

Immunogenic recombinant *Burkholderia pseudomallei* MprA serine protease elicits protective immunity in mice

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Burkholderia pseudomallei is resistant to a diverse group of antimicrobials including third generation cephalosporins whilst guinolones and aminoglycosides have no reliable effect. As therapeutic options are limited, development of more effective forms of immunotherapy is vital to avoid a fatal outcome. In an earlier study, we reported on the B. pseudomallei serine MprA protease, which is relatively stable over a wide pH and temperature range and digests physiological proteins. The present study was carried out to evaluate the immunogenicity and protective efficacy of the MprA as a potential vaccine candidate. In BALB/c mice immunized with recombinant MprA protease (smBpF4), a significantly high IgG titer was detectable. Isotyping studies revealed that the smBpF4-specific antibodies produced were predominantly IgG1, proposing that immunization with smBpF4 triggered a Th2 immune response. Mice were immunized with smBpF4 and subsequently challenged with B. pseudomallei via the intraperitoneal route. Whilst control mice succumbed to the infection by day 9, smBpF4-immunized mice were protected against the lethal challenge and survived beyond 25 days post-infection. In conclusion, MprA is immunogenic in melioidosis patients whilst also eliciting a strong immune response upon bacterial challenge in mice and presents itself as a potential vaccine candidate for the treatment of melioidosis.

Keywords: melioidosis, serine protease, immunotherapy

INTRODUCTION

Burkholderia pseudomallei is the Gram negative etiological agent for melioidosis, a potentially acute fulminating disease in humans and animals in Southeast Asia and Northern Australia (Cheng and Currie, 2005; Peacock, 2006). In areas where this bacterium is endemic, infection by B. pseudomallei has been estimated to be responsible for 20-30% mortality due to septicaemia and 40% of sepsis-related mortality (Wiersinga et al., 2006). Human melioidosis presents with an array of clinical symptoms ranging from acute or chronic localized infection involving a single organ, to fulminant septicaemia in multiple organs and septic shock (Cheng and Currie, 2005). Current management of melioidosis involves the full 20-week duration of third generation cephalosporins therapy. Unfortunately, treating a B. pseudoma*llei* infection is difficult as the bacterium is intrinsically resistant to many antibiotics and relapse following apparently successful therapy is well recognized (Cheng and Currie, 2005).

Production of an effective vaccine against *B. pseudomallei* is challenging due to the multi-factorial nature of this pathogen. Various vaccination strategies, including the use of attenuated stains of *B. pseudomallei* (Haque et al., 2006), heat-inactivated bacteria (Sarkar-Tyson et al., 2009), subunit vaccines (Harland et al., 2007), and DNA vaccines (Chen et al., 2006) have been extensively explored, but none were able to confer complete protection against the infection. Currently, the use of

protection-eliciting pathogen proteins, for example virulence factors, is looked upon as an alternative and viable approach as previously demonstrated with the caseinolytic protease of *Streptococcus pneumoniae* (Cao et al., 2009), parasite cysteine proteases (Jorgensen and Buchmann, 2011), cytotoxic serine protease of *Vibrio harveyi* (Cheng et al., 2010), outer membrane proteins of *Haemophilus parasuis* (Yuan et al., 2011), and outer membrane secretin PilQ of *Neisseria meningitidis* (Haghi et al., 2012).

A common group of virulence factors shared among bacterial pathogens is protease, and protease inhibitors have proven effective therapeutic agents in treating infectious diseases in vertebrates (Ribeiro-Guimaraes et al., 2007; Serrano-Luna et al., 2007). For example, in the pathogenic bacterium Enterococcus faecalis, both gelatinase and serine protease are required for systemic infections in mammalian hosts (Engelbert et al., 2004). Similarly, the extracellular alkaline serine protease produced by the nosocomial pathogen Stenotrophomonas maltophilia is an important pathogenicity factor and has been recognized as a potential therapeutic target (Windhorst et al., 2002). The primary function of proteases in the bacterial kingdom is to provide a source of free amino acids for bacterial survival and growth, but there is increasing evidence of the role of proteases in bacterial pathogenesis in creating a niche for colonisation, suppressing host defence mechanisms and supporting host dissemination. Furthermore, the involvement of proteases in antibody degradation or modulation

of cellular immune responses is also proposed (Jorgensen and Buchmann, 2011).

Many reports have alluded to the potential role of proteases in B. pseudomallei pathogenicity. The serine MprA protease has been implicated as a possible virulence factor (Lee and Liu, 2000) causing extensive damage to mammalian physiological proteins involved in circumventing the detrimental effects of bacterial secreted proteases (Chin et al., 2007). Previously, we identified an epitope of the *B. pseudomallei* protease by peptide biopanning against anti-B. pseudomallei antibodies, whereby selected peptides displayed a conserved motif of serine-methionine-alanine (S-M-A). This S-M-A motif mimics the catalytic serine and its flanking amino acids of the B. pseudomallei mprA gene (Chan and Nathan, 2005). As a potential antigen involved in B. pseudomallei pathogenesis, we further characterized the MprA protein and demonstrated that the expressed recombinant MprA protease maintained superior proteolytic activity over a wide pH (pH5-11) and temperature (4-68°C) range and partially digested immunoglobulins A and G, transferrin and myosin (Chin et al., 2007).

As the MprA protein carried an antibody-binding epitope and was implicated in proteolysis, we evaluated the potential of *B. pseudomallei* MprA protease as a vaccine candidate in a mouse model. We observed a significant delay in death of the immunized challenged animals indicating that this protease elicited a protective immune response in mice.

MATERIALS AND METHODS

ETHICS STATEMENT

All animal experiments were performed in accordance with the Universiti Kebangsaan Malaysia animal ethics guideline and approved by the Universiti Kebangsaan Malaysia Animal Ethics Committee (UKMAEC). BALB/c mice (6 weeks old, female) were obtained from the Animal House Facility, Universiti Kebangsaan Malaysia. Mice were maintained under specific-pathogen-free conditions, subjected to a 12 h light/dark cycle and were fed on a diet of commercial pellets and distilled water *ad libitum*.

BACTERIAL STRAINS AND PLASMID

B. pseudomallei human isolate (strain D286) was obtained from the Pathogen Laboratory, School of Biosciences and Biotechnology, Faculty of Science and Technology, Universiti Kebangsaan Malaysia. This bacterial strain was isolated from a patient with clinical manifestation of melioidosis at the Kuala Lumpur Hospital and previously characterized based on biochemical tests as well as by 16S rRNA sequencing (Lee et al., 2007). The recombinant MprA protease (BPSS1933) smBpF4 construct has been previously described (Chin et al., 2007).

BACTERIAL-CELL FREE CULTURE FILTRATE—*C. elegans* SURVIVAL ASSAY

B. pseudomallei bacterial cell-free culture filtrate was prepared and used to treat *Caenorhabditis elegans* to evaluate nematode killing. The assay and preparation of the cell-free filtrate was performed as previously described (Lee et al., 2011) with minor modifications. In brief, one *B. pseudomallei* colony was inoculated into 10 ml Brain Heart Infusion (BHI) broth and incubated overnight

at 37°C. Phenylmethylsulfonyl fluoride (PMSF, 10 mM) and ethylenediaminetetraacetic acid (EDTA, 10 mM) were added to the overnight bacterial culture and further incubated at 37°C for 30 min. Subsequently, the mixture was centrifuged at 4000 g for 15 min at 4°C and the supernatant collected and filtered through a 0.22 μ m pore size syringe membrane filter unit (Millipore, USA). The supernatant was supplemented with 1 ml S-basal medium. The assay was performed in six-well microtiter plates containing 2 ml of the supernatant and S-basal mixture. Thirty age-matched *C. elegans* Glp worms (Lee et al., 2011) were transferred into the conditioned liquid medium. Heat-killed *Escherichia coli* OP50 were supplied to prevent starvation. Mortality of the worms was monitored at 4-h intervals. *E. coli* OP50 was used in place of *B. pseudomallei* as the negative control.

HUMAN SERA SAMPLES

Culture-confirmed melioidosis patients sera (n = 5) from Malaysia presenting with acute and chronic disease manifestations and sera samples of patients presenting with Legionella infection (n = 5) but confirmed *B. pseudomallei*-culture negative were provided by the Institute for Medical Research, Malaysia. Healthy (confirmed HIV-negative) human sera samples (n = 5)were generously provided by Clinipath Malaysia Sdn. Bhd.

OVER-EXPRESSION AND PURIFICATION OF MprA PROTEASE, smBpF4

Over-expression and purification of recombinant MprA protease smBpF4 were performed as previously described (Chin et al., 2007).

REACTIVITY OF RECOMBINANT PROTEINS TOWARD MELIOIDOSIS PATIENTS' SERA

The 96-well microplates were coated with 0.3 μ g per well of either heat-killed (80°C, 1 h) and sonicated (15 min) (Vibracell, Sonics and Materials, USA) *B. pseudomallei* or recombinant smBpF4 purified protein in coating buffer (0.1 M sodium bicarbonate, p.H 8.6) and incubated overnight at 4°C. The plates were blocked, washed, then incubated with human sera samples (1:200 dilution) and incubated at 37°C for 1 h as previously described (Chin et al., 2007; Hara et al., 2009). Rabbit anti-human IgG conjugated with peroxidase (1:10,000 dilution, Sigma-Aldrich, USA) was added and incubation continued for 1 h at 37°C. At the end of the incubation, the plates were washed and 50 µl ABTS:peroxidase B (1:1; KPL, USA) was added as substrate. The absorbance was read at 405 nm on an ELISA reader (Dynex MRX, USA).

MOUSE IMMUNIZATION

BALB/c mice (n = 4, 6 weeks old, female) were immunized via three doses of 100 µg of heat-treated (80°C, 10 min) purified recombinant smBpF4 protein, via the intra-peritoneal (i.p.) route over 9 weeks as previously described (Hara et al., 2009). The first dose was administrated in Freund's complete adjuvant (Pierce Biotechnologies, USA) on day 0 and 2, boosts were administrated on days 14 and 28 with an identical dose of antigen in Freund's incomplete adjuvant (Pierce Biotechnologies, USA) (Harlow and Lane, 1988). Control mice (n = 3) were injected with Phosphate Buffered Saline (PBS) plus adjuvant as per the above schedule. The sera samples were collected and recovered prior to immunization and 1 week after the first and second booster according to the procedure of Harlow and Lane (1988) and antibody titres evaluated before the animals were challenged with *B. pseudomallei*. The resulting sera were subsequently used for antibody response analyses and protease neutralization assays by zymography

ANALYSIS OF ANTIBODY RESPONSE

Mouse antibody titers against recombinant proteins were measured by ELISA as previously described (Chin et al., 2007; Hara et al., 2009) with some modifications. Briefly, 96-well plates were coated overnight with 1 μ g of the purified recombinant protein used for immunization. Wells were blocked with 5% skimmed milk to prevent nonspecific binding. The plates were then washed thoroughly with distilled water. Serum samples were serially diluted two-fold (1:100 to 1:25,600 dilutions) across the microtiter plate in 5% skimmed milk and incubated for 1 h at 37°C. Rabbit anti-mouse IgG conjugated with peroxidase (1:4000 dilution, Pierce, USA) was added and incubation continued for 1 h at 37°C.

For the analysis of specific antibodies, Horseradish Peroxidase (HRP)-conjugated goat anti-mouse IgG (Promega, USA) (1:2500 dilution); rabbit anti-mouse IgM (μ chain) (1:2500 dilution), IgG₁ (gamma 1) (1:1000 dilution), IgG_{2a} (gamma 2a) (1:1000 dilution) or IgG_{2b} (gamma 2b) (1:1000 dilution) (Zymed[®] Laboratories, USA) was added prior to the addition of ABTS:peroxidase B (1:1; KPL, USA). The absorbance was read at 405 nm on an ELISA reader (Dynex MRX, USA).

PROTEASE NEUTRALIZATION BY ZYMOGRAPHY

The ability of anti-smBpF4 mice sera antibodies to neutralize MprA gelatinase activity was tested by zymography as previously described (Chin et al., 2007). Briefly, $10 \mu g$ of purified smBpF4 protein was pre-mixed with $30 \mu g$ of anti-smBpF4 at 37° C for 1 h. The mixtures were equilibrated to room temperature and subsequently subjected to zymography.

MOUSE PROTECTION STUDY

For the challenge experiment, smBpF4-immunized and PBS control mice were challenged with approximately 1×10^6 CFU *B. pseudomallei* D286 administered intraperitoneally 2 weeks after the second booster and mice were monitored daily for signs of mortality for 35 days post-challenge as previously described (Hara et al., 2009; Su et al., 2010).

ENUMERATION OF VIABLE B. pseudomallei IN THE BLOOD

smBpF4-immunized and challenged mice were tail-bled on days 6, 12, 18, and 25 post-infection. The blood was then plated on Ashdown agar and colonies were counted after 2 days incubation at 37° C.

STATISTICAL ANALYSIS

Statistical analysis on human and mice sera reactivity was performed using the Student's *t*-test. Data are expressed as the mean \pm standard deviation. Mortality of immunized vs. non-immunized groups was analyzed by Kaplan–Meier survival analysis and survival curves were compared using the Logrank test. Differences were considered statistically significant at

P = 0.05. All statistical analyses were performed using GraphPad Prism[®] version 4.0 (GraphPad Software, USA).

RESULTS

B. pseudomallei PROTEASES ARE IMPLICATED IN C. elegans KILLING

The prevalent view on the mode of action of extracellular proteases toward nematodes is that these proteases participate in cuticle penetration. To determine if B. pseudomallei proteases contribute to killing of C. elegans, a combination of serine protease inhibitors was added to the B. pseudomallei culture to inhibit the secreted proteases. Zymography was performed and the results confirmed that the B. pseudomallei cell-free culture filtrate gelatinase activity was indeed inhibited by 10 mM PMSF (serine protease inhibitor) and 10 mM EDTA (Zn2+ dependent metalloprotease inhibitor) (data not shown). Survival of the worms exposed to the laboratory food source E. coli OP50 remained at almost 100% over the course of the experiment (Figure 1). Survival of C. elegans exposed to bacterial cell-free culture filtrates treated with PMSF and EDTA was monitored over time and worms were observed to have an extended life span with a mean time to death (TD_{mean}) of 65.63 ± 3.70 h compared to worms exposed to B. pseudomallei D286 cell-free culture filtrates in the absence of inhibitors (TD_{mean} = 44.49 ± 1.37 h) [Logrank (Mantel-Cox) test, P < 0.0001] and B. pseudomallei D286 (TD_{mean} = 31.35 ± 0.91 h) (**Figure 1**). PMSF is a common inhibitor of protease hydrolytic activity and mortality of nematodes exposed to PMSF-treated culture filtrates was significantly delayed, indicating that the proteases most likely killed nematodes through protein hydrolysis and not via the typical ligand-receptor interaction mechanism as exemplified by the Bt toxin produced by Bacillus thuringiensis (Beale et al., 2006). The survival assay validated the importance of *B. pseudomallei* proteases as virulence factors in pathogenesis.

RECOMBINANT MprA PROTEASE (smBpF4) IS A PATIENT SERO-REACTIVE PROTEIN

To further evaluate and study the sero-reactivity of the smBpF4 protease, the full length recombinant 55 kDa smBpF4 protein was expressed and purified as previously described (Chin et al., 2007). The expressed smBpF4 protein was subsequently tested against melioidosis patient sera, legionellosis patient sera and healthy subject (HIV-negative) sera. ELISA analyses demonstrated that smBpF4 reacted with the melioidosis sera but not toward the legionellosis and healthy sera (**Figure 2**). The results indicated that the presence of anti-MprA antibodies is specific to melioidosis patients.

RECOMBINANT MprA PROTEIN IS IMMUNOGENIC AND INDUCES STRONG ANTIBODY RESPONSES

To assess the immunogenicity of the recombinant smBpF4, sera were collected and pooled from smBpF4-immunized mice prior to and following *B. pseudomallei* challenge, and tested for reactivity with purified smBpF4 protease by ELISA. The specific antibodies against smBpF4 protease were observed in immunized groups over several dilutions screened but not in the preimmunized sera (P < 0.001) (**Figure 3A**). In addition, the ELISA results (**Figure 3B**) demonstrated that the anti-smBpF4 also binds





melioidosis-confirmed, healthy control and legionellosis-confirmed sera. Significance was determined by Student's t-test (*P < 0.05; **P < 0.01).

to proteins in the *B. pseudomallei* lysate. To identify the type of immune response induced, we determined the levels of specific IgG subclasses (IgG_1 , IgG_{2a} , and IgG_{2b}) and IgM by ELISA. The predominant Ig subclass elicited by smBpF4 immunization was

IgG₁, followed by IgG_{2b} and to a lesser extent, IgG_{2a} (**Figure 3C**). The ratio of IgG_{2a}/IgG₁ was 0.817 in smBpF4-immunized mice (Figure 3C), indicating a Th2-type immune response in mice. The ELISA results indicate that recombinant MprA protease is an immunogenic protein able to trigger specific antibodies in the immunized mice.

To evaluate the ability of the anti-SmBpF4 antibodies to neutralize the protease, the polyclonal antibodies were reacted with smBpF4 protease and analyzed by zymography. smBpF4 gelatinase activity was successfully inhibited by anti-smBpF4 (Figure 3D). Our results thus far support the proposal that when smBpF4 is utilized as an immunogen, antibodies raised against the recombinant MprA protease should be able to react with and neutralize proteases secreted during an active infection, hence preventing the proteolytic digestion of the infected individual's key physiological proteins.

ANTIBODIES RAISED AGAINST smBpF4 ELICIT POTENTIAL **PROTECTIVE IMMUNITY IN MICE**

To determine if the immune response induced by immunization with smBpF4 could confer protection against B. pseudomallei infection, smBpF4 immunized-animals were challenged with $10 \times LD_{50}$ of *B. pseudomallei* D286 3 weeks after the second immunization boost and survival was observed. We had previously reported that the LD₅₀ of *B. pseudomallei* D286 was 10⁵ CFU following infection of BALB/c mice via an i.p. route (Hara et al., 2009). A challenge dose of $10 \times LD_{50}$ is expected to cause 100% mortality of the infected animals within the observation period. Thus, any reduction in the number of dead animals or a



significant delay in the time taken for all the animals to succumb to the challenge in immunized mice would indicate the potential protective efficacy of this recombinant smBpF4 protease.

Mice immunized with smBpF4 (n = 4) demonstrated better survival rates than control immunized mice (Figure 4A). In the control group of mice (n = 3) immunized with PBS in adjuvant and subsequently challenged with B. pseudomallei, one mouse succumbed on day-2 post challenge, the second on day-6 and the last mouse died on day-9 with a median survival of 6 days (Figure 4A). All of the smBpF4-immunized mice (4/4) survived beyond 25 days post-challenge. However, viable B. pseudomallei were still detected and the CFU counts increased in the blood of smBpF4-immunized and challenged mice over the 25 days postchallenge (Figure 4B). This eventually led to bacteremia in the smBpF4-immunized and challenged mice, a hallmark of acute melioidosis. Concomitantly, one mouse died on day-26, the second on day-27 and the third on day-30. Thus, recombinant smBpF4 demonstrated the ability to confer protection against B. pseudomallei infection where the median survival was prolonged by 22.5 days (P = 0.0101) in immunized animals when compared to the control group. However, immunization with a single B. pseudomallei immunogen was unable to provide complete protection probably due to the lack of T cell stimulation (Patel et al., 2011) as well as the fact that B. pseudomallei pathogenesis is most likely multi-factorial. The challenged surviving

mice also displayed splenomegaly and multiple abscess formation, demonstrating incomplete bacterial clearance from the system (**Figure 4C**) similar to that previously reported by Breitbach et al. (2008).

B. pseudomallei can produce different colony morphologies in vitro and in vivo. These morphotypes are associated with phenotypes that alter the ability of B. pseudomallei to survive in adverse environmental conditions, which could have major implications for the development of therapeutics and vaccines (Chantratita et al., 2007; Tandhavanant et al., 2010). Interestingly, we also observed changes in colony morphotypes in the smBpF4immunized and challenged mice over the infection period (Figure 4D). We identified the colony morphotypes observed in our study using the colony morphotyping scheme reported by Chantratita et al. (2007). B. pseudomallei D286 used in the challenge experiment presented a parental type 1 (the "cornflower head" morphology) colony morphology when cultured in vitro. Over time, this morphotype switched from parental type 1 to II (small and rough colonies) or type III (large and smooth colonies) (Figure 4D). The type III morphotype is reported to have a fitness disadvantage and lower intracellular replication rates in human macrophages when compared to parental type 1 morphotype (Chantratita et al., 2007; Tandhavanant et al., 2010). The switch to the type III colony morphotype predominantly seen throughout the infection period is most likely the contributing factor in the



observed prolonged survival (**Figure 4A**) and low bacterial counts (**Figure 4B**) in smBpF4-immunized and challenged mice relative to the initial inoculums.

DISCUSSION

Mortality associated with melioidosis remains high because of the lack of antimicrobial drugs, which could improve the outcome during the first 48 h after hospital admission (Peacock, 2006). This highlights the need for melioidosis-specific drugs to be administered immediately upon diagnosis to avoid a fatal outcome. Prophylactic recombinant *B. pseudomallei* proteins have been evaluated as potential candidate vaccines in animal models of melioidosis. The recombinant *B. pseudomallei* outer membrane protein A (OmpA, BPSL2522) (Hara et al., 2009) and Omp85 (BLSL2151) (Su et al., 2010) were shown to induce protective immunity against melioidosis. Pathogen surface-associated proteins have predominantly been the first pick for evaluation as potent antigens but recent findings suggest that secreted enzymes such as proteases may be effective targets as well (Jorgensen and Buchmann, 2011). Proteases from other pathogens, including parasites (Pechine et al., 2011) and bacteria (Murthy et al., 2009), are highly immunogenic and protective in animal models of infection. Thus, in this study, we extended the investigation to evaluate the prophylactic properties of recombinant *B. pseudomallei* serine MprA protease, a proposed virulence factor, which has been shown to cause extensive damage of mammalian physiological proteins although its role in the pathogenesis of melioidosis is still unclear.

B. pseudomallei MprA is located on Chromosome 2 of the *B. pseudomallei* K96243 genome at position 2,695,482–2,696,984. This protease is highly conserved among the Burkholderia spp. and subtilisins from other bacteria, particularly at the subtilisin-like catalytic triad residues (Asp 102, His 57, and Ser 195) (Lee and Liu, 2000). Our laboratory had previously utilized anti- *B. pseudomallei* protease polyclonal antibodies as a tool to investigate the protease-antibody interaction. We isolated phage displayed fusion peptides carrying a consensus peptide sequence, TKSMALSG, which closely resembled part of the *mprA* gene active site sequence, ⁴³⁵GTSMATPHVAG⁴⁴⁵ (Chan and Nathan, 2005). Homology modeling of the MprA protein

based on subtilisin demonstrated that the MprA catalytic binding pocket inclusive of the consensus sequence TKSMALSG could be the major potential antibody binding epitope (Chan and Nathan, 2005). Furthermore, we also proposed that B. pseudomallei MprA protease is produced within the bacterial cytosol as a pre-proenzyme that is subsequently processed utilizing a mechanism similar to subtilisin to produce the secreted mature active protease (Chin et al., 2007). The mprA gene is present in all 68 B. pseudomallei isolates tested by Lee and Liu (2000) as well as 11 local Malaysia B. pseudomallei animal isolates (Chin et al., data not shown). Two studies which utilized a PCR assay targeting the mprA gene of B. pseudomallei reported 100% sensitivity and specificity in detecting the pathogen in naturally infected dromedary and clinical samples, suggesting that MprA protease could be a potential drug target upon diagnosis of melioidosis (Neubauer et al., 2007; Suppiah et al., 2010).

Our group recently reported that different isolates of B. pseudomallei have divergent ability to kill the soil nematode C. elegans (Lee et al., 2011), thus providing us with a viable model to identify and characterize virulent determinants of B. pseudomallei. In this regard, the virulence of B. pseudomallei proteases toward C. elegans was assessed by inhibiting the gelatinase activity with EDTA and PMSF. The prolonged survival of worms exposed to inhibitor-treated bacterial cell free culture (Figure 1) confirms the pathogenic role of *B. pseudomallei* serine proteases in melioidosis. To follow up on these findings, preliminary characterization of the recombinant MprA protease as a potential vaccine candidate in controlling B. pseudomallei infection was initiated in a mouse model of B. pseudomallei infection. We clearly demonstrated that smBpF4 protein is immunogenic in mice whereby high levels of specific neutralizing IgG capable of inhibiting gelatinase activity were induced (Figure 3D), suggesting that immunization with MprA protease can potentially induce an appropriate neutralizing immune response to protect the B. pseudomallei-infected individual from the damages inflicted by MprA protease.

A key issue in vaccine development today is the type of immune response needed to best protect against organisms for which there are currently no effective vaccines. Most of the present generation of successful vaccines depends primarily on generating high antibody titres and many are administered with the Th2-biasing adjuvant alum. However, natural protection against many organisms, particularly intracellular bacteria, is mainly Th1 in nature (Beverley, 2002). B. pseudomallei is an intracellular facultative pathogen that can successfully evade the host immune system for an extended period of time. This has made vaccine development for melioidosis difficult and challenging and has driven researchers to develop a successful anti-B. pseudomallei vaccine that can strongly induce both humoral antibodies and a cellular-mediated immune response. Animal models of B. pseudomallei infection have been used for the development of effective vaccines; however the level of protection varied depending on the different vaccine strategies (Brett and Woods, 1996; Atkins et al., 2002; Engelbert et al., 2004; Chen et al., 2006; Haque et al., 2006; Hara et al., 2009; Su et al., 2010). For instance, immunization with attenuated B. pseudomallei mutants and active immunization with B. pseudomallei lipopolysaccharide resulted

in 60 and 50% survival at day-35 post-challenge of immunized mice, respectively (Nelson et al., 2004; Haque et al., 2006), whilst immunization with recombinant B. pseudomallei outer membrane protein Omp85 protected 70% of immunized mice up to 14 days (Su et al., 2010). Hence, data from our study is generally comparable to these previously reported efficacies. We demonstrated that all mice immunized with smBpF4 protease survived beyond 25 days of an active B. pseudomallei infection whereas the non-immunized mice succumbed within 9 days after infection (Figure 4A). Furthermore, candidate vaccines that achieved partial protection against the disease in mice suggests potential for human use and reduction in disease severity alone would be predicted to improve outcome of high mortality rates (Peacock et al., 2012). B. pseudomallei survival under adverse environmental conditions is likely to be facilitated by phenotypic adaptability and plasticity (Chantratita et al., 2007). The morphotype III observed in smBpF4-immunized mice was previously reported to be more resistant to the antimicrobial peptide LL-37 (Tandhavanant et al., 2010). This suggests that B. pseudomallei D286 undergoes a phenotypic change to adapt, survive, and replicate in the presence of bactericidal substances within the smBpF4-immunized host milieu.

The type of immune response induced by a vaccine antigen is very important for the success of the vaccine (Yuan et al., 2011). Different IgG subclasses have distinct effector functions. In the context of vaccination, different antigenic stimuli preferentially induce different antibodies (Findlow et al., 2006) and different domains of the same antigen can induce antibodies of different IgG subclasses (Soininen et al., 1999). In mice, immunoglobulin isotype switching from IgM to IgG₁ is associated with a Th2-type response, whereas IgG_{2a} is associated with a Th1-type response and promoted by interferon gamma (Chen et al., 2006; Yuan et al., 2011). Furthermore, IgG_{2a} has been shown to be the most effective IgG subclass antibodies in binding to FcyR1 on phagocytic cells (Li et al., 2007). Several studies have demonstrated that mice with an induced Th1-type immune response showed greater resistance to B. pseudomallei compared to mice with a Th2-type immune response (Hoppe et al., 1999; Ulett et al., 2000; Healey et al., 2005). The antibody subclass analysis of sera from smBpF4-immunized mice revealed production of both IgG2a and IgG₁, the higher levels of the latter indicate a dominant Th2 response over Th1. Furthermore, induction of a Th2-response was also noted for the prophylactic properties of the B. pseudomallei recombinant Omp85 protein (Su et al., 2010). Proteases from parasites (Jorgensen and Buchmann, 2011), Clostridium difficile (Pechine et al., 2011), V. harveyi (Cheng et al., 2010) and Vibrio parahaemolyticus (Liu et al., 2011) have also been proposed as promising vaccine candidates able to modulate Th1/Th2 responses in vaccinated animals.

In summary, our study has demonstrated that the *B. pseu-domallei* MprA protease is a promising prophylactic candidate against *B. pseudomallei* although the protection provided by this single protein was incomplete. Earlier studies have also noted that administration of single *B. pseudomallei* antigens is insufficient to confer total protection against melioidosis, therefore, a combination of multiple immunogens should be evaluated for a more complete anti-*B. pseudomallei* protection.

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