Histopathological and immunohistochemical characterization of Burkholderia pseudomallei lesions in an acute model of infection with BALB/c mice

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SUMMARY

Organ tissue damage is a key contributor to host morbidity and mortality following infection with microbial agents. Severe immune responses, excessive cellular recruitment and necrosis of cells all play a role in disease pathology. Understanding the pathogenesis of disease can aid in identifying potential new therapeutic targets or simply act as a diagnostic tool. Burkholderia pseudomallei is a Gram-negative bacterium that can cause acute and chronic diseases. The BALB/c mouse has been shown to be highly susceptible to aerosol challenge with B. pseudomallei and hence acts as a good model to study the acute and potentially lethal form of the disease melioidosis. In our study, BALB/c mice were challenged and culled at predetermined time points to generate a pathological time course of infection. Lung, liver and spleen were subjected to pathological and immunohistochemical analysis. The number and type of microscopic lesions within each organ, as well as the location and the mean percentage of neutrophils, B cells, T cells and Burkholderia capsule antigen within the lesions, were all characterized during the time course. Neutrophils were determined as the key player in tissue pathology and generation of lesions, with B cells playing an insignificant role. This detailed pathological assessment increases our understanding of B. pseudomallei disease.

Keywords
Burkholderia pseudomallei, immunohistochemistry, immunopathology, neutrophils

Burkholderia pseudomallei is the causative agent of the disease melioidosis. It is an intracellular Gram-negative bacterium that has the ability to infect numerous different cell types (David et al. 2015) and is naturally resistant to many antibiotics, due to efflux pumps (Mima & Schweizer 2010; Rhodes & Schweizer 2016).

Melioidosis is an emerging tropical disease, endemic to South-East Asia and Northern Australia (Aldhous 2005; Kaestli et al. 2009; Valade et al. 2009). Burkholderia pseudomallei is known to cause diseases in humans via the aerosol route and via cutaneous lesions (Leelarasamee 1998). Depending on challenge dose and route of infection, it can either cause an acute disease where the immune response is inadequate and mortality rates are high or a chronic infection where the pathogen can reside for long times in localized regions (Wiersinga et al. 2006, 2009).

The natural ability of B. pseudomallei to infect via the aerosol route and its low challenge dose have also made this bacterium a potential biothreat agent of concern (Gilad et al. 2007). Currently, there is no licensed vaccine to prevent infection, and antibiotic treatment is suboptimal and often time-consuming (Dance 2014). Single antibiotic dosing regimens are only partially effective, and combination treatments have to be given for prolonged periods, up to 6 months to clear the infection completely (Srinivasan et al. 2001). Therefore, there is a continued requirement to develop new therapies and thus a greater understanding of the disease.

Several animal models have been developed over the years to determine the host–pathogen interactions. The mouse model has been the most characterized with the BALB/c mouse being very susceptible to infection and the C57BL/6

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mouse strain being more resistant (Barnes et al. 2001). This has led to the BALB/c mouse being mainly used for acute disease studies and the C57BL/6 strain for chronic models (Leakey et al. 1998; Hoppe et al. 1999; Liu et al. 2002; Barnes & Ketheesan 2005; Conejero et al. 2011; Bearss et al. 2017).

Despite growing research into this pathogen, there are still some gaps in the knowledge of the organ pathology and lesion development associated with the disease, although it has been documented that excessive immune response postinfection with B. pseudomallei is a significant contributor to host mortality (Ulett et al. 2000; Tan et al. 2008). Therefore, controlling tissue pathology and organ damage caused by an overactive immune response could represent an appropriate therapeutic strategy where an immunomodulatory drug may control the excessive inflammation and an antibiotic can control the bacterial spread (Laws et al. 2015). Preliminary histological studies have demonstrated in the BALB/c mouse that there are foci of acute inflammation and necrosis in the lung, liver and spleen following infection with B. pseudomallei K96243 strain (Barnes et al. 2001; Lever et al. 2009). Additional histological studies have focused on the chronic C57BL/6 mouse model of infection with samples taken at day 20 and day 60 p.i., indicating that the spleen and liver are the most susceptible organs to bacterial growth and tissue destruction (Conejero et al. 2011).

In this study, we sought to carry out a detailed histopathological and immunohistochemical analysis using the BALB/c acute model and the B. pseudomallei K96243 human clinical isolate strain. Analyses were carried out at multiple time points during the short course of infection.

Material and methods

Experimental design

Forty male BALB/c mice (Charles River, UK), six to eight week old, were included in this study, split over two identical but independent experiments. Mice were randomly assigned into cages and transferred to a high-containment Class III rigid isolator, where they were given unlimited access to food and water. Thirty mice were challenged with the B. pseudomallei strain K96243 by aerosol, using a Henderson-type apparatus (Druett 1969) and a Collison nebulizer (May & Harper 1957), and ten were left as naive controls. Bacteria were grown in Luria broth at 37°C on a rotary platform. Animals received a lethal dose ~ 20 × LD50 with an average challenge dose of 146 CFU. The challenge dose is the average calculated retained dose per mouse. Mice were sprayed for a total of 10 minutes with a 2-minute subsamples (impinger) collected for bacterial enumeration. The impinger sample was serially diluted and plated in triplicates.

At 24 hpi, infected mice showed minor ruffling and hunching. At 48 hpi, ruffling and hunching were more pronounced and signs of conjunctivitis appeared. At 60 hpi, infected mice were either at or close to humane end point. Mice were culled at predetermined time points (TP1 = 24 hpi, TP2 = 48 hpi and TP3 = 60 hpi). Ten infected mice were culled at each time point while five naïve controls were culled at 24 hpi and five at 60 hpi.

Histopathology

The lung, liver and spleen were all processed within less than 1 hour post mortem and immersed in neutral sodium salt-buffered formalin. Samples were fixed for 1 month and embedded in paraffin wax, and 4-μm sections were cut, dewaxed and rehydrated through xylene and alcohols to be finally washed in running tap water. Haematoxylin and eosin (H&E) stain was used for the histopathological analysis.

Immunohistochemistry

The immunohistochemical detection of B. pseudomallei was performed with a murine monoclonal antibody (3VIE5) that reacts with a capsular polysaccharide of B. pseudomallei. For the characterization of cell composition of lesions, a rat monoclonal anti-Ly-6G (Invitrogen, Darmstadt, Germany), a rat monoclonal anti-CD45R-B220 (Invitrogen) and a rabbit polyclonal anti-CD3 (Dako) were used as immunomarkers for neutrophils, B cells and T cells respectively.

Consecutive 4-μm sections were dewaxed, rehydrated and placed in a fresh solution of 3% hydrogen peroxide in methanol for 15 minutes to block endogenous peroxidase activity. Samples were then washed for 10 minutes in tap water before being clipped into Shandon Sequenza clips (Thermo Scientific, Loughborough, Leistershire, UK). After slides were mounted, epitope demasking was performed through enzymatic digestion, using a solution of 2% proteinase K (Dako) in Tris-buffered saline (0.05 mol/L Tris–HCl pH 7.5 to 7.7). Then, the tissue sections were washed for 10 minutes with Tris-buffered saline (TBS) (0.85% NaCl, 0.0605% Tris, adjusted to pH 7.5 using 1 M HCl), and then, 190 μl of Universal BlockerTM Blocking Buffer in TBS (Thermo Scientific) was added to the slides as blocking agent. After twenty minutes, 190 μl of the primary antibody was added. The incubation time as well as the concentration varied for each primary antibody used (Table 1). The Vector M.O.M. Immunodetection kit (Vector, Peterborough, UK) was used for the anti-Burkholderia antibody following the manufacturers’ indications. For anti-Ly-G6, anti-CD45R-B220 and anti-CD3 antibodies, after two washes with TBS buffer, 190 μl of biotinylated link antibody and link block (Thermo Scientific) was added (Table 1), followed by two further buffer washes, 30 minutes later. Primary and secondary antibody binding was amplified using Ultra-Sensitive ABC Peroxidase Rabbit IgG Staining Kit (Thermo Scientific) and visualized using the Vector® NovaRED™ Substrate Kit (Vector Laboratories, Burlingame, CA, USA). Unbound conjugate was removed prior to chromogen application with two buffer washes. Slides were then washed in purified water, removed from the coverplates and placed in a rack. Samples were rinsed with tap water for five minutes, before being placed in Mayer’s Haematoxylin counterstain,
followed by a further wash in tap water. Finally, sections were dehydrated, cleared and mounted for analysis.

**Image analysis and statistics**

Immunolabelled sections were analysed using light microscopy and digital image analysis (Nikon NIS-Br, Nikon, Japan).

The area of tissue with lesions was calculated using a full section of lung, liver and spleen. The slides were examined with the ×40 objective to give a final magnification of ×400, to ascertain the percentage of immunostained cells within the lesion.

To compare the mean number and type of lesions per time point, nonparametric Mann–Whitney’s U-test was used. Data were considered statistically significant when \( P < 0.05 \).

For immunohistochemistry, the mean expression of anti-CD3 antibody was compared for type of lesions and time points. Differences were considered significant at \( P < 0.05 \).

The results of immunohistochemical analysis are expressed as mean ± standard deviation (S.D).

Due to the staining characteristics or the localization of the immunolabelled cells, it was not possible to perform the image analysis for the antibodies anti-*Burkholderia*, Ly-6G and CD45R-B220. For this reason, the analysis of these antibodies has been made in a qualitative way.

All the statistical analyses were conducted using SPSS 19 software package (SPSS Inc., Chicago, IL, 60606, USA) and GraphPad Prism 4.0 (San Diego, CA, USA).

**Ethical approval**

All procedures involving animals were conducted under project licences approved by internal ethical review and the Home Office, and in accordance with both the Animal (Scientific Procedures) Act (1986) and the 1989 Codes of Practice for the Housing and Care of Animals used in Scientific Procedures.

**Results**

**Time course histopathology in the lung**

Lung lesions were classified into three categories according to cellular composition and evolution stage: type I, characterized by a discrete inflammatory foci of macrophages, lymphocytes and some neutrophils, usually close to blood vessels and/or bronchioles (Figure 1a); type II, defined as round lesions with areas of granulomatous pneumonia with similar distribution to those in type 1, but containing epithelioid macrophages and more neutrophils, some of which were necrotic (Figure 1b); and type III defined as large pyogranulomas mainly formed by neutrophils and with a necrotic centre surrounded by epithelioid macrophages, some lymphocytes and plasma cells (Figure 1c). The total number of lesions of each type was counted in a full lung section per animal.

After 24 hpi, type II lesions were predominant in the lung followed by type I lesions. Type III lesions were exclusively observed in one animal at 24 hpi, while other animal did not show lesions at this time point. At 48 and 60 hpi, there was an increase in the total number of lesions, especially due to a statistically significant increase in type III lesions. Finally, at the last time point (60 hpi), the total number of lesions as well as the lesion types remained stable compared with 48 hpi (Figure 2).

**Time course histopathology in the liver**

Lesions in the liver were classified, similar to lung, into three categories according to necrosis extension. Type I was characterized by a minimal foci of necrosis mainly due to infiltrated neutrophils (Figure 1d); type II was defined as moderate foci of necrosis formed by mixed inflammatory cells (Figure 1e); and type III was characterized by more extensive foci of necrosis frequently due to coalescent lesions (Figure 1f).

The total number of lesions of each category was counted in a full liver section per animal. Only two animals showed type I lesion in the liver at 24 hpi. However, there was an exponential increase in the number of lesions at 48 hpi, with type I lesions being the most observed. There was an increase in the number of all types of lesions at 60 hpi when compared to 24 and 48 hpi (Figure 3).

**Time course pathology in the spleen**

The spleen showed a marked disruption of the normal tissue architecture. Lesions were typically pyogranulomatous, with...
abundant neutrophils and macrophages, mainly within the red pulp but also in the white pulp. No differentiation into types was performed for the splenic lesions. The percentage of tissue damaged in a full spleen section per animal was estimated to calculate the organ pathology score.

Only one animal showed a single necrotic focus at 24 hpi. There was a statistically significant increase in the percentage of affected tissue at 48 hpi and 60 hpi when compared to 24 hpi with all animals showing lesions at this time point. At 60 hpi, also there was an increase in the percentage of affected tissue compared with 48 hpi, which was not statistically significant (Figure 4).

Figure 1 (a) H&E lung section from a type I lesion (200×). Inflammatory foci of macrophages, lymphocytes and some neutrophils close to a blood vessel (*). (b) H&E lung section from a type II lesion (100×). Foci of granulomatous pneumonia containing some epithelioid macrophages and neutrophils, some of which were necrotic (arrow). (c) H&E lung section from a type III lesion (100×). Large pyogranuloma adjacent to a bronchiole (*) formed by neutrophils with a necrotic centre (arrow) surrounded by few macrophages, lymphocytes and plasma cells. (*). (d) H&E liver section from a type I lesion (200×). Minimal foci of necrosis mainly due to infiltrated neutrophils (arrow). (e) H&E liver section from a type II lesion (200×). Moderate foci of necrosis formed by mixed inflammatory cells (arrow). (f) H&E liver section from a type III lesion (200×). Extensive foci of necrosis due to coalescent lesions (*).

Figure 2 Time course histopathology in the lung. Total number of each type of lesions (I, II and III) observed per time point. At 48 and 60 hpi (TP2 and TP3), there was a statistically significant increase (P < 0.05) in the total number of type III lesions (*) when compared to 24 hpi (TP1).

Figure 3 Time course histopathology in the liver. Total number of each type of lesions (I, II and III) observed per time point. An exponential increase in the number of lesions was observed at 48 hpi when compared to 24 hpi. At 60 hpi, there was an increase in the number of lesions when compared to 24 and 48 hpi, significantly higher for type I and II lesions (*P < 0.05).
Immunohistochemical analysis of Burkholderia spp. antigen and cell populations within lesions

Anti-Burkholderia spp. antibody. Immunohistochemistry against Burkholderia spp. capsular antigen showed extensive positive staining within lung, liver and spleen lesions. The positive reaction was located mainly within the cytoplasm of macrophages in type I lung lesions at 24 hpi (Figure 5a). However, staining in the lung became more extensive over time as infection progressed and antigen was located within the cytoplasm of both macrophages and neutrophils associated with areas of consolidation and necrosis in type II and III lesions (Figure 5b).

Positive reaction was also observed within both degenerated and viable macrophages and neutrophils surrounding the areas of necrosis in liver and spleen pyogranulomas (Figure 5c, d). No Burkholderia spp. capsular antigen was detected in uninfected controls.

Neutrophils (Ly-6G). High numbers of Ly-6G+ cells were detected in all lesion types in the lung. Positive cells were mainly observed within the lesion but also in the adjacent tissue (Figure 6a). Fewer positive cells were observed within the centre of the lesions, especially in the larger ones (Figure 6b). Fewer Ly-6G+ cells were detected in hepatic and splenic lesions when compared to the lungs.

T lymphocytes (CD3). Few CD3+ cells were sparsely distributed within lesions independent of the lesion type. A small number of CD3+ cells were observed in lung lesions (Figure 6c, d). However, a statistically significant higher number of CD3+ cells were observed in type I and II lesions at 24 hpi when compared to 48 hpi and 60 hpi (Figure 7).

B lymphocytes (CD45R-B220). B lymphocytes do not seem to be involved in the development of B. pseudomallei lesions

Figure 4 Time course pathology in the spleen. Percentage of spleen area affected per time point. A statistically significant ($P < 0.05$) increase in the percentage of affected tissue at 48 and 60 hpi when compared to 24 hpi was observed (*).

Figure 5 Immunohistochemically stained sections against a capsular polysaccharide of Burkholderia pseudomallei. (a) Type I lung lesions (arrow) at 24 hpi (100×) with positive reaction mainly within the cytoplasm of macrophages and neutrophils. (b) Extensive type III lesion at 60 hpi (100×) showing the antigen within the cytoplasm of macrophages and neutrophils associated with areas of consolidation and necrosis (*). (c) Positive immunohistochemical reaction within both degenerated and viable macrophages and neutrophils surrounding a focus of necrosis in a type II lesion (200×) in the liver (arrow) and a small type I lesion (arrowhead). (d) Positive immunohistochemical reaction within the spleen at 60 hpi, mostly affecting the red pulp (arrows) (100×).
in our BALB/c model, as they were not observed in any type of lesions independently of the time point analysed. The number of B cells present in the adjacent healthy tissue was also small (Figure 6e), but occasionally they formed aggregates (Figure 6f). B cells were not observed in the hepatic lesions.

**Discussion**

The organ pathology associated with *B. pseudomallei* infection is still not clearly understood. The BALB/c mouse model has been proposed as an appropriate model to evaluate the acute form of melioidosis in humans. However, the histopathology and immunopathology associated with this disease in BALB/c mice has been analysed infrequently...
Our study shows a detailed histopathological and immunohistochemical characterization of lung, liver and spleen pathologies using the BALB/c mice acute infection model and the B. pseudomallei K96243 human clinical isolate strain.

In our study, we found that 9/10 animals showed pyogranulomatous lesions within the lung as early as 24 hpi. These results are in line with previous studies demonstrating the high susceptibility of this mouse strain and how quickly the disease can evolve to the development of severe lesion types even at the first 24 hrs, as we found one animal showing type III lesions at this time point (TP1).

Previous studies described an exponential bacterial multiplication within the lung over the first 24 hpi, before the rate of replication became more gradual (Jeddeloh et al. 2003; Lever et al. 2009). This disease progression pattern is reflected within the microscopic lesions observed in our study, with an exponential increase in the number of lesions at 48 hpi (TP2), being later more stable at 60 hpi (TP3).

Similar patterns of initial exponential growth have been described in the liver and spleen once the bacteria are seeded from the lung. At this point, animals died because the damage in these organs became more severe while the rate of microorganism replication remained steady (Lever et al. 2009; Conejero et al. 2011).

In the study from Lever et al. (2009), no hepatic lesions were detected at 24 hpi and only two small foci of necrosis were shown at 48 hpi in the liver. In the same study, the lesions in the spleen described as multifocal acute necrotizing splenitis were not shown until 48 and 72 hpi. In contrast, in our study, we found initial lesions in the liver of two animals as early as 24 hpi (TP1), including the one animal showing type III lesions in the lung and splenic lesions at this time point. Similar changes in the liver and spleen were observed by Barnes et al. (2001), with an early neutrophil influx to the organs causing small inflammatory foci at 24 hpi. Furthermore, we observed an exponential increase in the number of lesions in both liver and spleen, at 48 hpi which continued at 60 hpi although the challenge dose used in the Lever et al. (2009) study (100 cfu) was similar to the one used in our study.

Intense immunostaining for the capsular polysaccharide antigen of B. pseudomallei was observed within lesions in all tissues. The extension and intensity of the staining were higher in large lesions, which could reflect the sustained bacterial growth at these sites.

After analysing the cell composition of microscopic lesions by immunohistochemistry, we have found neutrophils (Ly-6G+ cells) as the most frequent cells. In a recent comparative virulence study (Massey et al. 2014), an important increase (around 300%) in the number of neutrophils in peripheral blood was observed in infected animals, Furthermore, previous studies have demonstrated that activated neutrophils are rapidly recruited to the lungs after a high-dose pulmonary challenge with B. pseudomallei using flow cytometry (Easton et al. 2007) or histopathology (Barnes et al. 2001; Kager et al. 2014). This has also been described in another study highlighting the presence of macrophages and natural killer cells in the lesions (Bears et al. 2017). In line with this finding, histological analysis of tissue from human patients with melioidosis has demonstrated the presence of abundant neutrophils together with a small number of macrophages within lesions (Wong et al. 1995).

It has been suggested that neutrophils play a role in early resistance against B. pseudomallei infection (Breitbach et al. 2006). However, our findings are more in agreement with those of the authors who have suggested that neutrophils are not sufficient to control the infection despite their recruitment to the infection site (Massey et al. 2014). In addition, Massey et al. (2014) found an important increase in IL-17 in infected animals compared with non-infected control animals. This cytokine is recognized as the main mechanism whereby Th17 cells contribute to tissue damage due to prolonged neutrophil recruitment (Fogli et al. 2013). This mechanism may also be involved in the development of microscopic lesions observed in our study.

There is growing literature supporting the view that an excessive immune response, leading to exaggerated cellular recruitment, is a major contributor to morbidity and ultimately mortality in a number of different infections (D’elia et al. 2013). Furthermore, it has been shown that dampening down the immune response during Burkholderia spp., infection can increase survival. Inhibition of IL-1beta, COX-2 and HMGB1 has been shown to be beneficial in models of melioidosis (Ceballos-Olvera et al. 2011; Asakrah et al. 2013; Charoensup et al. 2014; Laws et al. 2015; ). It has also been demonstrated that robust T-cell functions are paramount in protection against B. pseudomallei infection and that IFN-g and the IFN-g-inducing cytokines IL-12 and IL-18 are essential for initial resistance (Haque et al. 2006). In our study, the number of CD3 + cells within the lesions was quite low. We found a statistically significant higher presence of CD3 + cells within early lesions at 24 hpi, which could represent a first attempt to control the disease. However, the presence of CD3 + cells was minimal at 48 hpi and 60 hpi, where lesions were mainly formed by neutrophils and some macrophages. This could be one of the reasons why BALB/c mice fail to control the disease and die in <5 days after infection (Lever et al. 2009). Indeed, the importance of T cells later during infection has been shown in a B. mallei model (Rowland et al. 2010).

A recent study showed that strong CD4 + and CD8 + T-cell responses are required for patients to survive acute melioidosis (Jenjaroen et al. 2015). CD4 + T cells were also reported to protect mouse in the late stage of B. pseudomallei infection (Haque et al. 2006). We found a slight increase (not statistically significant) in the presence of CD3 + cells
in type I lesions at TP3 that would not seem to be enough to establish a late response.

B lymphocytes were not observed in any type of lesions in our study independent of the time point analysed, although occasional aggregates of B cells were observed within healthy lung tissue adjacent to lesions. This may be due to the fact that the lesional pattern is too acute for the intervention of this cell population.

This finding has been described in other diseases caused by intracellular pathogens such as Mycobacterium bovis (Salguero et al. 2016). However, the role of these ‘nests’ in the host immune response is not well understood.

The histopathological and immunohistochemical analysis of the lesions found in this BALB/c mice model demonstrates the high susceptibility of this mouse strain to acute melioidosis. This is corroborated by the highest presence of Burkholderia pseudomallei antigen within the largest lesions, which are deficient in a T-cell response and dominated by the presence of neutrophils. The high recruitment of neutrophils seems to be responsible for most of the tissue damage observed within histopathological lesions.

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Conflict of interest

Authors do not declare any conflict of interest.

Author contributions

WLG, FJS and RVD equally contributed to the writing of the manuscript. RVD was responsible for the experimental design and performing the animal studies. WLG and FJS were responsible for the histopathological and immunohistochemical characterization.

References


