Pulmonary Inflammation Triggered by Ricin Toxin Requires Macrophages and IL-1 Signaling

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Pulmonary Inflammation Triggered by Ricin Toxin Requires Macrophages and IL-1 Signaling

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Ricin is a potent ribotoxin considered to be a potentially dangerous bioterrorist agent due to its wide availability and the possibility of aerosol delivery to human populations. Studies in rodents and nonhuman primates have demonstrated that ricin delivered to the pulmonary system leads to acute lung injury and symptoms resembling acute respiratory distress syndrome. Increasing evidence suggests that the inflammatory effects triggered by ricin are responsible for its lethality. We demonstrated previously that ricin administered to the lungs of mice causes death of pulmonary macrophages and the release of proinflammatory cytokines, suggesting macrophages may be a primary target of ricin. Here we examined the requirement for macrophages in the development of ricin-mediated pulmonary inflammation by employing transgenic (MAFIA) mice that express an inducible gene driven by the e-fms promoter for Fas-mediated apoptosis of macrophages upon injection of a synthetic dimerizer, AP20187. Administration of aerosolized ricin to macrophage-depleted mice led to reduced inflammatory responses, including recruitment of neutrophils, expression of proinflammatory transcripts, and microvascular permeability. When compared with control mice treated with ricin, macrophage-depleted mice treated with ricin displayed a reduction in pulmonary IL-1β. Employing mice deficient in IL-1, we found that ricin-induced inflammatory responses were suppressed, including neutrophilia. Neutrophilia could be restored by co-administering ricin and exogenous IL-1β to IL-1α/β−/− mice. Furthermore, IL-1Ra/anakinra cotreatment inhibited ricin-mediated inflammatory responses, including recruitment of neutrophils, expression of proinflammatory genes, and histopathology. These data suggest a central role for macrophages and IL-1 signaling in the inflammatory process triggered by ricin. The Journal of Immunology, 2009, 183: 1419–1426.

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3Abbreviations used in this paper: ARDS, acute respiratory distress syndrome; BAL, bronchoalveolar lavage; EBD, Evans blue dye; qRT-PCR, quantitative real-time RT-PCR; SAPK, stress-activated protein kinase.

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Ricin is a 62-kDa protein consisting of two glycoprotein chains linked by a disulfide bridge. The 34-kDa B chain, a lectin that binds primarily to galactose-containing surface proteins, facilitates the internalization of the 32-kDa A chain, the toxic moiety. Following endocytosis and retrograde transfer through the Golgi apparatus, the A subunit of ricin enters the cytosol where it depurinates a single adenine (A4256 in mouse) in the 28S rRNA ribosomes. The depurination of the A4256 is directly responsible for the inhibition of protein translation (13–15) and the initiation of upstream events that lead to inflammatory responses (16). We reported the mechanism connecting the inhibition of protein synthesis and the activation of proinflammatory phenomena by demonstrating that ricin mediates activation of stress-activated protein kinases (SAPKs) by producing lesions in the peptidyl transferase center of 28S rRNA (17). The two classes of SAPKs in mammalian cells are the JNKs and the p38 MAPKs, both of which are activated by upstream kinases called SAPKKs and SAPKKKs (18–21). SAPKs belong to the family of MAPKs, which also include ERKs, and are important mediators through which stress signals are transduced to modulate expression of proinflammatory genes (22–24). Recent studies have identified ZAK/MLK7 as the MAP3K whose activation by ricin and related toxins leads ultimately to the phosphorylation and activation of the SAPKs (25). We demonstrated that delivery of ricin to both pulmonary and extrapulmonary tissues mediated the activation of JNK and p38. Another early consequence of ricin exposure is the activation of
NF-κB, a rapid-acting primary transcription factor that induces expression of genes encoding several proinflammatory cytokines and chemokines (12, 16, 26). Primary human airway epithelial cells and primary murine macrophages respond to ricin in vitro through activation of both MAPK and NF-κB (26, 27), but the specific cell types responsible for ricin’s lethal inflammatory effects in vivo remain unclear.

The pulmonary innate immune response is activated not only by pathogens but also by a wide variety of xenobiotic agents (28–30), and alveolar macrophages represent the first line of defense. Evidence from our laboratory suggests that macrophages play a role in mediating the proinflammatory effects of ricin in the lungs. Intra-tracheal administration of ricin leads to almost complete elimination of macrophages in the lungs (10) and intravascular administration of ricin leads to the death of macrophages in multiple organs (31, 32), suggesting that macrophages may serve as primary targets of ricin. Exposure of primary bone marrow-derived and alveolar murine macrophages to ricin in vitro leads to activation of SAPKs, increased mRNA transcripts encoding proinflammatory genes, and the increased production and release of TNF-α, suggesting the possibility that macrophages may mediate the early proinflammatory effects of ricin in vivo (26).

In the present study we investigated the role of pulmonary macrophages in ricin-mediated lung injury by employing a transgenic mouse line that harbors an inducible suicide gene in macrophages and their precursors (33). Using this model, we were able to achieve depletion of macrophages in the lungs of mice treated with the inducer. Compared with nondepleted mice, macrophage-depleted mice displayed markedly decreased inflammatory signs in response to ricin, including decreased recruitment of neutrophils, decreased expression of proinflammatory mediators, and reduced microvascular permeability. Employing mice deficient in IL-1α and IL-1β or wild-type mice co-treated with IL-1Ra, a recombinant human form of the IL-1 receptor antagonist, we determined that ricin-mediated inflammation was dependent on IL-1 signaling. Taken together, our data suggest that the proinflammatory effects of aerosolized ricin are dependent on IL-1 signaling and the presence of pulmonary macrophages, either as the source of IL-1 or as the target cells upon which ricin-induced IL-1 acts to amplify the inflammatory cascade.

Materials and Methods

Animals

C57BL/6J, MAFIA, and TNF-α−/− mice were purchased from The Jackson Laboratory. MAFIA mice are cataloged with The Jackson Laboratory as strain C57BL/6J-Tg(Csf1r-GFP, NGFR/FKBP12)2Bck/J. IL-1Ra/anakinra (Kineret; Amgen) were delivered in the same manner at IL-1Ra/anakinra with 50 mg/kg EBD diluted in saline to a total volume of 200 μL. Lyophilized AP20187 was a gift from Ariad Pharmaceuticals. Lyophilized AP20187 was dissolved in 100% ethanol at a concentration of 62.5 mg/mL stock solution and was stored at −20°C. As recommended by Ariad Pharmaceuticals, injection solutions consisted of 4% ethanol, 10% PEG-400, and 2% Tween 80 in water. All injections were administered i.p. within 30 min after preparation. The volume of injection solution was adjusted according to the average mouse body weight to deliver a dose of 10 mg/kg AP20187 per mouse in an average volume of 100 μL. Mice were injected daily for 5 days before ricin treatment.

Bronchoalveolar lavage (BAL) and neutrophil counts

At 48 h after treatment with ricin, lungs were lavaged four times with 1 ml of cold 0.7% PBS. After lavage, the population of cells from the BAL fluid was examined using Cytospin (Thermo Fisher Scientific) slide preparations and staining by Hema 3 stain set (Fisher Healthcare). Polymorphonuclear neutrophils were identified morphologically and were counted. BAL slides from at least five animals from each treatment group were analyzed to ensure reproducibility.

RNA isolation

Lung tissues were dissected and were immediately frozen and ground in liquid nitrogen. RNA was extracted using TRIzol reagent in accordance with the manufacturer’s instructions and was further digested with DNase. Both reagents were purchased from Invitrogen.

Real-time PCR analysis

Two micrograms of RNA were reverse-transcribed in the presence of SuperScript II and oligo(dT) primer (both reagents were purchased from Invitrogen). The amplification of the cDNA was accomplished using the ABI Prism 7900HT sequence detection system (Applied Biosystems) in the presence of the commercially available SYBR Green PCR Master Mix (Applied Biosystems) and 20 μmol/L of the corresponding sense and antisense RT-PCR primers for 120- bp amplicons in a 40-cycle PCR. Fold induction in gene expression was measured using absolute quantitation of a standard curve in arbitrary units and using levels of GAPDH for normalization. The nucleotide sequences of the primers used in this study have been previously published (16). The denaturing, annealing, and extension conditions of each PCR cycle were 95°C for 15 s, 55°C for 30 s, and 72°C for 30 s, respectively. RNA from five animals per group was analyzed by real-time PCR.

Histology and immunohistochemical analysis

Animals were sacrificed at 48 h unless otherwise indicated. After dissection, lungs were fixed in 4% paraformaldehyde solution for 24 h, at which time tissues were dehydrated and embedded in paraffin. For histology, 5-μm sections were mounted on glass slides, deparaffinized, and stained with H&E following standard procedures. For immunohistochemical analysis of activated caspase-3, Ag retrieval was performed by placing the deparaffinized slides in 10 mmol/L sodium citrate (pH 6) in a microwave oven for 10 min. Immunohistochemical detection of Gr-1/Ly6G did not require pretreatment for Ag retrieval. After blocking the slide, the slides were incubated with primary Abs overnight at 4°C at appropriate dilutions. Slides were further processed using the VectaStain Elite ABC kit (Vector Laboratories) according to the manufacturer’s recommendations using 3,3′-diaminobenzidine as substrate. Immunohistochemical analysis was performed on specimens from at least five animals in each group to verify reproducibility. For examination of embedded lungs, one lung from each animal was sliced transversely into 2-mm slices. To ensure thorough microscopic examination of the entire lung, all slices from each lung were processed, embedded, sectioned by microtomy, and mounted on a single glass slide for examination either by H&E or by immunohistochemistry.

Evans blue dye (EBD) assay

Forty-eight hours after ricin treatment, animals were anesthetized and injected with 50 mg/kg EBD diluted in saline to a total volume of 200 μL through the retro-orbital vein. The dye was allowed to circulate for 30 min, at which time the vasculature was perfused with 5 ml of saline through the right ventricle to remove residual EBD from the systemic circulation before dissection of the lungs. Lungs were removed, weighed, and homogenized; the dye was extracted in 1.5 mL of formamide. The OD of each
samples was determined spectrophotometrically (absorbance 620 nm), and EBD concentration was calculated using a standard curve of known EBD dilutions. Values are expressed in fold induction over saline-control after being normalized for the dry weight of each tissue, and they represent the average of three animals per group.

**ELISA**

Lung tissue was homogenized in lysis buffer in PBS containing 2% Nonidet P-40 and Complete protease inhibitor cocktail (Roche) and centrifuged at 10,000 × g for 20 min at 4°C. The resulting supernatant was analyzed for IL-1β levels using the BD OptEIA mouse IL-1β ELISA set from BD Biosciences per the manufacturer’s instructions. The detection limit for IL-1β was 4 pg/ml. Absorbance of standards and samples were determined spectrophotometrically at 450 nm using a microplate reader (Bio-Rad). Results were plotted against the linear portion of the standard curve. Results are expressed as pg/ml IL-1β and represent the average of three animals per group.

**Lung wet/dry weight ratio**

Left lungs were excised and rapidly weighed for wet weight. Samples were oven dried at 65°C for 72 h to a stable dry lung weight. Data are presented as the ratio of lung wet to dry weight 48 h after either saline or ricin treatment. Five animals from each treatment group were analyzed for wet/dry weight ratios.

**Statistical analysis**

Individual groups were compared using unpaired *t*-test analysis. To estimate *p* values, all statistical analyses were interpreted in a two-tailed manner. Values of *p* < 0.05 were considered to be statistically significant. Kaplan-Meier analysis was performed for survival curves. One-way ANOVA analysis was performed for EBD and wet/dry weight ratio experiments. All data are presented as means ± SEM.

**Results**

Exposure of primary murine macrophages to ricin in vitro causes activation of SAPKs in a dose-dependent manner and triggers the expression of key cytokines, chemokines, and cell-surface recognition molecules involved in proinflammatory signaling (26). To determine whether alveolar macrophages are required for ricin-mediated expression of proinflammatory transcripts in vivo, we compared the responses of wild-type mice with mice depleted of macrophages just before ricin treatment. Transgenic MAFIA mice were treated with AP20187 for 5 days, at which time apoptotic macrophages were detected in the lung tissue and in Cytospin preparations of the cells from BAL fluid (data not shown). Apoptosis was confirmed both by morphologic features and immunocytochemical detection of activated caspase-3, and, consistent with results of previous studies characterizing the MAFIA mouse model, we achieved 90% ablation of macrophages using this method (33). Control and macrophage-depleted MAFIA mice were exposed to either aerosolized saline or 0.3 μg/100 g of aerosolized ricin, a lethal dose that consistently caused death in 100% of wild-type mice within 48–96 h. Forty-eight hours later, lysates of lung tissue were examined by quantitative real-time RT-PCR (qRT-PCR) for expression of mRNA transcripts that encode a variety of proinflammatory genes. When compared with MAFIA mice receiving ethanol vehicle, MAFIA mice receiving AP20187 to deplete macrophages exhibited significantly reduced expression (*p* < 0.01) of several proinflammatory cytokines (IL-1β, TNF-α, IL-6), chemokines (CXCL-1/Gro-α and CCL-2/MCP-1), and the cell adhesion molecule E-selectin (Fig. 1). These data indicate that the presence of macrophages was indeed required for the ricin-induced expression of several proinflammatory mediators in vivo.

Recruitment of neutrophils to the lung parenchyma and alveolar spaces is a hallmark of acute pulmonary inflammation (35). In response to ricin, neutrophils accumulate in the BAL fluid and lung tissue of mice 48 h after administration of ricin (12). To determine whether macrophages were required for ricin to induce the recruitment of neutrophils to the lungs, lung sections and Cytospin preparations of BAL fluid were examined 48 h after administration of ricin. Compared with nondepleted mice, animals depleted of macrophages exhibited dramatically reduced numbers of neutrophils in both the lung parenchyma (Fig. 2a) and the BAL fluid (Fig. 2b), suggesting that the presence of macrophages was required for ricin-mediated neutrophil recruitment to the lungs.

Another hallmark of acute lung injury is enhancement of pulmonary capillary permeability to proteins and fluid across the endothelial cell barrier (36). EBD has been used as a marker of extravascular protein leakage due to its high affinity for binding albumin when injected into the bloodstream (37). To determine the role of alveolar macrophages in ricin-mediated leakage of extravascular proteins, MAFIA mice treated with ricin were injected i.v. with 50 mg/kg EBD 1 h before sacrifice. In lungs of MAFIA mice depleted of macrophages, the concentration of extravasated EBD after ricin exposure was reduced by 60% compared with MAFIA mice that had only received ethanol vehicle before ricin (Fig. 2c). This result suggests that ricin-induced vascular permeability to proteins across the lung endothelial barrier is at least partially dependent on the presence of macrophages.

We showed previously that ricin exposure triggers the production and release of early response cytokines TNF-α and IL-1β (26), which are linked to the generation of inflammatory cascades in multiple lung pathologies (38, 39). TNF-α and IL-1 are considered to be initiator cytokines, inasmuch as their release from macrophages has been shown to orchestrate inflammatory responses. In view of the increased expression of mRNA transcripts encoding TNF-α and IL-1β in response to aerosolized ricin in MAFIA mice (Fig. 1), we sought to determine whether these cytokines were required for ricin to elicit inflammatory responses. To address this question, we first examined TNF-α and IL-1β protein levels in lung homogenates of MAFIA mice treated with ricin. Both groups of MAFIA mice treated with AP20187 exhibited slightly higher basal levels of IL-1β in the lungs compared with nondepleted
mice. However, MAFIA mice whose macrophages had not been depleted displayed a 3- to 4-fold increase in IL-1β after exposure to ricin, while no ricin-mediated induction was observed in IL-1β levels of macrophage-depleted animals (Fig. 3). Meanwhile, TNF-α measured from lung homogenates did not change significantly between groups (data not shown), suggesting that IL-1β may play a larger role in the response to ricin in MAFIA mice.

To further probe the roles of IL-1 and TNF-α in ricin-mediated inflammation, we compared responses of mice harboring null mutations in IL-1α and IL-1β (IL-1αβ−/−) and TNF-α (TNF-α−/−). Lung tissue sections prepared from IL-1αβ−/− at 48 h after ricin treatment showed markedly reduced vascular congestion, destruction of alveoli, and accumulation of neutrophils in the airways compared with wild-type tissue sections (Fig. 4a). Furthermore, IL-1α-deficient animals had reduced neutrophil counts in the BAL fluid (Fig. 4b), improved survival compared with wild-type (Fig. 4c) and diminished pulmonary edema as measured by lung tissue wet/dry ratios (Table I). TNF-α−/− mice, however, failed to show improvement over wild-type animals in inflammatory responses after ricin treatment (data not shown). Taken together, the data in Fig. 3 and Table I indicate that IL-1 was required for ricin to mediate its proinflammatory effects in the pulmonary system and, furthermore, that TNF-α was dispensable for this response.

If the provoked release of IL-1 from alveolar macrophages were responsible for the proinflammatory effects of ricin, then the co-administration of exogenous IL-1 and ricin to IL-1αβ−/− mice should restore the inflammatory responses in these animals. To test this possibility, we administered 15 ng of IL-1β in the presence or absence of ricin and measured the accumulation of neutrophils in the BAL fluid. Whereas IL-1β alone failed to induce recruitment of neutrophils to the BAL fluid of IL-1αβ−/− mice, the co-administration of ricin and IL-1β to these mice resulted in the accumulation of BAL neutrophils to levels observed in wild-type mice treated with ricin alone (compare Figs. 5a and 2b). Taken together with the data shown in Figs. 3 and 4, these results support the conclusion that ricin-mediated pulmonary inflammation is dependent on the presence of IL-1.

To explore the possibility that reduced IL-1β production was a contributing factor in the suppressed inflammatory response observed in macrophage-depleted MAFIA mice, we examined the ability of macrophage-depleted animals to recruit neutrophils to the lungs when exposed to aerosolized ricin in conjunction with exogenous IL-1β. Co-administration of ricin and 15 ng of IL-1β to macrophage-depleted MAFIA mice resulted in a significant (p < 0.01) 10-fold increase in the number of neutrophils in the BAL fluid when compared with mice treated with either IL-1β or ricin alone (Fig. 5b). The data in Fig. 5 demonstrate that IL-1β can contribute to ricin-mediated recruitment of neutrophils in both IL-1αβ−/− and macrophage-depleted mice.

Experimental studies show that recombinant human IL-1R antagonist (IL-1Ra), a competitive inhibitor that interferes with binding of IL-1α and IL-1β to the IL-1R1, may limit the release of cytokines and the development of neutrophilia in animal models of lung inflammation (40–43). To determine the ability of IL-1Ra to block ricin-induced pathology, we administered IL-1Ra to wild-type mice at the time of ricin exposure and examined lung tissue sections after 48 h. Animals receiving IL-1Ra exhibited reduced vascular congestion, destruction of alveoli, and accumulation of neutrophils in the airways compared with animals receiving ricin alone (Fig. 6a). Lungs from IL-1Ra-treated mice had significantly lower expression of proinflammatory transcripts (p < 0.01; Fig. 6b) and reduced appearance of neutrophils in the BAL fluid (Fig. 6c).
These data confirm the central role of IL-1 signaling in the inflammatory response to ricin, and they suggest the possibility that blockade of the IL-1 pathway could be employed to suppress ricin-mediated inflammation.

Discussion

Here we show that acute inflammatory lung injury induced by aerosolized ricin was characterized by increased mRNA expression of proinflammatory genes, production of IL-1β, recruitment of neutrophils to the lungs, and elevated microvascular permeability, and that these responses were dependent on the presence of pulmonary macrophages and IL-1 signaling. Indeed, lung pathology was significantly reduced in animals depleted of macrophages or genetically deficient in IL-1, as well as in wild-type animals injected with IL-1Ra to block IL-1 signaling at the time of ricin exposure. Furthermore, at a dose insufficient to produce inflammatory effects on its own, exogenous IL-1β, when co-administered with ricin to mice deficient in IL-1, recapitulated neutrophil recruitment to the BAL fluid in numbers consistently observed in wild-type animals exposed to ricin alone. Taken together, these results demonstrate that pulmonary IL-1 signaling is essential for the acute inflammatory response to aerosolized ricin.

There is increasing evidence that the inflammatory effects triggered by ricin are responsible for ricin’s lethality (12, 44). Since aerosolized ricin represents the most plausible means of exposure for human populations in the event of a bioterrorist attack, understanding the pulmonary immune response to inhaled ricin toxin is critical for developing therapies aimed at managing symptoms and reducing mortality in case ricin is used as an agent of warfare. In the lung, immune responses to inhaled substances are orchestrated by resident macrophages, which initiate inflammatory cascades.

Table I. Lung wet/dry ratios

<table>
<thead>
<tr>
<th>Treatment</th>
<th>Wild Type</th>
<th>IL-1αβ−/−</th>
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<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>SEM</td>
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<tr>
<td>Saline</td>
<td>4.58</td>
<td>0.17</td>
</tr>
<tr>
<td>Ricin</td>
<td>5.72</td>
<td>0.08a</td>
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*p < 0.05 vs saline.

*p = 0.63 (NS) vs saline; p < 0.05 vs wild type.

FIGURE 4. IL-1αβ−/− mice have markedly reduced neutrophilia and pulmonary edema, and improved survival compared with wild-type mice. IL-1αβ−/− and wild-type animals were administered ricin and euthanized 48 h later. a, Representative photomicrographs of lung sections stained with H&E and labeled with anti-Gr-1/Ly6G. b, Total neutrophils counted in the BAL fluid (n = 5; **, p < 0.01). c, Survival curves comparing IL-1αβ−/− (circles) and wild-type (squares) survival percentages over 9 days following ricin treatment (n = 10; p = 0.002). Error bars represent SEM.

FIGURE 5. Exogenous IL-1β administered in combination with ricin restores ricin-mediated recruitment of neutrophils to the BAL fluid in both IL-1αβ−/− and macrophage-depleted MAFIA mice. Animals were treated with ricin and euthanized 48 h later. a, Total neutrophils counted in the BAL fluid of IL-1αβ−/− mice treated with ricin, 15 ng of IL-1β or ricin, and 15 ng of IL-1β (n = 3; * p < 0.05). b, Total neutrophils counted in the BAL fluid of MAFIA mice pretreated with ethanol vehicle (hatched bars) or AP20187 (white bars) for 5 days followed by administration of either 15 ng of IL-1β or ricin, or 15 ng of IL-1β and ricin (n = 3; * p < 0.05; ns, not significant). Error bars represent SEM.
through the production and release of immunomodulatory mediators like early response cytokines TNF-α and IL-1 (45). We previously demonstrated that ricin triggers the enhanced expression of genes encoding TNF-α and IL-1 from primary murine macrophages in culture (26). During microbial invasion, the release of cytokines and chemokines by macrophages serves to recruit and activate other leukocytes, mainly neutrophils, to help clear pathogens and resolve the infection. Inflammatory injury to local tissue by host immune cells is viewed as collateral damage in an infection (46). However, in the case of ricin exposure, in which there is no pathogen to clear, tissue damage occurs without benefit to the host. In view of this, we hypothesized that suppression of the innate immune response to ricin would serve to reduce inflammatory lung injury and lethality caused by ricin.

Depletion of macrophages resulted in diminished ricin-mediated expression of proinflammatory transcripts (Fig. 1), accumulation of neutrophils (Fig. 2, a and b), microvascular barrier permeability (Fig. 2c), and tissue IL-1β levels (Fig. 3), indicating that macrophages are required for these inflammatory processes. Not surprisingly, since macrophages are important for pulmonary homeostasis and constitute the cell type primarily responsible for the clearance of cell debris (47), the depletion of macrophages from mice caused some inflammation and weight loss that was independent of ricin treatment, consistent with studies published previously on MAFIA mice (33). We established 20% weight loss as a criterion for mortality and, to conform with ethical guidelines established by the Oregon Health and Science University Institutional Animal Care and Use Committee, sacrificed the animals that had reached this endpoint. For this reason, the weight loss consistently observed in MAFIA mice injected with AP20187 alone (10–15%) confounded our ability to perform survival studies. Both groups of macrophage-depleted animals, saline and ricin treated, exhibited some alveolitis (Fig. 2a) and increased IL-1β levels (Fig. 3) over control animals, and these effects were observed only in MAFIA and not AP20187-treated wild-type C57BL/6 mice (data not shown), indicating that the effect was specific to the induction of Fas-mediated apoptosis of macrophages in the transgenic animals. It is known that ligation of Fas on tissue macrophages induces proinflammatory cytokine release that can initiate acute inflammatory responses and tissue injury (48, 49), which may explain the higher level of “basal” inflammation in macrophage-depleted animals. Despite these limitations, inflammatory responses were drastically induced after ricin treatment, and we were able to glean from

FIGURE 6. Coadministration of IL-1Ra with ricin to wild-type mice prevents ricin-mediated inflammatory responses. Wild-type mice received ricin, 30 mg/kg IL-1Ra, or ricin plus 30 mg/kg IL-1Ra and were euthanized 48 h later. a, Representative photomicrographs of lung tissue sections stained with H&E at 10× magnification. b, Total neutrophils counted in the BAL fluid (n = 3; p < 0.05). c, mRNA transcript levels of proinflammatory genes measured from lung homogenates by real-time PCR (n = 3; **, p < 0.01). Error bars represent SEM.
these experiments that macrophages play an essential role in the development of ricin-mediated inflammatory disease.

Since the depletion of macrophages before ricin treatment resulted in an attenuation of inflammatory symptoms, we investigated the requirements for TNF-α and IL-1, the major proinflammatory mediators produced by macrophages. Early response cytokines such as TNF-α and IL-1 act on a variety of cells via cell membrane-bound receptors (50–52) to initiate a proinflammatory cascade resulting in chemokine production, up-regulation of adhesion molecules, transmigration of neutrophils into alveolar compartment and lung interstitium, and the release of proteases and reactive oxygen radicals, which are linked to tissue damage (36, 53). Mice deficient in both TNF-α and IL-1 signaling exhibit impaired neutrophilic inflammation in response to Streptococcus pneumoniae (54). To determine the requirement for these cytokines in the response to ricin, we obtained animals genetically deficient in TNF-α and IL-1α/β and administered aerosolized ricin. Notably, mice deficient in TNF-α showed no improvement over wild-type animals (data not shown), indicating that TNF-α, although induced by ricin (Fig. 1), is dispensable for this response. In contrast, animals lacking IL-1α/β had significant protection from ricin-mediated inflammatory indices (Fig. 4, a and b, and Table 1) and improved survival over wild-type animals (Fig. 4c), although they all eventually succumbed after several days. Furthermore, a 10-fold higher dose of ricin administered by aerosol to the pulmonary system led to death of all animals within 48 h regardless of genotype (data not shown). Because ricin ultimately leads to apoptosis of pulmonary epithelium (27), the death observed in IL-1-deficient animals in the absence of pulmonary inflammation may have resulted from the loss of barrier function and the subsequent entry of ricin into the vascular system.

IL-1β is thought to be a major participant in the pulmonary inflammatory cascade in ARDS; IL-1β, rather than TNF-α, was found to be the major inflammatory mediator in BAL fluid of patients with ARDS (55). IL-1 signaling mediates inflammatory lung injury induced by a variety of stimuli including endotoxemia (42), subacute ozone exposure (56), thermal injury (57), bleomycin administration (43), and experimental ventilation (40). Administration of IL-1 is sufficient to induce chemokine expression and recruitment of neutrophils to the airways (42). Neutrophils responding to cytokine and chemokine gradients accumulate in the airways and contribute to injury and loss of epithelial integrity that characterize ARDS/ALI (35). Indeed, in the present study, mice lacking IL-1α/β or pulmonary macrophages (and thus the major producers of IL-1) showed drastically reduced numbers of neutrophils in the BAL fluid following ricin exposure (Figs. 2b and 4b), and this reduction in neutrophils coincided with reduced edema in IL-1α/β−/− mice (Table 1) and diminished microvascular permeability in macrophage-depleted MAFIA mice (Fig. 2c). Additionally, mice treated with IL-1Ra to block IL-1R1 signaling at the time of ricin exposure exhibited reduced expression of proinflammatory transcripts (Fig. 6c) and accumulation of neutrophils in the BAL fluid (Fig. 6b), supporting a primary role for IL-1 signaling in ricin-mediated inflammation.

Precisely which cells respond to ricin-induced secretion of IL-1 remains unknown. Potential targets of IL-1 include pulmonary macrophages acting in an autocrine or paracrine fashion, microvascular endothelial cells, and alveolar epithelial cells. Type I (TI) and type II (TII) alveolar epithelial cells are known to be capable of responding directly to inflammatory stimuli as well as through inflammatory mediators produced by first-responders. For example, alveolar epithelial cells secrete chemokines MIP-2 (CINC-3/CXCL3) and MCP-1 (CCL2) in response to IL-1β and LPS (58) but require stimulation from macrophage-derived IL-1α to elicit chemokine secretion in response to ozone exposure (59). Similarly, human lung microvascular endothelial cells up-regulate the expression of adhesion molecules and release MCP-1, IL-8, and Gro-α/CXCL1 in response to IL-1 (60), suggesting that these cells may also play an active role in the propagation of inflammation initiated by ricin.

The finding that IL-1Ra suppressed the transcription of other genes following ricin exposure (Fig. 6c) suggests that IL-1 signaling is situated upstream of other events in the inflammatory process. Regulation of IL-1-responsive genes is mediated in large part through activation of the ubiquitously expressed transcription factor NF-κB (61). Nuclear NF-κB activity is rapidly induced in the presence of IL-1 in human lung epithelial cells, and this activation is inhibited by the addition of IL-1 receptor antagonist (61). Activation of NF-κB in airway epithelial cells has been implicated in the development of neutrophilic inflammation by multiple stimuli (54, 62, 63), including ricin (27). We reported previously that NF-κB is activated in human epithelial cells after exposure to ricin, and that targeted inhibition of NF-κB by small interfering RNA results in an inhibition of the expression of proinflammatory genes (27). The data presented herein suggest that ricin-mediated IL-1 signaling may contribute to the activation of NF-κB in the airways to promote the development of acute inflammatory disease.

Although many investigators have elucidated mechanisms of lung inflammation and injury following administration of pathogens and microbial products such as LPS, the mechanisms by which environmental xenobiotics induce lung inflammatory responses are not well understood. In addition to providing insight for development of therapeutic approaches to counteract the use of aerosolized ricin as a bioterrorist agent, an increased understanding of mechanisms underlying ricin’s toxicity could serve to provide greater understanding of how inhaled toxins can cause inflammation, injury, and death.

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Disclosures
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