CD11c⁺/CD11b⁺ Cells Are Critical for Organic Dust–Elicited Murine Lung Inflammation

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Organic dust exposure in the agricultural industry results in significant lung disease. Macrophage infiltrates are increased in the lungs after organic dust exposures, yet the phenotype and functional importance of these cells remain unclear. Using an established intranasal inhalation murine model of dust-induced lung inflammation, animals were treated once or daily for 3 weeks with swine confinement organic dust extract (DE). Repetitive DE treatment for 3 weeks resulted in significant increases in CD11c⁺/CD11b⁺ macrophages in whole lung-associated tissue. These cells displayed increased costimulatory molecule (CD80 and CD86) expression, enhanced phagocytic ability, and an increased production of IL-6, CXCL1, and CXCL2. Similar findings were observed with the CD11c⁺/CD11b⁺ macrophage infiltrate after repetitive exposure to peptidoglycan, a major DE component. To determine the functional importance of macrophages in mediating DE-induced airway inflammation, lung macrophages were selectively depleted using a well-established intranasal clodronate liposome depletion/suicide strategy. First, macrophage depletion by clodronate liposomes resulted in significant reductions in airway neutrophil influx and TNF- α and IL-6 production after a single exposure to DE. In contrast, after repetitive 3-week exposure to DE, airway lavage fluid and lung tissue neutrophils were significantly increased in clodronate liposome-treated mice compared with control mice. A histological examination of lung tissue demonstrated striking increases in alveolar and bronchiolar inflammation, as well as in the size and distribution of cellular aggregates in clodronate-liposome versus saline-liposome groups repetitively exposed to DE. These studies demonstrate that DE elicits activated CD11c⁺/CD11b⁺ macrophages in the lung, which play a critical role in regulating the outcome of DE-induced airway inflammation.

Keywords: macrophage; neutrophil; airway inflammation; peptidoglycan; organic dust

The chronic inhalation of organic dusts or bioaerosols can result in several respiratory diseases including chronic bronchitis, asthma, and obstructive lung disease (1, 2). Agricultural workers

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CLINICAL RELEVANCE

Inhalation exposure to organic dust extract from swine confinement facilities induces an influx of activated $CD11c^+/CD11b^+$ macrophages in the lung. Our findings established lung macrophages as key to the acute response to dust extract, and furthermore, lung macrophages are critical in down-regulating inflammatory responses after prolonged, repetitive dust extract exposures. This information may be important because potential therapies such as macrolides and vitamin D were shown to enhance macrophage phagocytic ability or autophagy.

are routinely exposed to organic dust environments, with the highest prevalence of airway disease associated with animal farming exposures, particularly swine confinement operations (2). Initial exposure to organic dust induces an intense proinflammatory response marked by fevers, bronchial hyperresponsiveness, and increases in bronchoalveolar lavage fluid (BALF) inflammatory cells and cytokines/chemokines that wane over time. However, persons chronically exposed to these environments are at increased risk of lung function decline, airway inflammation, and progressive respiratory impairment (1, 3–5). Although this paradigm, termed "chronic adaptation inflammatory response" (1), has been replicated and described in animal models by our group (6) and others (7), the critical cellular mechanisms underlying this response remain unclear.

Lung mononuclear phagocytes and macrophages are important innate immune lung effectors that serve a critical role in host defense against inhaled environmental insults and toxins (8–10). These cells can orchestrate inflammatory responses through the release of mediators that promote inflammation, activate epithelial cells, and elicit neutrophil recruitment (11, 12). Macrophages produce several mediators after in vitro exposure to organic dust extract (DE) that are associated with agriculturally induced disease, such as TNF- α , IL-6, and CXCL8 (1, 13). In the lung, alveolar macrophages are well recognized to represent the largest mononuclear phagocyte population (14). These cells differ from other tissue macrophages based on their high expression levels of CD11c, which is a molecule typically not present on other tissue macrophages, and is generally considered a dendritic cell marker (15-20). Furthermore, CD11b expression, which is high in other tissue macrophage populations, is quite low in alveolar macrophages, unless these cells are activated (16, 17). Lifetime nonsmoking swine farmers demonstrate signs of bronchial inflammation with neutrophils, alveolar macrophages, and lymphocytes present in lavage fluid (21). Furthermore, evidence suggests that the alveolar macrophages in these farmers may be activated, based on observations of increased

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macrophage chemotactic activity and ex vivo oxygen radical formation (21). Airway and lung parenchymal macrophage infiltrates are also increased in mice after organic dust exposure (6, 7). However, the phenotype and functional importance of these cells have not been well described. Therefore, to address the functional role of alveolar macrophages in the context of repetitive DE-induced lung inflammation, we analyzed CD11c⁺ lung macrophages to determine their activation phenotype. In addition, we compared these responses with those of peptidoglycan (PGN), a major component of organic DE previously shown to drive DE-elicited inflammatory responses in macrophages (13). Finally, we selectively depleted lung macrophages using a wellestablished intranasal clodronate liposome macrophage depletion/ suicide strategy to determine the functional importance of these cells in the context of both single and repetitive DE exposures (17, 22, 23).

Our studies demonstrate that both DE and PGN exposure elicited increased numbers of activated $CD11c^+/CD11b^+$ lung macrophages that were critical for regulating the extent of inflammation. Namely, airway inflammatory responses were attenuated after a one-time exposure to DE. However, after repetitive DE exposures, lung inflammatory and pathologic outcomes, primarily marked by neutrophil influx, were significantly worse when macrophages were depleted. Collectively, these studies demonstrate a critical role for lung macrophages in controlling DE-induced lung inflammation.

MATERIALS AND METHODS

Organic DE

DE was prepared as previously described (24, 25), with details provided in the online supplement.

Animal Model

Using an established intranasal instillation murine model of organic dust-induced and PGN-induced airway inflammation (6), C57BL/6 mice (6–8 wk old; Jackson Laboratory, Bar Harbor, ME) were treated once (acute/single exposure) or daily for 3 weeks (repetitive exposure) with 12.5% DE, PGN (100 μ g), or sterile PBS (diluent). The *Staphylococcus aureus* PGN (Sigma, St. Louis, MO) concentration comprises approximately half the protein in 12.5% DE, and elicits airway inflammatory responses similar to those of DE (26). Whole lung cells were isolated, as described previously (27) and in the online supplement. All animal experimental procedures were approved by the Institutional Animal Care and Use Committee of the University of Nebraska Medical Center, according to National Institutes of Health guidelines for the use of rodents.

Macrophage Depletion

The intranasal delivery of encapsulated clodronate liposomes is a wellestablished method to deplete lung macrophages selectively (17, 18, 28). Liposomes (30 μ l) were administered 48 hours before the first DE treatment, and every 3–4 days throughout the 3-week repetitive DE period to maintain macrophage depletion.

BALF

BALF was collected as previously described (6). The total cell number of pooled lavages (3×1 ml lavages) was enumerated, and differential cell counts were determined on cytospin-prepared slides (Cytopro Cytocentrifuge; Wescor, Inc., Logan, UT) stained with Diff-Quik (Dade Behring, Newark, DE). Cell-free supernatants for cytokine analysis from the first lavage were collected and stored at -20° C.

Lung Histology

Whole lungs were inflated to $10 \text{ cm H}_2\text{O}$ pressure with 10% formalin (Sigma) to preserve pulmonary architecture. Lungs were processed and

embedded in paraffin, and sections $(4-5 \ \mu m)$ were cut and stained with hematoxylin and eosin. Slides were semiquantitatively assessed, using a previously described lung inflammatory scoring system (6, 26) by a pathologist (W.W.W.) blinded to the treatment conditions.

Flow Cytometry

Whole lung cells from each animal were incubated with anti-CD16/32 (Fc Block; BD Biosciences, San Jose, CA), and then stained with monoclonal antibodies directed against CD11c, CD11b, IA-b, CD80, CD86, Ly-6G, and appropriate isotype control samples (BD Biosciences). The CD11c⁺ macrophage gating strategy is depicted in Figure E1 in the online supplement. The CD11c⁺ macrophage number was calculated by multiplying the percentage of gated cells measured by flow cytometry by the original hemocytometer count of total cells recovered for each animal.

Phagocytosis Assay

The phagocytic capacity of lung-associated $CD11c^+$ cells was assessed by flow cytometry, modified from methods described previously (10, 13) and in the online supplement.

Cytokine Assays

CD11c⁺ lung cells isolated by FACS from mice repetitively exposed to saline and DE were analyzed for *ex vivo* cytokine/chemokine (e.g., TNF- α , IL-6, KC/CXCL1, or MIP-2/CXCL2) responsiveness to subsequent DE restimulation by ELISA. Details are provided in the online supplement.

Statistical Methods

Data are presented as means \pm SEM. Statistical analyses were performed using a two-tailed, nonpaired *t* test and one-way ANOVA with the Tukey multicomparison *post hoc* test. We used GraphPad (version 5.01; GraphPad Software, La Jolla, CA) software. In all analyses, *P* values less than 0.05 were considered statistically significant.

RESULTS

Organic Dust Exposure Induces the Accumulation of Activated CD11c⁺/CD11b⁺ Lung Macrophages

The total number of lung-associated cells had increased after 3-week repetitive DE exposure compared with saline control samples, concomitant with significant elevations in the absolute number of CD11c⁺ cells (Figure 1A). The expression of costimulatory molecules (CD80 and CD86) and of the adhesion molecule CD11b was significantly up-regulated in lung-associated CD11c⁺ cells after DE challenge, compared with saline-treated control samples (Figure 1B). No change was observed in MHC Class II expression.

A significant increase was evident in the phagocytic ability of lung CD11c⁺ macrophages after repetitive DE challenge, compared with saline-treated mice (Figures 2A and 2B). In addition, IL-6 and CXCL1 production were significantly increased in ex vivo stimulated CD11c⁺ macrophages obtained from DE-treated animals compared with saline-treated mice, whereas TNF- α was significantly decreased (Figure 2C). Interestingly, CD11c⁺ cells from DE-treated mice produced elevated concentrations of IL-6, CXCL1, and CXCL2 at baseline, compared with saline-treated mice. Collectively, these data demonstrate that repetitive intranasal inhalation challenge with DE evokes a significant influx of CD11c⁺ lung cells, demonstrating an activated phenotype as revealed by increased CD11b and costimulatory molecule expression, phagocytic ability, and selected cytokine responsiveness to subsequent DE exposure.



Figure 1. Repetitive organic dust extract (DE) exposure leads to the activation of CD11c⁺ lung macrophages. Lung-associated cells were collected from C57BL/6 mice after 3 weeks of repetitive intranasal exposure to DE or saline control, and stained for FACS. (*A*) Results represent total lung cells and absolute CD11c⁺ cells. #, number of. (*B*) Mean fluorescence intensity (MFI) of surface marker expression on gated CD11c⁺ cells is shown. Results represent the mean \pm SEM (n = 6 mice/group), with statistical significance denoted by asterisks (*P < 0.05, **P < 0.01, and ***P < 0.001).

Repetitive Intranasal Challenge with PGN Induces Activated CD11c⁺/CD11b⁺ Macrophages

Because emerging evidence suggests that bacterial PGNs in industrial animal farming environments are highly prevalent and may be responsible for mediating airway disease (29, 30), and because our previous work demonstrated that PGN exposureinduced outcomes closely resemble those elicited by DE exposure (13, 26), we investigated the effects of intranasal PGN inhalation in separate experiments. Similar to what was observed among DE-treated mice, PGN induced an expansion in total lung cell infiltrates, with an increase in total number of CD11c⁺ macrophages (Figure 3A). Repetitive PGN exposure resulted in increased costimulatory molecule expression (CD80 and CD86) and increased adhesion molecule CD11b expression (Figure 3B) in CD11c⁺ lung macrophages, consistent with the observations already described for DE. However, unlike treatment with DE, PGN treatment significantly augmented MHC Class II expression in CD11c⁺ cells (Figure 3B). The phagocytic ability of lung CD11c⁺ cells from PGN-challenged mice was also significantly enhanced compared with saline-treated mice (Figure 3B). Collectively, these results show that repetitive PGN

treatment results in a significant influx of CD11c⁺ lung cells, demonstrating an activated phenotype, similar to findings observed after DE exposure.

Effects of Lung Macrophage Depletion in Acute DE-Induced Airway Inflammation

Next, we sought to assess the functional importance of lung macrophages in mediating DE-induced airway inflammation by selectively depleting these cells via the intranasal delivery of clodronate-encapsulated liposomes. To establish the efficacy and safety of this approach, we first investigated airway inflammatory outcomes in an acute DE-induced airway model (6). Clodronate liposomes (CL-LIP) and saline liposomes (SL-LIP) were administered 2 days before a one-time (acute) exposure to DE or saline control, whereupon mice were killed 5 hours after DE exposure. Macrophage reduction was confirmed by a microscopic review of the hematoxylin and eosin staining of lung sections. Decreases in macrophage infiltrates (> 80%) were observed in CL-LIP-treated mice, compared with SL-LIP-treated animals (Figure 4). Significant reductions in macrophage recovery (> 80%) from BALF in CL-LIP + saline and CL-LIP + DE animals were evident, compared with respective SL-LIP control mice (Figure 5A).

Macrophage reduction also produced changes in acute DEinduced airway inflammation. DE-induced airway neutrophil influx was nonsignificantly (P = 0.11) reduced in CL-LIP + DE-treated mice compared with SL-LIP + DE mice (Figure 5A). Significant decreases in TNF- α and IL-6 concentrations in CL-LIP + DE-treated animals were evident, compared with SL-LIP + DE-treated animals (Figure 5B). Interestingly, the expression of neutrophil chemokines was differently affected by macrophage infiltrates, because CXCL1 was significantly elevated in CL-LIP + DE-treated mice compared with SL-LIP + DE-treated mice, whereas CXCL2 concentrations were similar (Figure 5B). These studies demonstrate that macrophages play a selective role in regulating acute (single exposure) DE-induced airway inflammatory responses.

Effects of Alveolar Macrophage Depletion in Repetitive DE-Induced Airway Inflammation

Next, we investigated the role of macrophages in the repetitive exposure model to determine the functional importance of macrophages in mediating the chronic inflammatory adaptation-like response to DE. The administration of CL-LIP throughout the duration of repetitive DE exposure treatment (CL-LIP + DE) resulted in increased neutrophil numbers recovered from BALF, compared with SL-LIP + DE-treated animals (Figure 6A). As expected, macrophage recovery in BALF and whole-lung tissue was significantly reduced in CL-LIP + DE-treated mice compared with SL-LIP + DE-treated animals, confirming the efficacy of the clodronate liposome-mediated depletion approach (Figures 6A and 6B). CL-LIP did not alter the activation profiles of the few CD11c⁺ macrophages that were detected in the lungs after repetitive DE exposure, compared with SL-LIP-treated mice. Namely, CD11b, CD80, and CD86 expression had all increased (data not shown). The percentage and numbers of lung neutrophils from whole-lung tissue were significantly increased in animals treated with CL-LIP + DE, compared with SL-LIP + DE (Figure 6B). After 3 weeks of repetitive exposure, BALF cytokines and chemokines were at the lower limit of assay detection, consistent with the chronic inflammatory adaptation response (6, 7), and concentrations were not different between SL-LIP + DE and CL-LIP + DE (data not shown).

The histological examination of lung tissue demonstrated striking differences in lung parenchymal inflammation between the



or saline (0% DE) for 24 hours, and cytokine/chemokine concentrations were measured in cell-free supernatant by ELISA. Results represent the mean \pm SEM (n = 6-9 mice/group), with statistical significance between saline-and DE-treated denoted by asterisks (*P < 0.05 and ***P < 0.001).

CL-LIP + DE and SL-LIP + DE treatment groups (Figures 7A and 7B). Semiquantitative analysis of these DE-induced histopathologic changes were assessed, using an established lung pathology

scoring system (inflammatory score range of 0-3) (6). Significant increases were observed in the inflammatory scores for all parameters assessed in the CL-LIP + DE treatment group

Figure 3. Repetitive intranasal challenge with peptidoglycan (PGN) increases activated CD11c⁺ lung macrophages. C57BL/6 mice were repetitively exposed to PGN (100 µg) or saline for 3 weeks. and lung-associated cells were enumerated and stained for FACS. (A) Results represent the means \pm SEM (n = 6mice/group) of total lung cells and CD11c⁺ lung macrophages. (B) The mean fluorescence intensity (MFI) of surface marker expression on aated CD11c⁺ cells is shown. Next, lungassociated cells were exposed ex vivo to FITC-labeled Saccharomyces cerevisiae zymosan-A bioparticles at 0 and 60 minutes, to determine the phagocytic ability of gated CD11c⁺ lung macrophages. (C) Results represent fold changes in MFI (proportion of cells in the gated population at 60 min, compared with cells exposed for 0 min), expressed as means \pm SEM (n =6 mice/group). Statistical significance is denoted by asterisks (*P < 0.05 and ****P* < 0.001).



Figure 2. DE treatment increases phagocytosis and cytokine production by CD11c⁺ lung C57BL/6 cells. mice were repetitively exposed to DE or saline for 3 weeks, and lungassociated cells were exposed ex vivo to FITC-labeled Saccharomyces cerevisiae zymosan-A bioparticles at 0 and 60 minutes to determine the phagocytic ability of gated CD11c⁺ lung macrophages. (A) A representative dot plot depicts particle uptake in gated CD11c⁺Ly-6G⁻ macrophages from DE-treated mice and saline control-treated mice as a rightward shift in fluorescence. (B) The phagocytic ability of macrophages is shown as fold change in mean MFI (± SEM) of the proportion of cells in the zymosanexposed gated population at 60 minutes, compared with cells exposed for 0 minutes from DE and saline control mice (n = 6 mice/ group). (C) CD11c⁺ macrophages isolated by FACS from DE-treated mice and saline-exposed mice were restimulated ex vivo with DE (1%)



compared with the SL-LIP + DE treatment group. In particular, the CL-LIP + DE treatment group demonstrated larger and increased numbers of cellular aggregates that were distributed diffusely through the lung. Collectively, these studies demonstrate that lung macrophages play a key role in controlling repetitive DE-induced lung inflammatory responses.

DISCUSSION

Here we report that repetitive exposure to organic DE elicited the influx of activated lung CD11c⁺/CD11b⁺ macrophages, as Figure 4. Clodronate-encapsulated liposome treatment reduces lung alveolar macrophages. C57BL/6 mice were treated with clodronate liposomes (CL-LIP) and saline liposomes (SL-LIP), and exposed to saline or DE 48 hours later. Lung sections of all four mice per group were stained with hematoxylin and eosin, and lung macrophages in peripheral and central lung fields (10 total fields/section) were counted and averaged per individual mouse. (A) Results represent the means ± SEM of lung macrophage numbers per high-power field (HPF) in each group. (B) A representative 4- to 5-µmthick section from each group is shown at ×40 magnification. Arrows indicate alveolar macrophages. Statistical significance is denoted by hatch marks $(^{\#\#P} < 0.001)$ between SL-LIP-treated and CL-LIP DE-treated mice.

marked by enhanced costimulatory molecule expression, phagocytic ability, and selective cytokine/chemokine responsiveness. Exposure to PGN, a major bacterial cell-wall component in porcine and other large animal confinement facilities (29, 30), also resulted in an influx of activated CD11c⁺/CD11b⁺ macrophages. This corroborates previous studies from our group demonstrating a critical role for PGN as a bioactive component for eliciting lung inflammation. This work advances our understanding of the role that the lung macrophage plays in controlling repetitive DE-induced airway inflammation responses, because airway and lung parenchymal inflammation



Figure 5. Alveolar macrophages (Macs) are important in mediating acute airway inflammatory response to DE. C57BL/6 mice were treated with clodronate liposomes (CL-LIP) or saline liposomes (SL-LIP) 2 days before a one-time DE challenge. Mice were subsequently challenged with DE or saline, and bronchoalveolar lavage fluid (BALF) was collected 5 hours after exposure. (*A*) Results represent the means \pm SEM of total cells and cell differentials. (*B*) Results represent mean \pm SEM of cytokine/chemokine concentrations quantitated in cell-free BALF. PBS-only controls are also shown (n = 4 mice per group). Statistical significance is denoted by asterisks (*P < 0.05 and *P < 0.01) between respective saline-treated and DE-treated groups. "P < 0.05 and "P < 0.01 indicate statistical significance between SL-LIP-treated and CL-LIP DE-treated mice."



Figure 6. Repetitive DE-induced neutrophilic influx is increased when lung macrophages are depleted. C57BL/6 mice were treated with clodronate liposomes (CL-LIP) or saline liposomes (SL-LIP) beginning 2 days before the first DE challenge, and then every 3 to 4 days during the daily 3-week repetitive DE or saline exposure. (A) Results represent the means \pm SEM (n = 4 mice/ group) of the total cells and cell differentials recovered from the BALF of mice. Next, lungassociated cells were collected

from mice, and total cells were enumerated and stained by FACS. (*B*) Results represent means \pm SEM (n = 4 mice/group) of CD11c⁺ lung macrophages and neutrophils (percentage of cell type × total lung cell count). Statistical significance is denoted by asterisks (*P < 0.05, *P < 0.01, and ***P < 0.001) between respective saline-treated and DE-treated groups. Hatch marks ("P < 0.05, "#P < 0.01, and "##P < 0.001) indicate statistical differences between SL-LIP + DE-treated and CL-LIP + DE-treated mice.

marked by neutrophil influx became worse after macrophage depletion by clodronate liposomes. These experiments establish that DE-activated $CD11c^+/CD11b^+$ macrophages are, in part, responsible for down-regulating the chronic inflammatory lung response to organic dust exposures.

The chronic inflammatory adaptation response to organic dust environments is well-recognized (3), but the important underlying cellular mechanisms have not been clear. Alveolar macrophages are key innate immune cells that are rapidly activated and orchestrate immune responses after exposure to inhaled environmental toxins, such as organic dust, bacterial products, particulate air pollution, and ozone (9, 13, 31). Macrophages can produce inflammatory mediators and regulate and resolve chronic inflammatory responses by clearing bacteria, debris, and apoptotic cells (32). Activated CD11c⁺ lung macrophages are a common feature of inflammatory responses after exposure to various viral or bacterial pathogens (12, 16, 20, 33, 34). Our findings support that repetitive exposure to DE induces an analogous activated CD11c⁺ macrophage phenotype, with findings of high CD11b⁺ and costimulatory molecule expression and phagocytic ability (Figures 1 and 2). DE treatment did not augment MHC Class II expression, which is a feature

that was also observed in other infectious models (22). This may represent a regulatory signal or differences in the recycling of the MHC Class II complex (35). Although lung infections are not a common characteristic of lung disease in farmers, animal facilities are associated with a large diversity of microbial agents, and particularly Gram-positive bacteria (> 80-95%) (29, 30). Indeed, repetitive exposure to PGN, a major cellwall component of Gram-positive and, to a lesser degree, Gramnegative bacteria, also resulted in similar increases of CD11c⁺ macrophage activation.

We found evidence of a "priming effect," as revealed by enhanced IL-6 and CXCL1 after the stimulation of CD11c⁺ cells from DE-challenged mice (Figure 2C). Furthermore, the basal release of IL-6, CXCL1, and CXCL2 was significantly increased in CD11c⁺ cells isolated from DE-treated mice. This would indicate cellular responsiveness triggered by DE *in vivo*, because no additional stimulus was used. However, this "primed" or activated status was not global because TNF- α responsiveness was dampened, consistent with the so-called tolerant response/ adapted response (1, 36). However, a hyperresponsiveness in cytokine release with *ex vivo*, endotoxin-stimulated whole blood from farmers compared with healthy control subjects has been reported (37, 38).



Figure 7. Lung macrophages are important in mediating repetitive DE-induced lung inflammation. C57BL/6 mice were treated with clodronate liposomes (CL-LIP) or saline liposomes (SL-LIP) beginning 2 days before the first DE challenge, and then every 3 to 4 days during the daily 3-week repetitive DE or saline exposure. (A) Semiguantitative inflammatory scores (means ± SEM; n = 4 mice per group) of the degree and distribution of cellular aggregates, and of

alveolar and bronchiolar lung inflammation, are shown. PBS-only control scores are also shown. (*B*) A representative 4- to 5- μ m-thick, hematoxylinand-eosin–stained section of one of four mice per treatment group is shown at ×10 magnification. All lung specimens were inflated to 10 cm H₂O pressure during fixation to avoid atelectasis artifacts. Statistical significance is denoted by asterisks (**P* < 0.05, **P* < 0.01, and ****P* < 0.001) between respective saline-treated and DE-treated groups. **P* < 0.05 and ***P* < 0.01 indicate statistical difference between SL-LIP + DE–treated and CL-LIP + DE–treated mice.

The finding that CD11b expression was dramatically increased in DE-exposed CD11c⁺ cells is also important. Although CD11b is well recognized for its role in leukocyte adhesion and activation, phagocytosis, and transmigration (39), CD11b expression on mononuclear phagocytes within the respiratory tract has been correlated with airway neutrophilia (9, 40), and the correlation with airway neutrophilia is supported by our studies. Activated macrophages may contribute to DE-induced airway neutrophilia. However, CD11c⁺/ CD11b⁺ macrophages are more likely responding to the inflammatory insult in an attempt to control and resolve disease. This possibility is supported by our demonstration that without an adequate lung macrophage population, enhanced inflammatory consequences follow repetitive DE exposure (Figures 6 and 7). Previous work established that the mononuclear cellular aggregates induced by repetitive organic dust exposure comprise an admixture of T and B lymphocytes and phagocytes (6). Furthermore, DE-induced cellular aggregates consist of live cells (> 98.5%), as opposed to clumps of apoptotic cells based on the common method for detecting DNA fragmentation, terminal deoxynucleotidyl transferase dUTP nick end labeling (TUNEL) staining of tissue sections (data not shown).

In acute DE exposure studies within the setting of lung macrophage depletion, findings suggest that the alveolar macrophage is likely a major source for TNF- α and IL-6 production after acute DE exposure (Figure 5). In contrast, the macrophage does not appear to be an important source of neutrophil chemoattractants during DE-induced airway inflammation (Figure 5). This implies that other cell types, such as airway epithelium, may be important sources of DE-induced neutrophil chemoattractants. Furthermore, TNF- α expression was demonstrated in airway epithelial cells in all groups by an immunohistochemical staining procedure (Figure E3). Isolated airway epithelial cells produce substantial amounts of neutrophil chemoattractants after organic dust exposures (25, 41, 42). Without macrophages to phagocytize and remove the organic dust burden effectively, the airway epithelium may hyperrespond to an increased DE load.

Trends toward a reduction in DE-induced acute neutrophil counts were observed in the setting of macrophage depletion, despite the lack of reduction in murine neutrophil chemoattractants. With repetitive DE exposures, an adaptation-like response occurs, marked by decreased CXCL1 and CXCL2 expression, although neutrophil influx remains elevated (6, 7). Likewise, in our study, the cytokine/chemokine response diminished and did not differ between groups (data not shown), yet neutrophil influx remained elevated after repetitive DE exposures, and this elevation was further enhanced when macrophages were deleted (Figures 6 and 7). These observations highlight that the control and regulation of neutrophil recruitment after lung injury are not limited to chemokines, but may include a number of complex factors and networks (e.g., integrins, selectins, proteases, and reactive oxygen species) (43). Moreover, extracts of organic dusts from agricultural environments exhibit direct chemotactic activity in vitro, and this response has been shown to be independent of endotoxin (44, 45). This response may also be driven by N-formyl-methionyl peptides, motifs of microbial proteins, which can also trigger neutrophil recruitment via the formylated peptide receptor (46). Future lines of study should explore other mechanisms of neutrophil influx, and investigate and characterize subpopulations of DE-induced lung macrophages (e.g., M1, M2a, M2b, and M2c) (47).

Although we found no evidence of distress or mortality, mice receiving repetitive clodronate liposome treatment, regardless of subsequent DE or saline exposure, failed to exhibit equivalent weight gain, compared with their respective control mice. Although the reason for this observation is not entirely clear, repeated clodronate liposome treatment (i.e., 2–3 repeated challenges) was reported to cause subtle immunologic changes (18, 48). By extension, the host may elicit a compensatory mechanism to respond to the reduction in macrophages. Thus, we cannot completely eliminate the possibility of a nonspecific clodronate liposome effect in our repetitive delivery studies.

In conclusion, inhalation exposure to organic DE from swine confinement facilities induces an influx of activated CD11c⁺/CD11b⁺ macrophages in the lung. Moreover, PGN, a component of organic dust, elicited a similar lung response, suggesting that PGN may be an important player in complex organic dust-induced pathogenic lung inflammation. Our findings establish that lung macrophages are key in the acute response to DE, and furthermore, lung macrophages are critical in down-regulating inflammatory responses after prolonged, repetitive DE exposures. This information may be important because potential therapies such as macrolides and vitamin D were shown to enhance macrophage phagocytic ability or autophagy (49–52). Future studies are warranted to investigate whether promoting macrophage function might ultimately lead to novel approaches to reduce chronic organic dust-induced airway inflammatory consequences in exposed agricultural workers.

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