffer and quantitated using the BCA protein assay.
kit. For Western blot detection of TLR2, TLR4, TLR9, MyD-88 and NF-κB, samples (50 μg) were electrophoresed on a 10% SDS-polyacrylamide gel, electroblotted onto nitrocellulose membranes, and blocked at 4 °C overnight with 5% bovine serum albumin (BSA) dissolved in Tris-buffered saline (TBS) containing 0.1% Tween-20. Membranes were incubated for 2 h with rabbit anti-rat monoclonal antibodies (Cell Signaling Technology, Danvers, USA) against TLR2 (1:1,000), TLR4 (1:2,000), TLR9 (1:1,000), MyD-88 (1:1,000) and NF-κB (1:1,000). Blots were washed several times, and then incubated with horseradish peroxidase-conjugated mouse anti-rabbit antibody (1:5,000 in PBS containing 5% nonfat dry milk; Pierce, USA). Blots were again washed several times and visualized by enhanced chemiluminescence (Pierce, USA). Blots were also incubated with rabbit anti-rat GAPDH antibody (1:10,000, Santa Cruz Biotechnology, CA, USA) as a control for protein loading.

Sample size and statistical analysis
We estimated that a total of 30 rats would be needed to detect a difference between groups, with a 2-tailed α of 0.05 and a 1-β of 0.80, for a comparison of two independent proportions if there was an absolute decrease of 15% in the composite outcome measure. The Data and Safety Monitoring Board, blinded to treatment group, reviewed two formal interim analyses and regular reports of our primary composite outcome as well as serious adverse events. All data were analyzed using SPSS software (version 16.0; IBM, Chicago, USA). All quantitative data were expressed as mean ± SD. Multiple comparisons were analyzed using one-way ANOVA, followed by the Friedman test and the Wilcoxon signed-rank test. P values less than 0.05 were considered significant.

Results
Outcomes and overall pathophysiology assessment
All animals survived the 4-h period of spontaneous breathing or mechanical ventilation at low or high tidal volume. Lungs from animals ventilated with high tidal volume showed acute inflammatory infiltration and perivascular edema, whereas no major histological differences were observed between animals ventilated with low tidal volume or spontaneously breathing control animals. The lung histopathology score was higher in high Vₜ rats as compared with the controls and low Vₜ. However, no differences were noted between the low Vₜ and the controls (Figure 1 and Table 1).

Counts of isolated AMs
In the high Vₜ group, significantly more AMs were isolated than in the low Vₜ group and non-ventilated animals (Figure 2). And the macrophage percentages were more than 90% counted by the hemocytometer.

Wet/dry ratios and BALF analysis
Wet/dry ratios, BALF total protein levels were significantly higher in the group of high Vₜ than in the control and low Vₜ (Table 1). These three parameters were similar for rats ventilated with low tidal volume and for spontaneously breathing rats.
Pro-inflammatory cytokine concentrations in BALF and plasma

Cytokine profiles were similar in animals ventilated with low tidal volume and control animals (Table 1). IL-10 were undetectable (IL-10 detection threshold was 0.78 pg/mL) in either BALF or plasma in the three groups, and plasma IL-1β levels were below the level of detection (IL-1β detection threshold was 15.6 pg/mL) in control and ventilation groups. These animals showed significantly higher levels of IL-6 levels in BALF and plasma in the high VT group than did the other two groups of animals (Table 1).

Mechanical ventilation induced MyD-88 and NF-κB expression and up-regulated expression of TLR2, TLR4 and TLR9

Expression of mRNA and protein for TLR2, TLR4, TLR9, MyD-88 and NF-κB were similar for animals ventilated with low tidal volume and for spontaneously breathing controls (Figures 3,4). The mRNA and protein levels of all these genes, however, were significantly higher for animals ventilated with high tidal volume than the control and low tidal volume group. mRNA levels as follow, TLR2 (4.78±0.98 vs. 1±0.22 and 1.15±0.30, both P<0.001), TLR4 (5.15±0.78 vs. 1±0.19 and 1.23±0.35, both P<0.001), TLR9(10.75±1.80 vs. 1±0.27 and 0.91±0.28, both P<0.001), MyD-88 (3.97±0.75 vs. 1±0.20 and 0.96±0.24, both P<0.001), NF-κB (5.79±0.98 vs. 1±0.23 and 1.08±0.25, both P<0.001). Protein levels as follow, TLR2 (0.59±0.049 vs. 0.35±0.036 and 0.36±0.031, both P<0.001), TLR4 (0.845±0.0395 vs. 0.401±0.026 and 0.403±0.020, both P<0.001), TLR9 (0.727±0.074 vs. 0.383±0.039 and 0.367±0.043, both P<0.001), MyD-88 (1.01±0.060 vs. 0.485±0.045 and 0.507±0.046, both P<0.001) and NF-κB (0.776±0.067 vs. 0.448±0.043 and 0.481±0.047, both P<0.001).

Discussion

Here we investigated the hypothesis that mechanical ventilation activates alveolar macrophage TLR receptors that are normally involved in initiating innate immune responses to PAMPs. TLR activation should up-regulate expression of the adaptor protein MyD-88, stimulating production of NF-κB and ultimately the secretion of proinflammatory cytokines. As predicted, we found that 4-h mechanical ventilation of tracheotomized rats using a high tidal volume increased the expression of the major PAMP sentinels TLR2, TLR4 and TLR9, expression of

Table 1 Levels of total protein, IL-6 and IL-1β in BALF after ventilation-induced lung injury and lung histopathology score

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Non-ventilated control</th>
<th>Low tidal volume</th>
<th>High tidal volume</th>
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<tbody>
<tr>
<td>Wet/dry ratio</td>
<td>4.88±0.18</td>
<td>4.93±0.26</td>
<td>5.74±0.21*</td>
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<tr>
<td>Protein content (μg/μL)</td>
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<td>593±57</td>
<td>988±39*</td>
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<td>BALF (pg/mL)</td>
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<td>IL-6</td>
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<td>2.44±0.32</td>
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<td>IL-1β</td>
<td>53.63±5.26</td>
<td>53.55±6.63</td>
<td>139.95±9.37*</td>
</tr>
<tr>
<td>Plasma</td>
<td></td>
<td></td>
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<tr>
<td>IL-6 (pg/mL)</td>
<td>1.70±0.23</td>
<td>1.67±0.21</td>
<td>2.08±0.11*</td>
</tr>
<tr>
<td>VILI-score</td>
<td>0.30±0.10</td>
<td>0.50±0.15</td>
<td>3.5±0.5*</td>
</tr>
</tbody>
</table>

Data are presented as mean ± SD (n=10 in all groups). *, P<0.001 vs. non-ventilated animals and animals ventilated with low tidal volume. BALF, bronchoalveolar lavage fluid; IL-1β, interleukin-1β; IL-6, interleukin-6; VILI, ventilator-induced lung injury.
MyD-88 and NF-κB, and concentrations of IL-1β and IL-6 in the BALF. Our findings implicate TLR-mediated signaling in ventilation-induced lung injury and suggest that the pathogenesis occurs through a mechanism similar to that induced by infectious stimuli.

Our rat model of mechanical ventilation with high tidal volume reproduced the pathologic changes consistent with VILI, including increased alveolar permeability (Figure 1) and increased alveolar macrophage cell counts (Figure 2) and protein levels in BALF (Table 1) in the absence of infection. In this way, mechanical ventilation appears to stimulate the immune system akin to severe bacterial infections (12). These findings are consistent with various experimental models of VILI, in which the authors have proposed that alveolar overdistension triggers an innate immune inflammatory reaction similar to that caused by pathogens (20-22).

**Figure 3** Fold changes in TLR2, TLR4, TLR9, MyD88 and NF-κB mRNA levels in alveolar macrophages of healthy rats after 4 h of spontaneous breathing (non-ventilated) or 4 h of mechanical ventilation with tidal volumes of 7 mL/kg (low VT) or 40 mL/kg (high VT) are shown in (B). Data are mean ± SD, n=10. Results were normalized to the values in GAPDH control amplifications, and expressed as fold changes relative to the non-ventilated control animals. *, P<0.001 vs. non-ventilated animals. TLR, Toll-like receptor; MyD88, myeloid differentiation factor 88; NF-κB, nuclear factor-κB.

**Figure 4** Changes in TLR2, TLR4, TLR9, MyD88 and NF-κB protein levels on alveolar macrophages of healthy rats after 4 h of spontaneous breathing (non-ventilated) or 4 h of mechanical ventilation with tidal volumes of 7 mL/kg (low VT) or 40 mL/kg (high VT). Representative Western blot results are shown above, and quantitation of TLR2, TLR4, TLR9, MyD88 and NF-κB protein levels independent experiments is shown in (Figure 3A). Data are mean ± SD, n=10. Results were normalized to levels of GAPDH protein detected on the same blots. *, P<0.001 vs. non-ventilated animals; †, P<0.001 vs. animals ventilated with low tidal volume; ‡, P>0.05 vs. non-ventilated animals. TLR, Toll-like receptor; MyD88, myeloid differentiation factor 88; NF-κB, nuclear factor-κB.
reaction involves changes in the growth, differentiation, migration, remodeling, and gene expression of various pulmonary cell types, including alveolar macrophages, alveolar epithelial cells, endothelial cells and fibroblasts. One consequence of these changes is an increase in pulmonary and systemic pro-inflammatory cytokine levels, as we observed in our measurements of cytokine levels in BALF and serum after ventilation with high tidal volume.

Our results are consistent with previous studies suggesting that mechanical ventilation, alone or in combination with infection, induces inflammation by the synthesis of NF-κB and proinflammatory cytokines (12,23-25). For example, a study with isolated, perfused mouse lung showed that combining 150-min lung overinflation with a tidal volume of 32 mL/kg and lipopolysaccharide treatment activated NF-κB and induced the release of cytokines known to be dependent on NF-κB (24). Indeed we focused on alveolar macrophages in our study because they are involved in initiating innate immune responses and may activate NF-κB in acute lung injury and inflammation (26). Removing alveolar macrophages from rats exposed to mechanical ventilation at high tidal volume attenuates the resulting alveolar barrier dysfunction and inflammatory lung injury (4,8,9).

Our animals ventilated with low tidal volume were indistinguishable from control animals breathing spontaneously. This is consistent with the widespread use of ventilation with low tidal volume to protect against VILI (16,27,28). Our findings contrasts with studies in animals suggesting that even ventilation performed with low tidal volume can induce lung inflammation and increase the risk of ventilator-induced injury. For example, computed tomography studies suggest that low tidal volume can still cause tidal hyperinflation in patients with acute respiratory distress syndrome because of the anatomical heterogeneity of the damaged lung (16). These differences are likely due to differences between laboratory animals and humans, and should be explored further to optimize tidal volume for vulnerable patients.

Our study indicates that mechanical ventilation with high tidal volume upregulates the expression of TLR2, TLR4 and TLR9. TLRs have long been recognized to play a predominant role in the innate immune response to infection and tissue injury (16,29,30). TLR2 is widely expressed in first-line host defense cells, including monocytes, macrophages, dendritic cells, and polymorphonuclear leukocytes. TLR2 is thought to mediate inflammatory responses to components of Gram-positive and -negative bacteria and mycobacteria such as peptidoglycan, lipoteichoic acid, bacterial lipoproteins, lipopeptides, and lipoarabinomannan (31). TLR4 activates MyD88-dependent and Toll/IL-1 receptor domain containing adapter inducing interferon (TRIF)-dependent signaling pathways. It is activated in response to mechanical ventilation with low tidal volume (32), and it plays an important role in acute lung injury induced by ventilation with high tidal volume (33), lipopolysaccharide (34), acid aspiration (35), hemorrhage (36), and ischemia and reperfusion injury (37). TLR9 mediates cellular responses to CpG ODN and to endogenous ligands released during noninfectious tissue injury, including high-mobility group box 1 proteins released from necrotic cells, oxidized phospholipids created by locally generated reactive oxygen species, low molecular weight hyalurans and fibrinogens released from degraded extracellular matrix, heat shock proteins released from necrotic cells, and surfactant protein-A (16,38,39). In this way, our study extends the proinflammatory role of TLRs to the specific case of VILI.

Upon binding to endogenous activators, TLR2, TLR4 and TLR9 recruit the downstream adaptor molecule MyD88, which activates NF-κB and thereby induces transcription of pro-inflammatory genes, ultimately leading to cytokine production (40). Consistent with this sequence of events, we found that mechanical ventilation with high tidal volume upregulated TLR expression, NF-κB and MyD88 activation in the lung, and levels of IL-1β and IL-6 in BALF and serum. Our findings suggest that signaling involving TLR2, TLR4, TLR9 and MyD88 may contributes to the pro-inflammatory response during ventilation induced lung injury.

One limitation of our approach is that stresses due to the experimental manipulations may have triggered the same TLR-mediated signaling that we wished to study. To minimize this risk, we continuously monitored oxygen saturation and hemodynamic stability in the anesthetized rats. We also administered oxygen to the rats under anesthesia and maintained the inspired oxygen fraction below 55% to avoid both hypoxia and hyperoxia (16,41,42). The fact that all outcomes that we measured were similar between the group ventilated with low tidal volume and the control group suggests that our precautions were effective.

Although our data may imply roles for the TLR2, TLR4 and TLR9/MyD88 signaling in regulating multiple pro-inflammatory cytokines during MV, we acknowledge some limitations to this study. First, we did not explore whether repression of MyD88 expression during high Vₜ MV could be reversed by returning to low Vₜ. However in the patients with ALI, pulmonary and systemic inflammatory responses induced by temporary application of high Vₜ...
mechanical ventilation can be reversed by reinstitution of lung protective MV (12), at least over the time frame of a few hours. Second, we do not know whether inhibition of TLR2, TLR4 and TLR9 with blocking antibodies affect the MyD88 response. We can not say that our data fully demonstrate that TLR2, TLR4 and TLR9 pathway is conclusively involved in increased inflammation associated with the use of high-V̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̇̆