Burkholderia pseudomallei aerosol infection results in differential inflammatory responses in BALB/c and C57Bl/6 mice

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INTRODUCTION

Burkholderia pseudomallei is the causative agent of melioidosis, a disease that is endemic in South-East Asia and northern Australia. Human infection results in many possible clinical manifestations (Wiersinga et al., 2006), with pneumonia being the most common presentation in approximately half of all reported cases (Cheng & Currie, 2005). Though lung involvement is likely to be due to haemato-gonous spread following percutaneous inoculation, it is likely that, under certain environmental conditions, inhalation is a significant mode of infection (Currie, 2003). This is evidenced by reports of soldiers exposed to dust from helicopter rotor downwash in Vietnam (Howe et al., 1971) and the association between heavy rainfalls and increased incidence of pneumonic and septicaemic presentations with poor clinical outcomes (Cheng et al., 2006; Currie, 2003; Currie & Jacups, 2003; Liu et al., 2006). Furthermore, B. pseudomallei is also a potential bioterror agent and the most likely mode of dissemination during a biological attack is through an infectious aerosol. It is thus important to understand the pathogenesis of melioidosis following an inhalational exposure to B. pseudomallei.

Currently, by adopting artificial routes of infection, many research groups have demonstrated that differential susceptibility to B. pseudomallei infection exists between BALB/c and C57Bl/6 mice, with the latter being more resistant (Hoppe et al., 1999; Leakey et al., 1998; Liu et al., 2002). However, Jeddeloh et al. (2003) reported that the two strains of mice displayed similar sensitivities to nebulized B. pseudomallei and ruled out the contribution made by genetic differences in the host to the disease. This
finding suggests that an exposure to aerosolized *B. pseudomallei* results in fundamentally different pathogenesis and disease outcome, compared with inoculation by intranasal droplets of bacterial culture. Should this observation be correct, it would invalidate many previous studies on the pathogenesis of inhalational melioidosis. Jeddeloh et al. (2003), however, had used a whole-body aerosol system, which could have resulted in deposition of bacteria on the animals in addition to entry into the lungs. In order to provide an accurate model for pneumonic melioidosis in humans and to resolve the current discrepancy in the various murine models, we established a nose-only aerosol infection system to characterize accurately the early events that occur in post-inhalational challenge. Furthermore, we also characterized for the first time the induction of 17 chemokine and cytokine responses at the protein level.

**METHODS**

**Animals.** Female 4- to 8-week-old BALB/c and C57Bl/6 mice were purchased from the Center for Animal Resources of the National University of Singapore and housed in polystyrene cages with a bedding of wood shavings. The mice were fed with commercial pellets (PMI Nutrition International) and potable water ad libitum. The cages were sterilized for the first time the induction of 17 chemokine and cytokine responses at the protein level.

**Bacteria.** Three different strains of *B. pseudomallei* (KHW, EB6103 and K96243) were used in this study. KHW was isolated from a 19-year-old Singaporean serviceman who died from melioidosis in 1989. EB6103 was isolated from a 19-year-old Singaporean male who succumbed to melioidosis in the 2004 outbreak (Liu et al., 2006). K96243 was isolated from a 34-year-old Thai diabetic woman in 1996. For infection experiments, bacteria were cultured on trypticase soy agar (TSA; BD Diagnostics) for 24 h at 37 °C. Colonies were resuspended in sterile PBS to an OD600 of 1.9 (equivalent to 1 × 10⁹ c.f.u. ml⁻¹). The bacterial suspension was diluted to the desired concentrations for inoculation into mice and aliquots of these suspensions were plated on TSA in duplicate for retrospective determination of the number of viable bacteria inoculated into the mouse.

**Aerosol generation.** The aerosol chamber was housed within a flexiglass glove box. Bacterial suspensions of 10⁻¹⁻¹⁰⁶ c.f.u. ml⁻¹ in PBS were placed in three collision air jet nebulizers. Nebulization of each suspension was driven by compressed air at a pressure of 1–1.5 bar for 10 min to generate aerosol particles of size 1–5 μm. Aerosol density was measured by bubbling 10 l of the air in the aerosol chamber into 20 ml PBS in the impinger. Aliquots of the suspensions were diluted and plated in duplicate on TSA for viable counts. Good linear correlation between the bacterial concentrations in PBS (c.f.u. ml⁻¹) and bacterial density (c.f.u. per litre of air) in the aerosols was observed and the results were highly reproducible. Enumeration of viable *B. pseudomallei* showed that the bacterial counts in the aerosol were consistently 10⁻³-fold lower than those in the corresponding bacterial suspensions.

**Aerosol infection of mice.** Mice were restrained in 50 ml Falcon tubes in the aerosol chamber and exposed in a nose-only manner to the aerosols of *B. pseudomallei* for 30 min. To determine the actual inhaled dose (AID) after aerosol challenge, three mice from each group were sacrificed immediately post-exposure, and their lungs were aseptically removed. The actual number of bacteria that entered the lungs was determined as described below. The theoretical inhaled dose (TID) was calculated according to the method of Guyton (1947). Control mice were subjected to aerosols of PBS.

**Intranasal, intraperitoneal and subcutaneous infection of mice.** For intranasal inoculation of bacteria, the mice were lightly anaesthetized with 3% isoflurane in oxygen. The desired dose of bacteria was then delivered through one nostril of each mouse with a pipette tip in a total volume of 20 μl PBS. For intraperitoneal infection of mice, the appropriate dose of bacteria in a total volume of 100 μl PBS was injected while, for subcutaneous infection, a volume of 50 μl PBS containing the appropriate number of bacteria was injected at the left inguinal location of the mice. Control mice received PBS only.

**Determination of LD₅₀.** Ten- and 20-day LD₅₀ values were determined by the method of Reed & Muench (1938). Mice were divided into six groups of six per group. Five different doses of bacteria were used to infect the mice and the last group received PBS as a control. For aerosol experiments, the AID was used for the calculation of LD₅₀ values whereas, for intranasal, intraperitoneal and subcutaneous experiments, the concentration of bacterial suspensions used for infection of the mice were adopted for the LD₅₀ value calculation.

**Enumeration of viable *B. pseudomallei* in organs, nasal cavity and blood.** After aerosol infection with a low dose of *B. pseudomallei*, three mice were euthanized 1, 3 and 6 days post-challenge. Blood was collected aseptically in lithium heparin by cardiac puncture. The nasal cavity was washed with 1 ml sterile PBS. Lungs, spleens and livers were removed aseptically and placed in 2 or 4 ml sterile PBS. The organs were homogenized through a 100 μm nylon mesh sieve in a 6-well plate, after which they were serially diluted and plated in duplicate on Ashdown’s agar plates (Ashdown, 1979). After incubation at 37 °C for 48 h, colonies were counted from plates of appropriate dilutions and the mean colony count was derived to give the bacterial load in the different organs.

**Measurement of cytokine and chemokine production.** Lung and spleen homogenates from aerosol-infected mice were spun at 18 000 g for 5 min in a Microfuge 18 centrifuge (Beckman Coulter) before they were filtered through a 0.22 μm Millex-GV filter unit (Millipore). The homogenates were then assayed for gamma interferon (IFN-γ), tumour necrosis factor (TNF)-α, interleukin (IL)-2, IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12p70, IL-13, granulocyte–macrophage colony-stimulating factor (GM-CSF), monocyte-chemotactic protein (MCP)-1, keratinocyte-derived chemokine (KC), monocyte induced by gamma interferon (MIG), macrophage inflammatory protein (MIP)-1β and RANTES (CCL5) using the FACS Array Bioanalyzer (BD Biosciences) and CBA Flex Set System, according to the manufacturer’s instructions. Samples from lungs and spleens of uninfected mice were tested as negative controls to obtain cut-off values. The mean values (pg ml⁻¹) from three negative control mice (BALB/c/C57Bl/6) are as follows: (lungs) TNF-α, 0.21; IFN-γ, 0.1; IL-6, 0.3; KC, 2.2; RANTES, 249/263; MCP-1, 0.27; MIG, 35/19; MIP-1β, 8/9; IL-10, 39/36; (spleens) TNF-α, 41/0; IFN-γ, 4/0; IL-6, 6/0; KC, 4/0; RANTES, 4284/3728; MCP-1, 55/0; MIG, 94/91; MIP-1β, 29/23; IL-10, 60/41. The assays were done in duplicate.

**Statistical analysis.** Data were analysed using Student’s unpaired t test; P<0.05 was considered to be significant.

**RESULTS AND DISCUSSION**

**Differential susceptibility of BALB/c and C57Bl/6 mice to *B. pseudomallei* via different routes of infection**

To mimic the different ways in which humans may acquire melioidosis, both BALB/c and C57Bl/6 mice were infected
with three different strains of *B. pseudomallei* via intranasal (infection through inhalation in humans), intraperitoneal (systemic melioidosis) or subcutaneous (percutaneous inoculation) routes. Mice were monitored daily to follow their progression towards severe disease/mortality. BALB/c mice infected with any of the three strains via the intranasal route succumbed to the disease within 48 h of challenge, and represented an acute form of melioidosis, whereas those infected via the intraperitoneal route only displayed disease symptoms from day 6 post-challenge. For BALB/c mice infected subcutaneously, the onset of disease only appeared on day 23 post-infection. A similar trend in the development of disease via the intranasal, intraperitoneal and subcutaneous routes of infection was observed with the C57Bl/6 mice, with the main difference being a much delayed onset of disease. Hence, acute and chronic forms of melioidosis are established in mice infected intranasally and subcutaneously, respectively.

Table 1 summarizes the LD$_{50}$ values obtained for the different routes of infection in both strains of mice with the three strains of bacterium. The data unambiguously demonstrated that BALB/c mice were more susceptible to infection by all strains via the different portals of entry, than the C57Bl/6 mice, by a factor of 10- to 100-fold. Our results are in accordance with studies performed by various groups for the intranasal, intravenous, intraperitoneal and subcutaneous routes of infection (Barnes & Ketheesan, 2005; Hoppe et al., 1999; Liu et al., 2002). Collectively, from day 10 (intranasal and intraperitoneal) and day 20 (intraperitoneal and subcutaneous) LD$_{50}$ data, the order of the lethality of the route of infection, in decreasing order, was intranasal $>$ intraperitoneal $>$ subcutaneous. Furthermore, *B. pseudomallei* strain KHW was the most virulent of the three strains tested. Hence, variations in the genetic background of hosts and bacterial strains encountered in the environment, as well as the different portals of entry, are important factors and contribute to the varying disease outcomes.

**Table 1. Comparison of 10 day and/or 20 day LD$_{50}$ values (c.f.u.) of BALB/c and C57Bl/6 mice infected with three strains of *B. pseudomallei* via four different routes**

Both strains of mice were infected with strains KHW, EB6103 or K96243 via aerosol, intranasal, intraperitoneal or subcutaneous routes. Mice were monitored daily for progression of disease for 10 days (aerosol and intranasal) or 20 days (intraperitoneal and subcutaneous). Mortality of the infected mice was scored and LD$_{50}$ values were calculated. For aerosolization experiments, the AID was used for the calculation of LD$_{50}$ values; for intranasal, intraperitoneal and subcutaneous experiments, the concentration of bacterial suspensions used for infection of the mice were adopted for the LD$_{50}$ value calculation. Aerosolization and intranasal experiments were performed twice and similar values were obtained. ND, Not done.

Infection via aerosolization mimics closely the inhalational route in humans

Infection of mice via aerosolization was examined in order to establish an accurate murine model that more closely mimics the inhalational route in humans. *B. pseudomallei* strain KHW was chosen for this route owing to its increased virulence demonstrated above. Mice were infected with a nebulized suspension in a nose-only exposure system. The LD$_{50}$ values obtained for BALB/c and C57Bl/6 mice were $<$20 and 423 c.f.u., respectively (Table 1); C57Bl/6 mice were therefore at least 20-fold more resistant to nebulized *B. pseudomallei* than BALB/c mice. This contrasts with the results obtained by Jeddleh et al. (2003) using a whole-body aerosol exposure system in which both BALB/c (LD$_{50}$ 10 c.f.u.) and C57Bl/6 (LD$_{50}$ 27 c.f.u.) mice exhibited similar sensitivities to nebulized *B. pseudomallei* strain 1026b. The discrepancy in these results is most likely due to the utilization of different aerosol exposure systems in the two studies. In a whole-body aerosol exposure system, nebulized *B. pseudomallei* are not only inhaled through the nose, but also through the mouth, and a large proportion of the bacteria would also be trapped in body fur. Subsequently, following aerosol exposure, ingestion or inhalation of additional nebulized bacteria on the fur could occur, leading to much higher exposure doses than would otherwise be accounted for. Hence, non-pulmonary routes of infection may complicate the infection. In contrast, in a nose-only aerosol exposure system, the mice are subjected to the nebulized bacteria only during the actual duration of aerosolization. Hence, the number of bacteria delivered to the mice can be accurately determined. Bearing in mind that the bacterial strains used in the two studies are highly virulent, the addition of even a few bacteria would be enough to induce a very different clinical outcome. Furthermore, the LD$_{50}$ values obtained by Jeddleh et al. (2003) were based on the TID values, whereas in our system, the LD$_{50}$ values were based on the AID values in the mice. The TID was derived on...
Table 2. AID of viable aerosolized B. pseudomallei in the lungs of BALB/c and C57Bl/6 mice

Both strains of mice were exposed to varying concentrations of nebulized B. pseudomallei strain KHW for 30 min. Immediately after aerosol exposure, three mice from each group were euthanized. Their lungs were removed aseptically and homogenized and homogenates were plated to determine the number of viable bacteria retained in the lungs. Three independent experiments were conducted for bacterial suspension concentration $10^9$ c.f.u. ml$^{-1}$, while experiments with bacterial suspensions of concentrations of $10^6-10^8$ c.f.u. ml$^{-1}$ were performed twice independently. Results obtained from all three experiments performed under same conditions are shown here.

<table>
<thead>
<tr>
<th>Bacterial suspension (c.f.u. ml$^{-1}$)</th>
<th>Aerosol density (c.f.u. Γ$^{-1}$)</th>
<th>Mean AID (c.f.u. per pair of lungs)</th>
<th>Ratio BALB/c: C57Bl/6</th>
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<tr>
<td></td>
<td>BALB/c (individual results)</td>
<td>C57Bl/6 (individual results)</td>
<td></td>
</tr>
<tr>
<td>$10^5$</td>
<td>10$^2$</td>
<td>18 (20, 20, 14)</td>
<td>1.3 (0, 0, 4)</td>
</tr>
<tr>
<td>$10^6$</td>
<td>10$^3$</td>
<td>137 (133, 140)</td>
<td>89 (67, 110)</td>
</tr>
<tr>
<td>$10^7$</td>
<td>10$^4$</td>
<td>2212 (1753, 2670)</td>
<td>847 (633, 1060)</td>
</tr>
<tr>
<td>$10^8$</td>
<td>10$^5$</td>
<td>14289 (12577, 16000)</td>
<td>8617 (8553, 8680)</td>
</tr>
<tr>
<td>$10^9$</td>
<td>10$^6$</td>
<td>185 000 (196 000, 174 000)</td>
<td>96 092 (103 433, 88 750)</td>
</tr>
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</table>

the basis of the weight-dependent respiratory minute volume of the mouse (Guyton, 1947), but the AID represented the actual number of nebulized bacteria inhaled by the mouse and would thus be more accurate for the calculation of LD$_{50}$ values. Hence, the murine model of acute pneumonic melioidosis, established by the nose-only aerosol exposure system, more closely resembles that of human pneumonic melioidosis, rather than a whole-body exposure system (Jeddeloh et al., 2003) or the intranasal route of infection (Liu et al., 2002).

Comparing the LD$_{50}$ values obtained from the aerosol and intranasal routes of infection, there was no difference for BALB/c mice infected via either inhalational route. However, for C57Bl/6 mice, the day 10 LD$_{50}$ values were 423 and 6230 c.f.u. for the aerosol and intranasal routes of infection, respectively. These results suggest that the aerosol route of infection is more lethal than the intranasal route, and this could be explained by the fact that aerosols generated in the range of 1–5 μm can enter and be retained in the lungs, whereas droplets of bacteria delivered intranasally tend to get lodged in the nasal passage. Hence, the order of lethality for the different routes of infection, in decreasing order, was aerosol > intranasal > intraperitoneal > subcutaneous.

Bacterial colonization in B. pseudomallei aerosols-infected BALB/c and C57Bl/6 mice

To investigate other factors which could contribute to the differential susceptibility observed between the two murine hosts, the kinetics of bacterial clearance or colonization in both BALB/c and C57Bl/6 mice was determined at four different time points after aerosol challenge (immediately, day 1, day 3 and day 6). Both strains of mice were subjected to nebulized B. pseudomallei KHW with an aerosol density of 30 c.f.u. Γ$^{-1}$ for 30 min. This dose was chosen because it induced 100% mortality in BALB/c mice within 10 days, but had no apparent effect on C57Bl/6 mice. Colony counting at the earliest time point revealed a mean of 14 and 4 c.f.u. of B. pseudomallei detected in the lungs of BALB/c and C57Bl/6 mice, respectively. To exclude that this difference, observed at such an early time point, was not due to the low concentration of the bacterial suspension used, two separate experiments were performed using five different bacterial suspensions of $10^5$ to $10^9$ c.f.u. ml$^{-1}$. Both strains of mice were subjected to each aerosol density for 30 min. The results showed that a mean of 1.5- to 2-fold more viable bacteria were retained in the lungs of BALB/c mice than those of C57BL/6 mice (Table 2). The difference in bacterial numbers observed at the earliest time point was also consistent with the kinetics of bacterial growth over days 1, 3 and 6 following aerosol infection, in the lungs and nasal washes (i.e. the primary sites of infection), as well as the spleen, liver and blood (examined to determine the extent of bacterial dissemination) (Fig. 1). Collectively, BALB/c mice had overwhelming bacterial loads, which increased tremendously over the 6 day post-infection period, displaying marked organotropism for the lungs, spleens and livers. In contrast, C57 Bl/6 mice had much reduced bacterial loads, with some mice clearing the infection.

Induction of cytokines and chemokines

To characterize the early immune events which take place following aerosol challenge, both locally at the pulmonary site and systemically at distant sites, 17 cytokines and chemokines (IFN-γ, TNF-α, IL-2, IL-3, IL-4, IL-5, IL-6, IL-9, IL-10, IL-12p70, IL-13, GM-CSF, MCP-1, KC, MIG, MIP-1β and RANTES) were measured at the protein level in the lungs and spleens of BALB/c and C57Bl/6 mice following infection with aerosols of B. pseudomallei strain KHW (Fig. 2). Low levels of TNF-α, IL-6, IFN-γ, KC, RANTES, MCP-1, MIP and MIP-1β were detectable at day 1 post-infection in the lungs of C57Bl/6 mice and they increased to a modest level at either day 3 or day 6 (Fig. 2, left column). In contrast, very high levels of most of these cytokines and chemokines were found in the lungs of BALB/c mice on days 3 and day 6 (Fig. 2, left column). A modest level of IL-10 was detected in the lungs of C57Bl/6
mice on days 1 and 6, but it remained undetectable in the lungs of BALB/c mice on days 1, 3 and 6. Protein levels in spleens followed trends similar to those observed in the lungs of both strains of mice. GM-CSF and IL-12p70 were found to be at very low levels in both the lungs and spleens of BALB/c and C57Bl/6 mice, whereas IL-2, IL-3, IL-4, IL-5, IL9 and IL-13 were undetectable in both organs of both strains of mice (data not shown). Cellular sources of these cytokines have been reported: natural killer (NK) cells have been shown to be the principal source of IFN-γ, with contributions by CD8⁺ T cells through a bystander effect (Lertmemongkolchai et al., 2001); monocytes and neutrophils have been shown to be the major producers of TNF-α (Easton et al., 2007).

The increased protein levels of cytokines and chemokines in BALB/c mice paralleled the corresponding increase in their bacterial loads. No cytokine polarization was found in the lungs and spleens of either BALB/c or C57Bl/6 mice, as we were able to detect levels of proinflammatory cytokines as well as IFN-γ and chemokines. Therefore, the susceptibility of BALB/c mice to B. pseudomallei infection was not due to an absent or inadequate inflammatory response. Similarly, this observation has been reported by other groups using either the intranasal, intraperitoneal or intravenous models (Barnes et al., 2001; Liu et al., 2002; Ulett et al., 2000). However, data from these groups consisted mainly of mRNA responses, and not actual protein expression levels (Barnes et al., 2001; Ulett et al.,...
Since an increase in the transcriptional activity of cytokine and chemokine genes does not necessarily translate into release of functional protein products, our study is the first report of a comprehensive investigation of the cytokines and chemokines induced at the protein level for the nose-only aerosol infection model, which, in itself, is a better model of acute pneumonic melioidosis in humans than the intranasal model.

Both local and systemic production of IFN-γ were higher in BALB/c mice than in C57Bl/6 mice, suggesting that hyperproduction of IFN-γ is not protective and may lead to septic shock (Car et al., 1994; Heremans et al., 1990). Studies in patients with melioidosis have also demonstrated a link between severity of disease and the levels of cytokines and chemokines in serum, including IFN-γ, TNF-α, IL-6 and IL-8 (Brown et al., 1991; Friedland et al., 1992; Lauw et al., 1999).

Fig. 2. Induction of cytokines and chemokines in lungs and spleens of aerosol-infected BALB/c and C57Bl/6 mice. Mice were infected with KHW aerosols of 30 c.f.u. l⁻¹ for 30 min. At days 1, 3 and 6 post-infection, lungs and spleens from three mice per group were excised and homogenized. The lung (left column) and spleen (right column) homogenates of BALB/c (open bars) and C57Bl/6 (filled bars) mice were assayed for the presence of cytokines and chemokines (note the different scales in the two columns). Data shown are mean values with error bars showing the SEM from three mice at each time point. Samples from uninfected mice were tested as negative controls to obtain cut-off values. The assays were done in duplicate. Two independent experiments were performed and similar values were obtained. Results from one experiment are shown here. *, P<0.05; **, P<0.01.
Hence, the early production of local proinflammatory cytokines observed in BALB/c mice appears to spill over into the systemic compartment, and thus contribute to the immunopathogenesis of the disease (Liu et al., 2002). In contrast, C57Bl/6 mice responded to the infection with a moderate and transient induction of proinflammatory response, and they were able to control and clear the infection. Although too much of a good thing, i.e. IFN-γ in this case, is not ideal, the obligatory role of this cytokine for the survival of infected mice is undisputed (Santamirand et al., 1999). Our observation that IL-10 was only found in C57Bl/6 mice, and not BALB/c mice, is in contrast to those reported by Koo & Gan (2006) who observed higher IL-10 production by BALB/c splenocytes than C57Bl/6 splenocytes to live B. pseudomallei infection in vitro. The discrepancy in IL-10 production could be explained by the different systems used in the model used by these authors (in vitro) and our model (in vivo), the latter being a closer and more accurate representation of human melioidosis. The lack, or inadequate production, of anti-inflammatory responses in our BALB/c mouse model could also contribute to the development of acute infection, unlike in the C57Bl/6 mouse model, where the subclinical infection is correlated with a more balanced and modest production of pro- and anti-inflammatory cytokines. It is noteworthy that greater infiltration and accumulation of neutrophils were observed in organs of BALB/c mice, whereas early lesions in C57Bl/6 mice were predominantly comprised of macrophages (Barnes et al., 2001). This observation correlates with the acute inflammation seen in BALB/c mice and the better control of infection, possibly by macrophages, in C57Bl/6 mice (Gan, 2005). However, data from Easton et al., (2007) demonstrated that neutrophils are critical to the generation of the early proinflammatory cytokine environment in the lungs, and suggested an important cytokine-mediated immunomodulatory role for neutrophils in B. pseudomallei infection.

Because the development of the acute form of this disease is not due to a lack, but rather, an excess of inflammation, our nose-only aerosol model, together with the simultaneous detection of a comprehensive list of cytokine and chemokine responses at the protein level, will be an extremely useful platform, critical for future studies to investigate host interactions with bacterial pathogens and will assist in the development of better and/or novel therapeutic options for melioidosis patients.

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