Anthrax Lethal Toxin Impairs Innate Immune Functions of Alveolar Macrophages and Facilitates \textit{Bacillus anthracis} Survival

Wilson J. Ribot,1 Rekha G. Panchal,2 Katherine C. Brittingham,1 Gordon Ruthel,1 Tara A. Kenny,2 Douglas Lane,2 Bob Curry,1 Timothy A. Hoover,1 Arthur M. Friedlander,1 and Sina Bavari1*

\textit{U.S. Army Medical Research Institute of Infectious Diseases, 1425 Porter Street, Fort Detrick, Maryland 21702, 1 and Target Structure-Based Drug Discovery Group, SAIC-Frederick, Inc., NCI-Frederick, Frederick, Maryland 21702-1201}

Received 20 February 2006/Returned for modification 27 April 2006/Accepted 14 June 2006

Alveolar macrophages (AM) are very important for pulmonary innate immune responses against invading inhaled pathogens because they directly kill the organisms and initiate a cascade of innate and adaptive immune responses. Although several factors contribute to inhalational anthrax, we hypothesized that unimpeded infection of \textit{Bacillus anthracis} is directly linked to disabling the innate immune functions contributed by AM. Here, we investigated the effects of lethal toxin (LT), one of the binary complex virulence factors produced by \textit{B. anthracis}, on freshly isolated nonhuman primate AM. Exposure of AM to doses of LT that killed susceptible macrophages had no effect on the viability of AM, despite complete MEK1 cleavage. Intoxicated AM remained fully capable of \textit{B. anthracis} spore phagocytosis. However, pretreatment of AM with LT resulted in a significant decrease in the clearance of both the Sterne strain and the fully virulent Ames strain of \textit{B. anthracis}, which may have been a result of impaired AM secretion of proinflammatory cytokines. Our data imply that cytolysis does not correlate with MEK1 cleavage, and this is the first report of LT-mediated impairment of nonhuman primate AM bactericidal activity against \textit{B. anthracis}.

Due to their overwhelming presence in the lung, alveolar macrophages (AM) are the principal immune effector cells in the airways. AM and other cells in the lung, such as dendritic cells (DC), detect microbes through a family of receptors called pattern recognition receptors, which detect pathogen-associated molecular pattern components of microbes. AM are avidly phagocytic cells that are capable of directly killing engulfed microbes when pathogen-associated molecular patterns are detected (2). AM play a key role in maintaining anti-inflammatory conditions within the lung during steady state, helping to prevent chronic inflammatory responses to harmless inhaled antigens. However, during microbial invasion, it is primarily the responsibility of AM to initiate innate immune responses in the lung by releasing immunomodulatory products, such as prostaglandins, leukotrienes, chemokines, and cytokines. The secreted products act in concert to recruit and activate other leukocytes from the blood to the site of infection. Activated AM transmigrate to draining lymph nodes, where they present antigens and secrete cytokines to facilitate activation of the adaptive immune system (14).

The innate protective functions provided by AM appear to be manipulated by the dangerous pathogen \textit{Bacillus anthracis} during inhalational anthrax, the most lethal form of the disease. A previous report suggested that inhaled spores are avidly phagocytosed by AM (25), and recently DC were implicated in anthrax pathogenesis (3). Phagocytosis of spores did not stimulate significant inflammation in the lung until at least 5 h postinfection (16). It has also been proposed that during this time spores are transported inside AM (16, 23) or DC (3) to the mediastinal and peribronchial lymph nodes. En route to the lymph nodes, the spores germinate into vegetative bacteria (10, 23), which are capable of secreting anthrax lethal toxin (LT) and edema toxin. In the lymph nodes, bacteria quickly escape from macrophages and are released into the lymphatics (8, 16). From there bacilli are disseminated into the circulatory system and throughout the host (15).

Lethal toxin, a combination of protective antigen and lethal factor secreted by \textit{B. anthracis}, disrupts the mitogen-activated protein kinase (MAPK) signaling pathway of innate immune cells, such as AM and DC, and impairs intracellular signaling required to initiate adaptive immune function (1). To date, no direct causal relationship between disruption of MAPK signaling and LT-mediated cell death has been determined, and the importance of such a relationship for pathogenesis is unclear. Extensive studies of mouse macrophage cell lines demonstrated that LT induces cell death, suggesting that macrophages may be the primary target of LT activity. This hypothesis was supported by a study which showed that mice depleted of macrophages survived lethal doses of LT (11). In contrast, mouse strains whose macrophages were resistant to LT-induced cell death exhibited a higher mortality rate after \textit{B. anthracis} infection than strains with susceptible macrophages (27). Furthermore, human peripheral blood mononuclear cells are resistant to LT-mediated cytotoxicity, suggesting that cell death may not be a primary function of LT in humans (22). A recent study indicated that the cytotoxic effect of LT on three different human monocytic cell lines was dependent on the differentiation state of the cells (13).

The notion that LT modulates innate and adaptive immunity is gaining momentum, but the exact mechanism of action is not known. To further elucidate the effect of LT on immune cells,
**ABSTRACT**

Alveolar macrophages (AM) are very important for pulmonary innate immune responses against invading inhaled pathogens because they directly kill the organisms and initiate a cascade of innate and adaptive immune responses. Although several factors contribute to inhalational anthrax, we hypothesized that unimpeded infection of *Bacillus anthracis* is directly linked to disabling the innate immune functions contributed by AM. Here, we investigated the effects of lethal toxin (LT), one of the binary complex virulence factors produced by *B. anthracis*, on freshly isolated nonhuman primate AM. Exposure of AM to doses of LT that killed susceptible macrophages had no effect on the viability of AM, despite complete MEK1 cleavage. Intoxicated AM remained fully capable of *B. anthracis* spore phagocytosis. However, pretreatment of AM with LT resulted in a significant decrease in the clearance of both the Sterne strain and the fully virulent Ames strain of *B. anthracis*, which may have been a result of impaired AM secretion of proinflammatory cytokines. Our data imply that cytolysis does not correlate with MEK1 cleavage, and this is the first report of LT-mediated impairment of nonhuman primate AM bactericidal activity against *B. anthracis*. 

**SUBJECT TERMS**

*Bacillus anthracis*, anthrax, macrophages, lethal toxin, immune function, laboratory animals, nonhuman primates
we examined the link between toxin exposure and antibacterial responses of nonhuman primate AM.

**MATERIALS AND METHODS**

Cultivation and infection of alveolar macrophages. Nonhuman primate alveolar macrophages from cynomolgus macaques (Macaca fascicularis) were obtained as described previously (6). AM were washed twice in phosphate-buffered saline (PBS) supplemented with 1 mM EDTA and 20 μM of gentamicin. The cells were cultured in Dulbecco modified Eagle medium supplemented with 7.5% fetal bovine serum, 1% minimum essential medium, nonessential amino acids, and 1% glutamine (referred to below as cDMEM). AM were purified by adherence to cell culture plates or glass coverslips and were cultured in cDMEM supplemented with 100 U of penicillin and 100 μg/ml of streptomycin. The purity of AM was determined by May-Grünwald-Giemsa staining of air-dried cytosin smears. Lavaged cells were 97 to 100% macrophages, and trypan blue exclusion indicated that more than 80% of the AM were viable. Before use in experiments, AM cultures were washed four times to remove antibiotics and any dead cells. AM cultured in cDMEM were treated with recombinant LT or spores at a multiplicity of infection (MOI) of 1 to 5, unless indicated otherwise. For all infections of AM, cell culture plates were centrifuged for 5 min at 500 × g to synchronize infection.

**B. anthracis** spores. Culturing, spore preparation, and engineering of green fluorescent protein (GFP)-labeled Sterne spores and nonfluorescent Ames spores were performed as described previously (12, 24). Briefly, to generate GFP-labeled Sterne spores, plasmid pApF8gfp was used in transfections of *B. anthracis* bacilli to produce bacilli that constitutively expressed GFP. Plasmid pApF8gfp contains a kanamycin resistance gene and a synthetic gfp promoter sequence ligated upstream of gfp. The extrachromosomal plasmid replicated during proliferation and segregated with progeny cells. During sporulation, GFP was incorporated inside spores (as determined by immunoelectron microscopy [not shown]), which resulted in GFP-labeled spores. Spores were purified by centrifugation through Renografin (Renold-76; Braaco Diagnostics, Princeton, N.J.). The purified spores were washed three times with cold sterile water and stored at 4°C. Endotoxin-free reagents were used for spore preparation. Spore preparations were incubated at 65°C for 30 min to kill any vegetative cells and to activate the spores before infection of AM. Spores were resuspended in cDMEM immediately before infection.

**Electron microscopy.** For transmission electron microscopy (TEM), AM infected with Ames or Sterne spores for 30 min were fixed at 25°C with 2.5% glutaraldehyde for 1 h. The samples were treated with 1% osmium tetroxide in modified Millonig's buffer (0.1 M PBS, 0.5% dextrose; pH 7.3) (17), stained en bloc with 0.5% uranyl acetate in ethanol, dehydrated in graded ethanol and propylene oxide, and embedded in Poly/Bed 812 resin (Polysciences, Inc., Warrington, PA). Ultrathin sections were placed on 200-mesh nickel grids and stained with 5% uranyl acetate and 0.2% lead citrate.

**Fluorescence microscopy.** For detection of early endosomal antigen-1 (EEA1) on phagosome membranes, AM were purified by adherence to microscope coverslips. AM were incubated with GFP-labeled Sterne spores at an MOI of 10, centrifuged at 500 × g for 5 min at 4°C, washed with cDMEM, and incubated for 15 min at 37°C. Samples were then fixed with 4% paraformaldehyde in PBS for 30 min, washed with PBS, and permeabilized with cold methanol for 10 s. After incubation for 5 min in blocking buffer that contained 2% donkey serum (Sigma), samples were stained with rabbit anti-EEA1 (Affinity BioReagents, Golden, CO), followed by Alexa 594-conjugated goat anti-rabbit antibody and 4′,6-diamidino-2-phenylindole (DAPI) (Molecular Probes, Eugene, OR). EEA1 immunofluorescence and intraphagosomal GFP-labeled bacteria were detected either by wide-field epifluorescence microscopy or by confocal microscopy.

To count ingested GFP-labeled Sterne spores, *B. anthracis* cells located inside AM were distinguished from noninternalized *B. anthracis* cells after 30 min of infection. Nonpermeabilized cells were incubated with a mixture of monoclonal antibodies specific for *B. anthracis* spore exosporum and for bacillus polysaccharide (kindly provided by T. Abshire and J. Ezzell, USAMRIID) to label all spore (Fig. 1C). Occasionally, more than one spore was detected within a single phagosome (Fig. 1B). Similar results were obtained with Ames spores (data not shown). Antibody labeling of *B. anthracis* exosporum and polysaccharide after mild fixation of samples provided an additional way to determine whether GFP-labeled Sterne spores, EE1A, a protein recruited to early phagosomes after endosome fusion, was detected on phagosomes containing *B. anthracis*. This confirmed the endosomal localization of spores (Fig. 1B).

**RESULTS**

**Phagocytosis of *B. anthracis* spores by AM.** To examine whether AM were able to phagocytose spores via the normal endocytic pathway, the cells were infected with Ames or Sterne spores for 30 min to allow phagocytosis. The intracellular location of both Sterne (Fig. 1A) and Ames (data not shown) spores was determined by TEM, which revealed spores surrounded by a phagosomal membrane. Confocal microscopy showed that within 15 min of infection with GFP-labeled Sterne spores, EE1A, a protein recruited to early phagosomes after endosome fusion, was detected on phagosomes containing *B. anthracis*. This confirmed the endosomal localization of spores (Fig. 1B).

**Viability of *B. anthracis* strains after phagocytosis.** We next determined whether AM could kill *B. anthracis*. After infection of AM, bacterial viability was monitored at different time intervals. A continuous rapid decline in the viability of Ames and Sterne until 6.5 h postinfection was observed (Fig. 2). Within 6.5 h, more than 60 to 80% of Ames and Sterne spores were killed by AM. Although we found no statistically significant difference between killing of Ames spores by AM and killing of Sterne spores by AM, the AM tended to kill lower numbers of...
Ames spores than Sterne spores. It is noteworthy that during the 6.5-h incubation, AM were unable to kill all of the spores. This was not due to AM death during Ames and Sterne infections (data not shown).

**Effect of LT on AM viability and MEK1 cleavage.** Since anthrax LT can kill susceptible macrophages, we next examined the sensitivity of nonhuman primate AM to LT-mediated cell death. AM and J774A.1 macrophages were exposed to different concentrations of LT for 4 h, and cell viability was determined by the MTT assay. At the lowest concentration of LT (0.1 μg/ml), less than 10% of J774A.1 cells were viable, compared to the 98% survival of similarly treated AM (Fig. 3A). Increasing the concentration of LT to 10 μg/ml had little additional effect on the viability of AM, as revealed either by microscopy or by the MTT assay. These results demonstrated that LT is not cytolytic for AM compared to J774A.1 indicator macrophages. To determine if endogenous MEK1 is sensitive to LT-mediated cleavage, AM were incubated with different concentrations of LT. Western blot analysis revealed that MEK1 was cleaved both 4 and 16 h following LT treatment (Fig. 3B).

**AM cytokine responses after LT treatment.** Because LT treatment did not induce the death of AM despite cleavage of critical signaling proteins, we next investigated functional impairment of AM by LT. Treatment of AM with LT alone for 4 h resulted in a slight increase in secretion of interleukin-1β (IL-1β) compared to the secretion by untreated AM. There was essentially no effect on tumor necrosis factor alpha (TNF-
addition of LPS to AM elicited production of TNF-α, IL-1β, IL-6, and IL-8, indicating that the AM were capable of producing these cytokines. Simultaneous treatment of AM with LT and LPS resulted in reduced levels of secreted TNF-α, IL-1β, IL-6, and IL-8 after 4 h compared to the levels for AM treated with LPS alone (Fig. 4). These results suggested that the consequence of MEK1 cleavage for AM was impaired induction of proinflammatory cytokine genes partially regulated by signaling in this pathway.

**Effect of LT on AM bactericidal activity against B. anthracis.** Since LT treatment impaired MAPK signaling and suppressed cytokine production, we next tried to determine whether LT-treated AM were phagocytic. The numbers of phagocytosed Sterne or Ames spores for AM that were treated for 16 h with LT or LPS were not significantly different from the numbers of phagocytosed Sterne or Ames spores for untreated AM (Fig. 5A). Therefore, AM phagocytosis was not impaired by LT despite MEK1 cleavage and cytokine suppression.

Next, we examined whether pretreatment of AM with LT impaired their ability to kill B. anthracis. When AM were treated for 16 h with 0.1 or 1 μg/ml of LT and then infected with B. anthracis spores for 4.5 h, the bactericidal activity was substantially reduced, as indicated by enhanced bacterial survival (Fig. 5B). Microscopic examination of AM cultures (Fig. 5C) after 4.5 h of infection with Sterne revealed that the bacteria were able to replicate more efficiently in LT-treated AM cultures than in untreated AM cultures.

**DISCUSSION**

Alveolar macrophages are a component of sentinel immune cells that protect against infections after inhalation of microbial pathogens. These phagocytic cells can engulf microorganisms and/or other foreign particles and mediate an innate immune response through secretion of cytokines and chemokines. The fate of AM and their response to B. anthracis spores...
deposited in the lungs are fundamental aspects of the progression of inhalational anthrax. To characterize events in the lungs immediately after aerosol exposure, we examined the interactions between spores of \(B.\) \(anthracis\) and purified nonhuman primate alveolar macrophages. AM readily phagocytosed \(B.\) \(anthracis\) spores. The fully virulent Ames spores and the attenuated Sterne spores were both observed in phagosomes surrounded by EEA1 within a few minutes after infection. While the mechanism has not been established yet, mouse (21, 26) and guinea pig peritoneal macrophages (7) can kill \(B.\) \(anthracis\). Mouse macrophages phagocytosed germinated Sterne strain 7702 spores and inhibited their growth (9). However, in contrast to nonhuman primate AM (data not shown), murine macrophages did not survive infection.

The MAPK pathway relays environmental signals to the transcriptional machinery in the nucleus. The signaling pathways involving MEK1, MEK2, and MEK3 play a crucial role in activation of macrophages and are directly involved in production of proinflammatory cytokines, such as TNF-\(\alpha\), IL-1\(\beta\), and IL-6. In our study, nonhuman primate AM viability was not affected, but MEK1 cleavage was detected 4 to 16 h following the exposure of AM to LT (Fig. 3B). In addition, LT impaired the ability of LPS-treated AM to secrete cytokines, which was probably a consequence of MEK1 cleavage. There are conflicting data concerning the modulation of cytokines by LT. Sublethal doses of LT induced production of TNF-\(\alpha\) and IL-1\(\beta\) (11) in mouse macrophages, but an absence of this effect has also been reported (4). Additionally, LT has been shown to inhibit the production of nitric oxide (NO) and TNF-\(\alpha\) induced by LT in murine macrophage lines and mouse peritoneal macrophages (20). Excessive amounts of nitric oxide, TNF-\(\alpha\), and IL-1\(\beta\) secreted by macrophages can lead to the development of shock. Mice depleted of macrophages were shown to be insensitive to LT-induced shock, a consequence of the inability of the remaining immune cells to secrete IL-1\(\beta\) (11). It was proposed previously that the impairment of macrophage and dendritic cell (3, 21) cytokine responses by LT may be a means by which \(B.\) \(anthracis\) inhibits the innate immune response, thereby enhancing its ability to establish an infection.

Despite MEK1 cleavage, impairment of cytokine secretion, and loss of bactericidal activity following LT treatment, AM were still capable of phagocytosis (Fig. 5). The inability of AM to completely eradicate \(B.\) \(anthracis\) after 6.5 h of infection with spores may have been due to accumulation of LT to critical intoxicating levels inside AM or in the tissue culture medium. Therefore, the primary function of LT in anthrax pathogenesis may be to facilitate bacterial survival and outgrowth from AM, but not AM death. A similar LT-induced loss of guinea pig AM bactericidal activity was described by Gimenez et al. (7), who attributed the effect to LT-inhibited group IIA secreted phospholipase A2. However, we did not detect group IIA secreted phospholipase A2 in the \(B.\) \(anthracis\)-infected AM culture media (data not shown). It is not known exactly how much LT is produced by \(B.\) \(anthracis\) during infection, and the amount may depend on the stage of infection. In our study, we were able to observe modulation of AM in the presence of submicromolar amounts of LT. The concentration of LT may be important in pathogenesis, and larger amounts of LT may be relevant during the septicemic stage of anthrax, at which time measurable amounts of LT have been found in the plasma of infected animals (5, 19).

Taken together, these findings suggest that modulation of cytokine responses by LT is a means by which \(B.\) \(anthracis\) might survive the bactericidal activity of macrophages in the later stages of an infection. The inability of AM to completely eradicate bacteria suggests that intracellularly secreted lethal
factor, edema factor, or other virulence factors help overcome the lethal consequences of ingestion by AM. The inhibitory effects of LT on the bactericidal activity of AM may be similar to the effects that LT has on other resident and circulating macrophages in later stages of infection. Our data suggest that following AM infection spores quickly reach endosomal compartments. Subsequent phagosome maturation may lead to killing of some of the organisms. However, in numerous experiments we have observed the presence of apparently dormant B. anthracis spores residing in phagosomes within macrophages of various origins. Such spores can be observed readily by confocal microscopy of RAW cells (24). The inherit ability of spores to withstand harsh conditions may help prevent their killing by AM. Delayed germination of B. anthracis spores, until transport to the lymph nodes, may facilitate their survival until a more favorable environment is encountered. Release from alveolar macrophages and subsequent germination within the lymph fluids might allow sufficient synthesis of LT and edema toxin, which may help overcome the bactericidal activity of circulating macrophages. This would further enhance survival, growth, and dissemination in an infected mammalian host. It would therefore be critical that antitoxin therapeutics be administered before LT disables the host innate immune response.

ACKNOWLEDGMENTS

We thank Kathleen Kuehl for excellent help with electron micrograph studies and David Heath for critical reading of the manuscript. This work was partially funded by DTRA-JSTO project B X003 04. RD B. This project was funded in whole or in part with funds from the National Cancer Institute, National Institutes of Health, under contract N01-CO-12400. This research was supported in part by the Developmental Therapeutics Program in the Division of Cancer Treatment and Diagnosis of the National Cancer Institute. K.C.B. is a recipient of National Research Council fellowships.

The contents of this paper do not necessarily reflect the views or policies of the Department of Health and Human Services, nor does mention of trade names, commercial products, or organizations imply endorsement by the U.S. Government. The opinions, interpretations, conclusions, and recommendations are those of the authors and are not necessarily endorsed by the U.S. Army.

REFERENCES


Editor: V. J. DiRita