GM-CSF–deficient mice are susceptible to pulmonary group B streptococcal infection

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Granulocyte-macrophage colony-stimulating factor (GM-CSF) gene-targeted mice (GM+/+) cleared group B streptococcus (GBS) from the lungs more slowly than wild-type mice. Expression of GM-CSF in the respiratory epithelium of GM+/+ mice improved bacterial clearance to levels greater than that in wild-type GM+/+ mice. Acute aerosolization of GM-CSF to GM+/+ mice significantly enhanced clearance of GBS at 24 hours. GBS infection was associated with increased neutrophilic infiltration in lungs of GM+/+ mice, while macrophage infiltration predominated in wild-type mice, suggesting an abnormality in macrophage clearance of bacteria in the absence of GM-CSF. While phagocytosis of GBS was unaltered, production of superoxide radicals and hydrogen peroxide was markedly deficient in macrophages from GM–/– mice. GM-CSF plays an important role in GBS clearance in vivo, mediated in part by its role in enhancing superoxide and hydrogen peroxide production and bacterial killing by alveolar macrophages.


Introduction
Granulocyte-macrophage colony-stimulating factor (GM-CSF) is a 23-kDa glycoprotein produced by multiple cell types including monocytes/macrophages, B lymphocytes, neutrophils, eosinophils, pulmonary type II cells, and other respiratory epithelial cells (1, 2). The functions of GM-CSF include regulation of hematopoietic cell proliferation and differentiation (1), and modulation of the function of mature hematopoietic cells. These effects include enhanced antigen presentation (3), increased complement- and antibody-mediated phagocytosis (4), augmented microbicidal capacity (5, 6), and heightened leukocyte chemotaxis and adhesion (7, 8).

Alveolar macrophages are thought to play a critical role in host defense of the lung. Alveolar macrophages bind, phagocytose, and kill bacteria in association with cellular activation, release of intracellular proteases and reactive oxygen species. Reactive oxygen species are released by activated alveolar macrophages directly killing bacteria. Phagocytic cells exposed to GM-CSF demonstrate increased phagocytosis (6, 9), oxidative metabolism (9, 10), activation of Fc receptors (7), release of chemotactic factors (4, 11), and stimulation of mechanisms for killing intracellular viruses, fungi, bacteria, and protozoa (12, 13).

GM-CSF administration to mice before burn injury and infection with Escherichia coli significantly reduced systemic infection and enhanced survival (14). GM-CSF improved survival in neonatal rats treated before Staphylococcus aureus infection (15). However, survival did not improve in rats treated with GM-CSF after the onset of infection from cecal ligation and puncture (16). Although there is compelling evidence that GM-CSF enhances host defense, its role in the clearance of respiratory pathogens in vivo, has not been demonstrated. To assess the function of GM-CSF in vivo, mice carrying a null allele of the GM-CSF gene were generated through the use of gene targeting techniques in embryonic stem cells (17, 18). Mice homozygous for the GM-CSF null allele (GM–/–) have normal hematopoietic function but develop lung pathology characteristic of pulmonary alveolar proteinosis (17, 18). Lung lavage from GM–/– mice revealed markedly increased tubular myelin figures, surfactant proteins A, B, and C, phospholipid and enlarged, foamy alveolar macrophages. Alveolar proteinosis and alveolar macrophage morphology were corrected in the GM–/– mice by direct expression of mouse GM-CSF in lung epithelial cells using the human surfactant protein C (SP-C) gene promoter (SP-C-GM transgenic mice) (19).

Group B streptococcus (GBS) is the most common bacterial infection causing pneumonia and sepsis in newborn infants. Host responses to GBS include activation of both alveolar macrophages and polymorphonuclear leukocytes (PMNs). Phagocytosis and killing of GBS in the lungs is enhanced by surfactant protein A, which increases phagocytosis and reactive oxygen species–mediated killing (20–22). Because macrophage function is strongly influenced by GM-CSF, the present study tested whether GBS clearance from the lungs was influenced by GM-CSF in vivo. Chronic and acute exposure to GM-CSF protected mice from GBS pneumonia mediated in part, by oxygen radical production.

Methods
Animal husbandry. GM-CSF–deficient mice (GM–/–) were bred from the C57BL/6/129Sv F2, homozygous GM-CSF–deficient mice described previously by Dranoff et al. (17) and were maintained in the C57BL/6 background for several years. GM–/– mice.
have alveolar proteinosis with elevated phospholipid and surfactant proteins in the lung. The human surfactant protein C (SP-C) gene promoter was used to direct expression of mouse GM-CSF in lung epithelial cells in GM-CSF null mutant (GM−/−) mice, resulting in correction of the alveolar proteinosis (SP-C-GM mice) (19). SP-C-GM mice have increased bronchoalveolar lavage (BAL) GM-CSF, normal concentrations of surfactant proteins A and B and phospholipid in BAL fluid and increased numbers of alveolar macrophages under vivarium conditions. The C57BL/6 strain of mice (Taconic Farms, Germantown, New York, USA) used for comparison (GM+/+), GM−/− and SP-C-GM mice were housed and studied under Institutional Animal Care and Use Committee-approved protocols in the animal facility of the Children’s Hospital Research Foundation. Male and female mice weighing 20–25 grams (35–42 days old) were used.

Preparation of bacteria. A stock culture GBS was obtained from a clinical isolate from a newborn infant with systemic infection. Bacteria were suspended in sterile PBS containing 20% glycerol and frozen in aliquots at −80°C. Bacteria from the same passage were used to minimize variations in virulence related to culture conditions. Before each experiment, an aliquot was thawed and plated on tryptic soy/5% defibrinated sheep blood agar and then inoculated into 4 ml of Todd-Hewitt broth (Difco Laboratories, Detroit, Michigan, USA) and grown for 14–16 h at 37°C with continuous shaking. The broth was centrifuged, and the bacteria were washed in PBS at pH 7.2 and resuspended in 4 ml of the buffer. To facilitate studies, a growth curve was generated so the bacterial concentration could be determined spectrophotometrically which was confirmed by quantitative culture of the intratracheal inoculum.

Labeling of bacteria with FITC. Bacteria were harvested from agar plates 24 h after streaking, suspended in 5 ml PBS (pH 7.2), and centrifuged 1 min at 220 g to remove any large aggregates of bacteria. The OD at 660 nm of the bacterial culture was measured to determine bacterial concentration. The suspension was then pelleted at maximum speed in a microfuge and the pellet resuspended in 0.9 ml PBS (pH 7.2) and heated to 95°C for 10 min to kill the bacteria. The heat-killed bacteria were then pelleted and resuspended in 1 ml 0.1 M sodium carbonate (pH 9.0). FITC (Molecular Probes, Eugene, Oregon, USA) was added as a 10 mg/ml stock in DMSO to a final concentration of 0.01 mg/ml, and the suspension was incubated for 1 h in the dark at room temperature with gentle agitation. Labeled bacteria were washed four times for 5 min each time with PBS (pH 7.2) to remove unconjugated fluorophore and finally diluted in PBS and stored in aliquots of 100 μl at −80°C.

Intratracheal inoculation. Administration of GBS into the respiratory tract was performed by intratracheal inoculation of 106 CFU/ml diluted in sterile PBS. Mice were anesthetized with iso-flurane, and an anterior midline incision was used to expose the trachea. A 30-gauge needle attached to a tuberculin syringe was inserted into the trachea and a 100 μl inoculum was dispersed into the lungs. The incision was closed with one drop of Neo-Sang. Nonpyogenic PBS was injected intratracheally as control.

Bacterial clearance. Quantitative cultures of lung and spleen homogenates were performed 6, 24, and 48 h after inoculation of the animals with bacteria. Mice were exsanguinated after a lethal intraperitoneal injection of sodium pentobarbital. The animal was sacrificed via intratracheal instillation of 104 GBS/ml at −80°C. The lungs were retrieved, weighed, and each homogenized in 2 ml of sterile PBS. Homogenate (100 μl) and further dilutions were plated on blood agar plates to quantitate bacteria.

Pathology. Lungs were inflated via a tracheal cannula at 20 cm pressure with 4% paraformaldehyde and removed en bloc from the thorax. Lungs were dehydrated and embedded in paraffin.
every 15 min until 2 h, at 37°C. OD was converted to nanomoles of cytochrome C reduced using a molar extinction coefficient of 21.1 mM/cm. Each measurement was the mean of at least two replicates with eight determinations at each time. Data were expressed as nanomoles cytochrome C reduced per 10^5 cells for total oxygen radical production. Superoxide production was assessed by subtracting activity in the presence of superoxide dismutase from total oxygen radical production. Hydrogen peroxide production was assessed by subtracting activity in the presence of catalase (200 U/10^5 cells) from total oxygen radical production.

**Superoxide anion generation by neutrophils.** Alveolar neutrophils were collected by BAL. Mice infected with 10^6 CFU of GBS had >95% neutrophils in BAL. Because previous studies have demonstrated that surfactant can decrease superoxide generation by neutrophils (24), BAL neutrophils were studied directly after lavage and, in separate experiments, after removal of the surfactant. To separate neutrophils from surfactant, BAL fluid was placed on neutrophil isolation medium (Cardinal Associates, Santa Fe, New Mexico, USA), centrifuged at 1,200 rpm for 30 min, and the band of neutrophils collected. Superoxide generation by neutrophils was analyzed as described for macrophages. Superoxide levels were slightly lower after separation of the surfactant, most likely resulting from partial activation with the factant, most likely resulting from partial activation with the

Data were expressed as nanomoles of cytochrome C reduced using a molar extinction coefficient of 21.1 mM/cm. Each measurement was the mean ± SEM with n = 10 mice per group. *P < 0.05 compared with GM+/+ mice. GBS, group B streptococcus; GM+/+, GM-CSF gene-targeted mice; SP-C-GM, human surfactant protein C (SP-C) gene promoter expressing GM-CSF.

**Results**

**Pulmonary pathology after GBS administration.** To determine an appropriate bacterial dose for study, wild-type mice were inoculated intratracheally with GBS at concentrations of \(10^5\)–\(10^6\) colony-forming units (CFU) (four mice per group). The \(10^6\) CFU dose resulted in 50% mortality with the deaths occurring after 24 hours at this dose. A sub-lethal dose of GBS was chosen for the study. Intratracheal administration of \(10^4\) CFU GBS was well-tolerated, and all animals survived the 48-hour study period. No alterations in activity or physical appearance of the animals were detected throughout the 48 hours of study.

In GM+/+ mice, GBS caused pulmonary infiltrates consisting of both macrophages and polymorphonuclear leukocytes. In contrast, GBS infection caused predominately macrophage infiltrates in GM+/- and SP-C-GM mice 24 and 48 hours after infection. GBS administration to GM+/- mice caused mild-to-moderate pulmonary inflammation in four of five mice studied at 48 hours. A
similar infiltrate was detected in only one of five GM–/– mice 48 hours after infection (data not shown). GM–/– mice have baseline histologic changes in the lung consisting of accumulation of eosinophilic material in the alveolar spaces and mononuclear cell infiltrates that progress with age. Uninfected GM–/– mice used in the current study (five to six weeks of age) had minimal pulmonary infiltrates. Pulmonary infiltrates were not observed in GM–/– mice inoculated with sterile PBS (n = 6).

Decreased bacterial clearance in GM–/– mice. GBS proliferated in the lungs of GM–/– mice and bacteria were more numerous in lung homogenates compared with GM+/+ controls (Fig. 1). GBS clearance was markedly improved in the bitransgenic SP-C-GM mice, to levels better than that seen in GM+/+ mice at 6 hours (Fig. 1). Systemic dissemination of the bacteria was assessed by quantitative culture of the spleen. At 6 hours, no GBS was isolated from the spleens of GM+/+, SP-C-GM, or GM–/– mice. GBS was detected in spleen homogenates in both GM–/– and GM+/+ mice at 24 hours; however, colony counts were not different (data not shown).

Phagocytosis of GBS by alveolar macrophages. GBS infection (10⁴ CFU) increased the number of macrophages and PMNs in BAL fluid from GM+/+ and GM–/– mice. However, BAL fluid from GM–/– mice infected with GBS contained a greater percentage of neutrophils at 6 (5.4 ± 1.2 vs. 1.8 ± 0.8), 24 (27.8 ± 0.8 vs. 0), and 48 hours (34.0 ± 2.4 vs. 0) compared with BAL fluid from GM+/+ mice, respectively; mean ± SEM, n = 5 mice per group, P < 0.05 compared with GM+/+ mice.

Control experiments in which sterile 0.9% NaCl was injected intratracheally showed that the inoculation procedure did not alter the cell counts in BAL fluid. BAL fluid from uninfected GM–/– mice contained a slightly lower percentage of macrophages than BAL fluid from uninfected GM+/+ mice (93.4 ± 0.4 vs. 98.6 ± 0.6, respectively; mean ± SEM, n = 5 mice per group, P < 0.05). FITC-labeled bacteria and stained macrophages from the infected animals were visualized under confocal microscopy with three-dimensional imaging. Serial sections through the cell confirmed the presence of intracellular bacteria. Phagocytosis of GBS by alveolar macrophages was similar in GM+/+ and GM–/– mice and mice inoculated with sterile PBS (n = 6).

Decreased superoxide production by alveolar macrophages from GM–/– mice after GBS infection. Superoxide production was assessed in cells isolated from BAL fluid 18 hours after intratracheal administration of GBS. BAL fluid from both GM–/–, GM+/+, and SP-C-GM mice contained >90% macrophages after GBS infection (10⁴ CFU). After stimulation with PMA, superoxide radical, and hydrogen peroxide production by alveolar macrophages was significantly decreased in GM–/– compared with GM+/+ mice. Macrophages from SP-C-GM mice generated greater superoxide and similar amounts of hydrogen peroxide compared with macrophages from GM–/– mice (Table 1).

Superoxide production by neutrophils is unaltered in GM–/– mice. Superoxide production was assessed in cells isolated from BAL fluid 18 hours after intratracheal administration of GBS (10⁶ CFU). At this concentration of GBS, BAL fluid from both GM–/– and GM+/+ mice contained > 90% neutrophils. Stimulation of BAL neutrophils with PMA produced a maximal level of superoxide at 60–75 minutes. Superoxide radical production by neutrophils was similar in cells from GM–/– and GM+/+ mice (4.6 ± 1.8 and 8.5 ± 3.1 nm cytochrome C reduced per 10⁵ cells, respectively; mean ± SEM, n = 8). Similar results were obtained when neutrophils were separated from surfactant with neutrophil isolation medium; superoxide radical production was 2.5 ± 0.9 for GM–/– and 2.8 ± 1.1 nm cytochrome C reduced per 10⁵ cells (mean ± SEM, n

### Table 1

<table>
<thead>
<tr>
<th>Source of macrophages (Genotype)</th>
<th>Peak superoxide radical</th>
<th>Peak hydrogen peroxide</th>
<th>n</th>
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<tbody>
<tr>
<td>GM–/–</td>
<td>0.46 ± 0.7a</td>
<td>Not detectablea</td>
<td>8</td>
</tr>
<tr>
<td>SP-C-GM</td>
<td>8.68 ± 2.1a</td>
<td>2.17 ± 0.6</td>
<td>8</td>
</tr>
<tr>
<td>GM+/+</td>
<td>3.69 ± 1.6</td>
<td>4.43 ± 1.3</td>
<td>8</td>
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Superoxide and hydrogen peroxide production was assessed in cells isolated from BAL fluid 18 hours after intratracheal administration of GBS. Cells in BAL fluid from GM–/–, GM+/+, and SP-C-GM mice consisted of >90% macrophages. The reduction of cytochrome C in the presence and absence of superoxide dismutase or catalase was measured from PMA-stimulated macrophages. Data are expressed as nanomoles cytochrome C reduced per 10⁵ cells. Superoxide production was assessed by subtracting activity in the presence of superoxide dismutase and hydrogen peroxide production was assessed by subtracting activity in the presence of catalase from total oxygen radical production. Macrophages from SP-C-GM mice generated significantly more superoxide than macrophages from GM–/– and GM+/+ mice. Hydrogen peroxide was not detectable from PMA-stimulated macrophages from GM–/– mice. Data are mean ± SEM for the number of determinations. aP < 0.05 compared with GM+/+ mice. BAL, bronchoalveolar lavage; GBS, group B streptococcus; GM–/–, GM-CSF gene-targeted mice; SP-C-GM, human surfactant protein C (SP-C) gene promoter expressing GM-CSF.
- 8) for GM<sup>+/+</sup> mice. Thus, abnormalities of superoxide production by GM<sup>–/–</sup> neutrophils were not a determinant for the impaired clearance of GBS.

Decreased 8-iso-PGF<sub>2α</sub> in lung homogenates and BAL fluid from GM<sup>+/+</sup> mice. Eighteen hours after GBS infection, 8-iso-PGF<sub>2α</sub> levels were greater in lung homogenates and BAL fluid from GM<sup>+/+</sup> compared with GM<sup>–/–</sup> mice (Fig. 2). F<sub>2</sub>-isoprostanes are prostanoids produced independently of cyclooxygenase by free radical–catalyzed peroxidation of arachidonic acid–containing lipids. These findings are consistent with a role of GM-CSF in oxygen radical generation in the lung.

Cytokine levels in lung homogenates. Twenty-four hours after GBS infection, proinflammatory cytokines TNF-α, IL-6, and MIP-2 were significantly increased in lung homogenates from GM<sup>–/–</sup> compared with GM<sup>+/+</sup> mice (see Fig. 3). TNF-α concentrations in lung homogenates were increased in SP-C-GM mice, while MIP-2 and IL-6 were similar to those in GM<sup>+/+</sup> mice 24 hours after infection (Fig. 3). In contrast, IFN-γ levels were increased in GM<sup>–/–</sup> (27.2 ± 1.3 pg/ml) compared with GM<sup>+/+</sup> (6.7 ± 0.8 pg/ml; mean ± SEM, P < 0.05) mice 24 hours after infection. Increased concentrations of TNF-α in lung homogenates from SP-C-GM mice were associated with enhanced bacterial clearance in these animals. However, increased TNF-α was associated with decreased bacterial clearance in the GM<sup>–/–</sup> mice. TNF-α was undetectable in lung homogenates from uninfected GM<sup>–/–</sup> (data not shown).

Increased nitrite in BAL fluid from GM<sup>–/–</sup> mice. Nitric oxide production after GBS infection was measured as nitrite in BAL fluid. Six and 24 hours after infection, increased nitrite levels were found in BAL fluid from the GM<sup>–/–</sup> compared with GM<sup>+/+</sup> and SP-C-GM mice (Fig. 4). To estimate nitrate production, nitrate reductase was used to convert nitrate to nitrite and total nitrite/nitrate measured in BAL fluid (20). Six and 24 hours after infection, increased nitrite/nitrate levels were found in BAL fluid from the GM<sup>–/–</sup> compared with GM<sup>+/+</sup> and SP-C-GM mice (Fig. 4). Nitrite was undetectable in BAL fluid from uninfected GM<sup>–/–</sup> mice. The observed impairment of bacterial clearance in GM<sup>–/–</sup> mice, despite increased nitrite concentrations in BAL fluid, suggests that the nitrite production was not a critical determinant of clearance of GBS.

Aerosolized GM-CSF enhances GBS clearance. Wild-type mice were treated with exogenous GM-CSF to determine whether acute administration of GM-CSF to the lung enhanced bacterial clearance. Two doses of aerosolized GM-CSF (4 ng/mouse/dose, 20 hours before and at time of infection) did not increase clearance of GBS from the lungs of GM<sup>+/+</sup> mice at 6 hours. Treatment of GM<sup>–/–</sup> mice with three doses of aerosolized GM-CSF (additional dose 6 hours after infection) significantly enhanced clearance of GBS at 24 hours (Fig. 5).

Discussion

Pulmonary clearance of intratracheally administered GBS was reduced in GM<sup>–/–</sup> mice when compared with GM<sup>+/+</sup> mice. Expression of mouse GM-CSF in the respiratory epithelium of GM<sup>–/–</sup> mice under control of the SP-C promoter and acute administration of GM-CSF to GM<sup>–/–</sup> mice improved bacterial clearance to levels greater than wild-type mice. While phagocytosis of bacteria by macrophages was similar in GM<sup>–/–</sup> and GM<sup>+/+</sup> mice, superoxide radical and hydrogen peroxide production were markedly decreased in alveolar macrophages from GM<sup>–/–</sup> mice.

Figure 4
Increased nitrite concentrations in BAL fluid from GM<sup>–/–</sup> mice after GBS infection. Nitrite in the BAL fluid was measured by the Griess reaction as described in Methods. Six and 24 h after intratracheal inoculation of 10<sup>4</sup> CFU GBS, nitrite (a) and nitrite/nitrate (b) levels in BAL fluid from GM<sup>–/–</sup> mice (solid bars) were increased compared with GM<sup>+/+</sup> (hatched bars) and SP-C-GM (open bars) mice. Data represent mean ± SEM with n = 5 mice per group. *P < 0.05 compared with GM<sup>+/+</sup> mice.

Figure 5
Exogenous GM-CSF enhances GBS clearance from the lung. Concentration of GBS was determined by quantitative cultures of lung homogenates. Colony counts were significantly decreased 24 h after administration of 10<sup>4</sup> CFU GBS to wild-type mice (solid) treated with three doses of aerosolized recombinant mouse GM-CSF (4 ng/mouse) compared with untreated controls (hatched bars). Data are mean ± SEM with n = 15 mice per group. *P < 0.05 compared with untreated wild-type mice.
mice. These findings support the concept that GM-CSF plays an important role in pulmonary host defense, mediated in part by increased alveolar macrophage oxygen radical production and bacterial killing.

Surprisingly, impaired pulmonary clearance of GBS in GM+/+ mice was not associated with abnormalities in macrophage phagocytosis. Previous in vitro studies demonstrated that GM-CSF enhanced phagocytosis of serum opsonized Staphylococcus aureus by neutrophils (6) and Cryptococcus neoformans by macrophages (4, 25). The present study did not demonstrate abnormalities in phagocytosis, suggesting that other mechanisms operative in vivo may compensate for the lack of GM-CSF. For example, surfactant proteins A and D (SP-A, SP-D) are markedly increased in BAL from GM+/+ mice (17). These surfactant proteins play an important role in host defense against bacterial pathogens. Surfactant proteins A and D stimulate macrophage chemotaxis (26, 27) and SP-A enhances binding of bacteria to macrophages (20).

Phagocytic cells generate reactive oxygen species that are involved in bacterial killing. Phagocytosis of invading microorganisms leads to increased oxygen consumption, reduction of oxygen to superoxide, which is then secreted into the phagosome where it dismutates to hydrogen peroxide. Production of superoxide and hydrogen peroxide contributes to microbial killing. In the present study, stimulated macrophages from GM+/+ mice generated less superoxide and hydrogen peroxide than GM–/– macrophages, demonstrating a critical role of GM-CSF in oxidant production. In previous studies, GM-CSF enhanced superoxide anion generation from monocytes in vitro (29), and inhibited replication of Trypanosoma cruzi with enhanced oxidative metabolism (30).

Elevated superoxide production (both basal state and after stimulation) was also observed in peritoneal and pleural macrophages isolated from transgenic mice expressing GM-CSF with a retroviral promoter (31), consistent with the present observations that superoxide production was decreased in GM–/– mice. Likewise, local expression of GM-CSF in the respiratory epithelium of the transgenic SP-C-GM mice was associated with enhanced superoxide generation by alveolar macrophages. Concentrations of 8-iso-PGF2α in the lung, a measure of free radical–induced lipid peroxidation, were decreased in GM–/– mice, consistent with the role of GM-CSF in oxygen radical production. F2-isoprostanes are biologically active prostaglandin F2 (PGF2)–like compounds formed nonenzymatically by free radical–catalyzed peroxidation of arachidonic acid (32). Macrophages from GM+/+ mice had impaired production of oxygen radicals (superoxide, hydrogen peroxide), consistent with the finding of decreased lipid peroxidation. The defects in superoxide production and lipid peroxidation by alveolar macrophages could be primary defects resulting from the lack of GM-CSF signaling or alternatively could be secondary defects resulting from the abnormal alveolar macrophages with characteristic enlarged, foamy cytoplasm with lipid inclusions.

Oxidative killing of bacteria can be mediated by the production of nitric oxide, which in the presence of reactive oxygen species is converted to peroxynitrite, a potent bactericidal free radical (33). However, decreased bacterial clearance was associated with increased nitrite concentrations in BAL fluid from GM+/+ mice. Therefore, nitrite production was not a determinant in the clearance of GBS from the lungs. In vitro, NO production by lipopolysaccharide (LPS)-stimulated peritoneal macrophages from GM–/– mice was decreased; however, LPS and interferon-γ increased NO production similarly in wild-type and GM–/– macrophages (34). Nitric oxide has antimicrobial effects against a variety of intracellular pathogens (35); however, it remains unclear whether NO contributes to the ability of murine macrophages to eliminate pyogenic microorganisms in vivo.

Clearance of GBS from the lungs of GM+/+ mice was deficient in spite of increased neutrophil recruitment. Furthermore, differences in superoxide radical generation by neutrophils in GM+/+ and GM–/– mice were not a determinant in the impaired pulmonary clearance of GBS. Increased infiltration of neutrophils after GBS administration was associated with increased cytokine production in the lungs of GM–/– mice. MIP-2, a neutrophil chemoattractant, was elevated, consistent with increased neutrophils in GM–/– BAL fluid. In vitro studies have suggested that GM-CSF may regulate cytokine release, in particular, TNF-α and IL-6 (36, 37). GM-CSF deficiency had no apparent effect on GBS-induced peak TNF-α response, because both GM+ and SP-C-GM mice had greater TNF-α levels in BAL fluid compared with GM–/– mice. Previous studies found decreased serum IL-6 and IFN-γ in GM–/– mice treated with LPS (34), in contrast to the current study demonstrating elevated IL-6 and IFN-γ in BAL fluid from GM–/– mice after GBS infection. It is unclear from the current study whether these differences are directly related to the absence of GM-CSF or to the severity of infection and failure of early bacterial clearance.

Pulmonary clearance of GBS was improved acutely by administration of aerosolized GM-CSF and by genetic replacement of GM-CSF in the respiratory epithelium of transgenic mice. This finding is consistent with previous studies demonstrating that intraperitoneal GM-CSF increased survival and decreased bacterial load in mice infected with Salmonella typhimurium (38). Additionally, GM-CSF decreased the severity of Pneumocystis carinii pneumonia in vivo (39). Although GM–/– mice had greater pulmonary bacterial burden, systemic dissemination was similar in GM–/– and GM+/+ mice. Enhanced infection was observed in GM–/– mice infected intraperitoneally with Listeria monocytogenes (a facultative intracellular bacterium) (40), suggesting that GM-CSF may have a role in host defense outside of the lung. SP-A and SP-D enhance phagocytosis (20), chemotaxis (26, 27), macrophages activation (21, 41), and enhance GBS clearance from the lung (28). The increased concentrations of SP-A and SP-D (17) failed to correct pulmonary infection or to limit systemic spread of GBS in GM–/– mice.

In summary, the present study demonstrates the important role of GM-CSF in pulmonary clearance of GBS in
Effects of GM-CSF on GBS clearance were associated with increased superoxide radical and hydrogen peroxide generation by alveolar macrophages, demonstrating the critical role of GM-CSF in oxidant generation by these cells. Exogenous administration of GM-CSF enhanced clearance of GBS in wild-type mice and therefore, may represent a strategy to prevent or treat pulmonary infections.

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