

In search of a combined brucellosis and tuberculosis vaccine for cattle

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ABSTRACT. Bovine brucellosis is caused by *Brucella abortus*. The bacterial pathogen causes economic losses because it induces abortion in cattle. Vaccination of calves with live *B. abortus* strain 19 induces a certain level of protection but induces persistent antibodies against cell envelope lipopolysaccharide that make it difficult to Distinguish Infected from Vaccinated Animals (DIVA). Live vaccine *B. abortus* strain RB51 was developed to eliminate such interfering antibodies and therefore, facilitate the differentiation of infected from vaccinated animals and help in the eradication of the disease. Vaccination with strain RB51 induces levels of protection similar to strain 19 but neither of the two vaccines give complete protection. We have been working to enhance protection induced by strain RB51 vaccine. Protective *Brucella* antigens can be over-expressed in strain RB51 by introducing a plasmid containing the *leuB* gene and the genes encoding such antigens. To avoid the expression of antibiotic resistance genes, we produced a *leuB* deficient strain RB51 and introduced a plasmid containing the *leuB* gene and the genes to be over-expressed. This new strain maintains the plasmid and has induced significantly high protection levels in mice. In addition, it allowed the construction of an RB51 vaccine strain able to express *Mycobacterium bovis* protective antigens so that the vaccine could protect against brucellosis and tuberculosis simultaneously.

Key words: *Brucella*, *Mycobacterium*, vaccine, RB51, protection.

Animal brucellosis is a disease affecting various domestic and wild life species and is caused by an infection with bacteria belonging to the genus *Brucella*. The genus has several species and each species has a preference for specific animal hosts, for example, *B. abortus* mainly infects cattle, *B. melitensis* infects mainly goats and sheep and *B. suis* infects mainly pigs. These small Gram negative bacteria form a genetically coherent taxon which is related most closely to *Ochrobactrum* and more distantly to *Agrobacterium* and *Rhizobium* within the alpha-2 subgroup of the *Proteobacteriaceae*.

Most of the *Brucella* nomen species are transmissible to humans where it can cause serious acute and chronic disease with some cases resulting in death, making brucellosis an important zoonosis. *Brucella* is also a facultative intracellular parasite; the pathogenesis of brucellosis and the nature of the protective immune response are closely related to this property (Cheers 1984).

Because of the serious economic and medical consequences of brucellosis for both cattle farmers and humans in general, efforts have been made to prevent the infection through the use of vaccines (Nicoletti 1990). These were initially developed on an empirical basis, but with our current ability to manipulate the genome of the bacteria, more rational designs are being used for the development of better vaccines against the disease.

The lipopolysaccharide (LPS) molecule is closely associated with the phenotype of *Brucella* colonies in culture. *Brucella* can present itself upon culture with either a “smooth” or a “rough” colony morphology, with some

strains presenting a “mucoid” phenotype (White and Wilson 1951). It is possible for smooth colonies to become rough spontaneously and some rough *Brucella* strains may revert to the smooth morphology making the degree of stability an important consideration in vaccine development. Virulence of *B. abortus* is associated with smooth morphology and therefore, *B. abortus* field strains producing acute and chronic brucellosis in cattle (and humans) are smooth. Smooth organisms have an LPS molecules containing a polysaccharide O-chain made from a homopolymer of perosamine (N-formyl-4-amino,4,6-dideoxy mannose), while rough organisms lack this O-chain on their LPS molecule; some strains possess only a greatly truncated portion of it (Caroff *et al* 1984, Moreno 1984). The O-chain of smooth *Brucella* is located on the surface of the organism. Essentially all animals infected with smooth strains respond immunologically to this O-chain by making antibodies against it. This is a very important point because, all serological tests used in the diagnosis of infected cattle are based on the detection of O-chain antibodies (Diaz R *et al* 1968, Nielsen *et al* 1988, Vemulapalli *et al* 2000). Also, it is very important to be aware that antibodies against the O-chain do not protect cattle against infection with *B. abortus* and in general, a clear role of any other *Brucella* antibodies in protection against cattle brucellosis has not been demonstrated. An effective immune response against *Brucella* infection in cattle requires a strong Cell Mediated Immune (CMI) response based on both, active T-helper1 and T-cytotoxic cells (Schurig *et al* 2002). The current understanding is that protection in cattle is highly dependent on the induction of these specific T lymphocytes and the concurrent activation of macrophages by T-helper1 cells secreting interferon-gamma (Schurig *et al* 2002, Vemulapalli *et al* 2000^c).

The vaccine that has been most widely used to prevent bovine brucellosis is *B. abortus* strain 19 (Nicoletti 1990).

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This strain was first described in 1930 and was originally isolated from the milk of a Jersey cow as a virulent strain in 1923, but after being kept in the laboratory at room temperature for over a year, was found to have become attenuated (Buck 1930). This strain was able to induce a certain level of protective immunity in cattle. Effectiveness fluctuated depending on a variety of variables including age of vaccination, dose, route and prevalence of brucellosis in vaccinated herds (Nicoletti 1990). Strain 19 is an attenuated organism with a smooth morphology normally unable to grow in the presence of erythritol (Jones *et al* 1965). Although strain 19 is of low virulence for cattle, vaccination of pregnant cows can result in abortions. This event is rather rare, ranging from less than 1 up to 2.5% under field conditions and between 10-12% under experimentally controlled conditions (Mingle *et al* 1941, Moore 1950, Manthei 1952, Beckett and Mc Diarmid 1985, Tabynov *et al* 2016). Intravenous injections of pregnant cattle with strain 19 induces 100% abortions, while RB51 did induce up to 25% abortion (Palmer *et al* 1996). Strain 19 is pathogenic for humans and can lead to chronic infections if cases are not treated with appropriate antibiotics (Revich *et al* 1961, Wallach *et al* 2008). The most common route of human infections with strain 19 is accidental inoculation during vaccination of cattle. Strain 19 can also be found in the milk of cattle if vaccination is carried out during adulthood (Samartino *et al* 2000).

Since strain 19 is a smooth species of *Brucella*, the presence of LPS with O-chain on its surface is the reason for the appearance and persistence of O-chain antibodies in serum following administration of this vaccine (Diaz *et al* 1968, Nielsen *et al* 1988). Since these antibodies are the basis of all diagnostic tests, strain 19 vaccination prevents easy differentiation of vaccinated from infected animals and therefore, delays eradication efforts and leads to over condemnation of cattle. In order to decrease this problem to some extent, calves are vaccinated rather early in life since their antibody response to *Brucella* is weaker at that time. The problem becomes worse if calves are vaccinated for the first time later in life or are re-vaccinated for booster purposes since antibody levels will increase even more and be very persistent (Manthei 1952, Samartino *et al* 2000).

Several undesirable characteristics of strain 19, particularly its confusing effect on sero-diagnosis, led to the development of *B. abortus* strain RB51 vaccine (Schurig *et al* 1991). Strain RB51 is a highly attenuated mutant of *B. abortus* strain 2308. It is a very stable strain meaning that it does not change colony morphology upon multiple passages in culture or animals (Schurig *et al* 1991, Colby 1997). Strain RB51 has a rough colony morphology and is essentially devoid of O-chain (Schurig *et al* 1991). The *wboA* gene of strain RB51, a gene involved in the biosynthesis of the O-chain, is interrupted by an IS711 element impeding the production of O-chain (Vemulapalli *et al* 1996, 2000^a). Since strain RB51 lacks O-chain, vaccinated cattle do not respond to the O-chain and do not develop

antibodies to this antigen. Therefore, vaccinated animals remain serologically negative in all serological diagnostic tests. Furthermore, re-vaccination at any age does not induce O-chain antibodies in the animals allowing the application of “booster” vaccinations without affecting serological diagnostic tests necessary for disease eradication (Samartino *et al* 2000, Dorneles *et al* 2015). When used in cattle, one vaccination with strain RB51 induces protection levels similar to those induced by strain 19 and protection of the vaccinated animals can vary from 65% to 100% depending on the conditions prevailing in a specific herd or experiment (Schurig *et al* 2002). Revaccination with strain RB51 for a second time appears to increase immunity and represents an additional advantage of RB51 over strain 19. Since 1996, millions of cattle have been immunised with strain RB51 and reversion to a virulent form has never been observed testifying to its extreme stability.

As mentioned before, CMI responses play a critical role in resistance against intracellular bacterial infections (Schurig *et al* 2002, Vemulapalli *et al* 2000^c). It is therefore critical that vaccines are presented to the immune system in a way that they induce a T cell response that can protect against the infection. Live bacterial vaccines are considered essential to induce an appropriate T cell mediated CMI protective response. Although the exact reasons for needing live organisms to induce the right response are debatable and are probably multiple, the synthesis of antigens by the live organisms during the immune response induction phase appears critical. Replication of the vaccine strain may be less critical than synthesis of new antigens since irradiated vaccines, where the bacteria do not replicate but are able to synthesize antigens after the irradiated vaccine application, are protective. Irradiated *Brucella* are able to induce protection through CMI responses while killed organisms are not (Montaraz and Winter 1968, Magnani *et al* 2009, Moustafa *et al* 2011). Since live bacteria are able to induce protective CMI responses, attenuated strains of bacteria are often used as live vaccines to protect against intracellular bacterial infections. Both *B. abortus* strain RB51 and strain 19 are live, attenuated vaccines able to induce protection while the same vaccines rendered metabolically inactive are not unless potent adjuvants are used (Montaraz and Winter 1968). This observation has important practical implications since live, attenuated vaccines have to be handled carefully (kept cold, use only shortly after reconstituting in the field, etc) to maintain their effectiveness.

In many cases live, attenuated vaccines do not provide high levels of protection, particularly if animals are confronted with high numbers of infectious bacteria (Manthei CA 1968). For example, a *Brucella* vaccinated cow will have a very high likelihood of being well protected against infection if she is exposed to a low number of field *B. abortus* bacteria (Manthei CA 1968). Not only does level of exposure affect the protective ability of the vaccine, vaccine dose will also affect the protective outcome (Manthei CA

1968, Confer *et al* 1985). In contrast, a cow may not be protected when exposed to the placenta and fetus of a *B. abortus* abortion where *Brucella* number in the hundreds of billions. Therefore, eradication of brucellosis not only depends on vaccination, it depends on a sustained vaccination program and good management where vaccinated animals are separated as much as possible from high level infection sources and where seropositive animals are consistently removed from the herd.

The fact that full protection in cattle is probably not achieved under most field conditions, made it relevant to work on approaches that could improve effectiveness of the current strain RB51 vaccine without changing its positive characteristics of being highly attenuated and leaving animals seronegative after one or multiple vaccinations. We hypothesized, some time ago, that over-expression (production of more than normal quantities) of a *Brucella* protective antigen by vaccine strain RB51 would result in enhancement of the vaccine's efficacy (Vemulapalli *et al* 2000^b, Vemulapalli *et al* 2002). Our studies demonstrated that we could over-express *B. abortus* Cu/Zn superoxide dismutase (SOD) protein in vaccine strain RB51 (Vemulapalli *et al* 2002). SOD is considered to be one (He *et al* 2002) of probably many protective *Brucella* antigens and significantly increases the vaccine's protective capabilities in the murine model of brucellosis without altering the attenuation, stability or serological characteristics of the vaccine. This stimulated our interest in pursuing this approach to produce a more effective RB51 vaccine. Interestingly, even though homologous over-expression of Cu/Zn superoxide dismutase SOD enhanced protection significantly in mice, it did not enhanced protection against *B. abortus* infection in bison and elk (Olsen *et al* 2009, Nol *et al* 2016). In bison, the overexpressing RB51-SOD vaccine was actually less efficacious and was cleared faster from the vaccinated animals than the parenteral RB51 strain (Olson *et al* 2009). Faster clearance may have resulted in a weaker immune response and may explain the decrease in efficacy. This indicates that the murine model is not always an indicator of what may happen in cattle or other animal species making protection experiments in cattle crucial before a new vaccine can be called more effective in the target species.

Since homologous over-expression of a protective antigen can lead to a more effective vaccine, it was logical to think that expression by strain RB51 of a protective antigen derived from an unrelated infectious agent, for which protection is mediated by a CMI response, would protect against the unrelated disease as well as against infection with *Brucella*. Consistent with this hypothesis, protection against infection with *Neospora caninum* was demonstrated in the mouse model using a strain RB51 strain expressing parasite protective antigens (Rajasekaran *et al* 2008). Therefore, it may be possible to create strain RB51 vaccines able to protect cattle against *Brucella*

infection as well as protect against unrelated diseases simultaneously. Our approach to create a strain RB51 vaccine able to protect against *Brucella* and *Mycobacterium bovis* infections simultaneously is described below.

In order to use strain RB51 as a platform vaccine able to induce specific immune responses against a variety of homologous and/or heterologous antigens, a plasmid containing the gene encoding the foreign antigen along with an antibiotic resistance gene (as a plasmid marker) had been employed during our developmental work carried out in the past (McQuiston *et al* 1995, Vemulapalli *et al* 2000^c, Vemulapalli *et al* 2002). Because *Brucella spp.* have no naturally occurring plasmids, we adapted a broad host range plasmid that had been shown to replicate in *Brucella* (Kovach *et al* 1994). We eliminated extraneous plasmid DNA sequences and introduced a variety of promoters to allow for expression of genes derived from *Brucella* (homologous overexpression) as well as derived from other bacterial species (heterologous overexpression) and the resulting plasmid was designated as pNS or pLeuB (Seleem *et al* 2004, see figure 1).

The approach of using an antibiotic resistance gene as a selection marker has been criticized as not being environmentally safe because in the vaccinated animals, it has the potential to introduce the antibiotic resistance gene into normal and pathogenic flora. Therefore, expressing the protective antigens from a plasmid that is not dependent on an antibiotic resistance gene for selection of the desired strain, would be an advantage, as it would have a minimum environmental risk.

The *leuB* gene, encoding isopropyl malate dehydrogenase, is one of the four genes essential for the biosynthesis of leucine in *B. abortus* (Essenberg *et al* 1993). *B. abortus* is known to survive and replicate in nutrient-limited environments inside a host. Therefore, a *B. abortus* mutant lacking a gene for the biosynthesis of an essential amino acid like leucine is unlikely to survive in that environment including being able to replicate in minimal media (Essenberg *et al* 1993, Bacon *et al* 1951, Bange *et al* 1996). The complementation of such an auxotroph with a plasmid carrying the wild-type *leuB* gene (encoding the enzyme necessary for the amino acid synthesis) would provide a means of selection and maintenance of this plasmid in *B. abortus* under minimum media conditions. Thus, we produced a *leuB* mutant in strain RB51 using allelic exchange (Rajasekaran *et al* 2008). The resultant *leuB* auxotroph (RB51*leuB*) cannot grow in leucine-deficient conditions but when complemented with a plasmid carrying the wild-type *B. abortus leuB* gene, the leucine deficiency of RB51*leuB* is eliminated (Rajasekaran *et al* 2008). Thus, the complemented RB51*leuB* strain can be used to over-express homologous and/or heterologous antigens and eliminate the environmental concerns related to antibiotic resistance. Importantly, the complemented RB51*leuB* strain retained the basic characteristics of the original strain RB51 which are attenuation, no induction

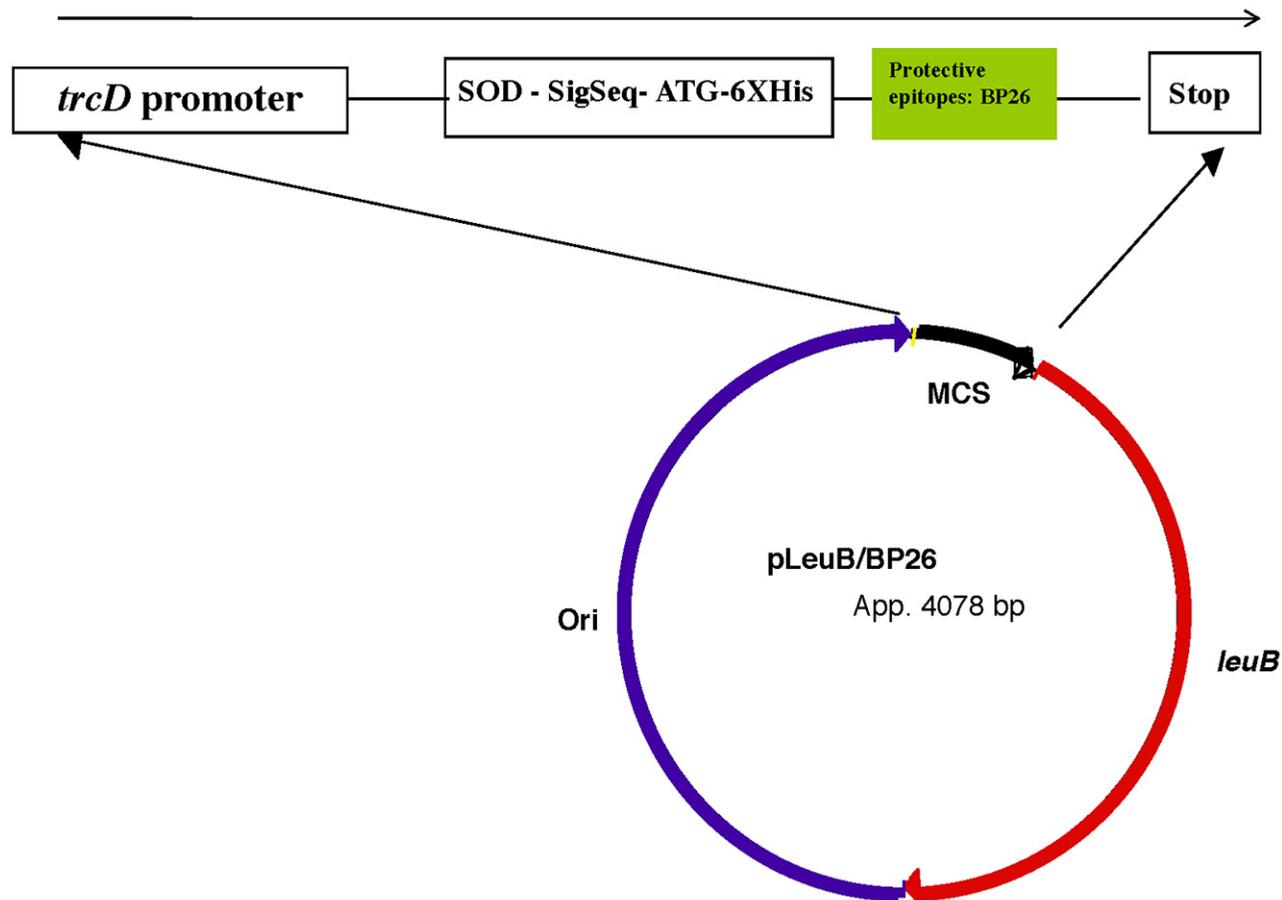


Figure 1. Schematic of pLeuB plasmid containing the *leuB* gene of *B. abortus*; a multiple cloning site (MCS) and an origin of replication (Ori). The *trcD* promoter is a very strong promoter as it is a hybrid between the *lacZ* and tryptophan promoter. The signal sequence (SigSeq) is from the SOD gene and followed by an ATG start codon and an in-frame histidine tag (6XHis) followed by a protective antigen from BP26 of *B. abortus*.

of O-chain antibodies and protection against *B. abortus* infection.

Using strain RB51 *leuB* as a platform and pNSLeuB, an antibiotic-resistance marker free plasmid, we constructed several strains over-expressing homologous antigens. For example, strains RB51 *leuB*/SOD (superoxide dismutase), RB51 *leuB*/SOD/L7/L12 (ribosomal proteins) and RB51 *leuB*/SOD/WboA (glycosyl transferase) were constructed to over-express the selected *Brucella* antigens: SOD alone, SOD and ribosomal protein L7/L12 or SOD and glycosyl transferase, respectively. The ability of these vaccine candidates to protect against a virulent *B. suis* challenge were evaluated in a mouse model. All vaccine groups protected mice significantly ($P < 0.05$) when compared to the control group. Within the vaccine groups, the mice vaccinated with strain RB51 *leuB*/SOD/WboA were significantly better protected than those that were vaccinated with either strain RB51 *leuB*/SOD or RB51 *leuB*/SOD/L7/L12. These results suggest that *Brucella* antigens can be over-expressed in strain RB51 *leuB* and can elicit enhanced protective immune responses against brucellosis at least in the murine model

of the disease (Rajasekaran *et al* 2011). The murine model for Brucellosis has been used by numerous authors to test protective abilities of vaccines (Montaraz and Winter 1968). Nevertheless, it is not clear if protective abilities observed in mice translate into protective abilities in cattle or other species. For example, work carried out with strain RB51 in mice indicating protection by strain RB51 was later demonstrated in cattle (Schurig *et al* 1991, Cheville *et al* 1993). On the other hand, RB51 overexpressing Cu/Zn superoxide dismutase showed higher protection than RB51 in mice but did not protect bison against *Brucella* challenge (Vemulapalli *et al* 2002, Olson *et al* 2009). It remains to be seen if similar results can be obtained in cattle.

Control of bovine tuberculosis is an important element in the control or eradication of human tuberculosis. Unfortunately, there is no practical vaccine available for the control of tuberculosis in cattle even so, the problem is very significant. To illustrate the dimension of the problem, long term data analysis of bovine tuberculosis in India suggests that approximately 7.3% of cattle are potentially infected. This means that there are 21.8 million

TB infected cattle, which amounts to more than all cattle in United States. This systematic analysis is even more alarming as approximately 10% of human tuberculosis is due to *M. bovis* infection. Therefore, control of bovine tuberculosis is an important component related to efforts of human tuberculosis eradication, especially in the developing world, as test and slaughter may not be an option due to economic and religious considerations (Srinivasan *et al* 2018).

Bacille Calmette Guerin (BCG) vaccine is the current TB vaccine used in humans. It is an attenuated strain of *M. bovis* (Behr *et al* 1997, Murray 2004). BCG was attenuated by serially passing a virulent *M. bovis* strain on ox-bile medium for 230 times in the laboratory. This process led to attenuation of the virulent *M. bovis* strain as indicated by self-limiting infection as well as partial resistance to reinfection with *M. bovis* (Takamura *et al* 2005). One of the serious disadvantages of BCG vaccine is that vaccinated humans and animals turn skin test positive. Therefore, BCG is rarely used in cattle to protect against bovine tuberculosis (Waters *et al* 2012). Recent, largescale field trials with cattle using very low doses of BCG demonstrated high levels of protection (vaccine efficacy between 85.7%-86.7%) without affecting their *ante mortem* skin or blood tests (Nugent *et al* 2018). This low dose approach may make vaccination of cattle with BCG more acceptable than in the past.

Following our previously defined strategy of over-expressing heterologous antigens in strain RB51, we speculated that expression of protective antigens from *M. bovis* in *B. abortus* strain RB51, a vaccine strain that is USDA approved and extensively used throughout the world, could make an effective dual vaccine able to protect against Brucellosis and Tuberculosis in cattle simultaneously. To prove this point, we selected the following protective *Mycobacterium* antigens: 85B, ESAT6 and Rv2660c for this purpose (Al Qublan, Dissertation, Virginia Tech, 2014). These genes were cloned and expressed in our auxotrophic *leuB* deletion mutant of strain RB51 (RB51*leuB*). As explained before, the advantage of using the RB51*leuB* deletion mutant is that one can have a plasmid encoding the protective antigens along with the *leuB* gene instead of a drug resistance gene for selection purposes

The reasons for selecting these antigens are outlined by characteristic that define them as protective antigens.

ANTIGEN 85B

Antigen 85B (Ag85B) is one of the most dominant protein antigens secreted by all mycobacterial species (Belisle *et al* 1997). Ag85B (30 kDa) belongs to the Ag85 complex, which is a family of three structurally related fibronectin-binding proteins (Ag85A, Ag85B and Ag85C) with mycolyl-transferase activity that is involved in the final stages of cell wall assembly. Ag85B protein is not only the major secretory protein of *M. tuberculosis* in broth culture,

but it is also among the major proteins of all *M. bovis* proteins expressed and, it is a major stimulator of T-cell proliferation and IFN- γ production in most healthy-looking animals infected with *M. bovis*. Studies have shown that immunisation with plasmid DNA encoding Ag85B can stimulate strong cell-mediated immune response and confer significant protection to mice challenged with virulent *Mycobacterium* (Lozes *et al* 1997). All these findings suggest that Ag85B is a promising protective antigen for vaccine use in other animal species.

EARLY SECRETORY ANTIGENIC TARGET 6

Early secretory antigenic target 6 (ESAT6) is a protein encoded by the region of difference 1 (RD1) of the *M. bovis* genome. RD1 has been shown to be a major virulence factor involved with membrane-lysing activity (Smith *et al* 2008, Gao *et al* 2004). Although the exact function of ESAT6 has not been determined, studies have shown that deletion of the ESAT6 protein results in abrogation of the necrosis-inducing effect of tuberculosis on human monocyte-derived macrophages; suggesting that ESAT6 is involved with causing necrosis (Welin *et al* 2011). ESAT6 is a strongly recognized T-cell antigen in the first phase of infection and has demonstrated protective efficacy as a subunit vaccine in animal models (Brant *et al* 2000), a DNA vaccine (Qingtao *et al* 2013), and a recombinant BCG vaccine (Pym *et al* 2003).

RV2660C

Rv2660c is a newly recognised antigen of unknown function that was first reported in a gene expression profiling study by Betts and colleagues (Betts *et al* 2003). In their study, it was reported that in nutrient-starved cultures, expression of Rv2660c increased 100 to 300 fold making it the most strongly up-regulated of all nutrient starvation-induced genes identified. In another gene expression profiling study in a mouse model, it was found that Rv2660c was expressed at high levels during early and late stages of infection (Aagaard *et al* 2011). More importantly, Govender and colleagues reported that Rv2660c was preferentially recognized by patients with latent TB compared with those with active tuberculosis (Govender *et al* 2010). These findings suggest that Rv2660c is involved in latency and therefore provides a promising vaccine candidate for targeting tuberculosis infection as it transitions into latency.

Preliminary studies carried out in our laboratories, indicated that it was crucial that the DNA sequences of the protective *Mycobacterium* antigens to be cloned into our plasmid were synthesised using the codon usage of *Brucella*. The original ESAT6 sequence had a low codon optimization Index (COI) of 0.47 in *Brucella* and when optimized for strain RB51 expression, the COI changed to 0.95. Similarly, the Ag85b COI changed from 0.51 to

0.94 and Rv2660C COI changed from 0.29 to 0.93. These optimised sequences were used in the generation of the synthetic genes encoding the three protective antigens (Genscript NJ, USA). BamHI, BglIII and XbaI sites were engineered into the sequences and the synthetic genes were then cloned into pLeub plasmid. Two strains were constructed one expressing 85B (RB51*leuB*/85B) and one expressing Rv2660c and ESAT6 as a fusion protein (RB51*leuB*/Rv2660c-ESAT6). The strain constructs were verified by PCR using specific primers for directional cloning.

After confirmation of expression of the tuberculosis protective antigens by Western blot, the above two vaccine constructs, RB51*leuB*/85B and RB51*leuB*/Rv2660c-ESAT6 were mixed in equal proportions and the vaccine denominated RB51TB and it was tested for its protective ability in one group of BALB/c mice. BCG immunisation testing was carried out in a second group of mice and PBS was used as the non-immunised control. The mice were vaccinated intraperitoneally (ip), boost vaccinated through the same route 4 weeks later and then challenged ip with *M. tuberculosis* four weeks after the booster vaccination. Four weeks after challenge, mice were killed and lungs and spleen were cultured to determine Colony Forming Units (CFUs) per organ. Vaccinated mice had significantly less CFUs in each cultured organ when compared to the non-vaccinated control group indicating good protection. There was no significant protection difference among the mice vaccinated with BCG and our strain RB51 vaccine constructs. These results clearly suggest that heterologous expression of protective tuberculosis antigens in strain RB51 can lead to protection against infection with *Mycobacterium* at least in the murine model. Mice were also protected against challenge with *B. abortus* indicating again that expression of heterologous antigens by strain RB51*leuB* does not alter its protective effect against brucellosis.

If the protective ability against mycobacterial infection in the murine model can be replicated in cattle, strain RB51TB vaccine could be used to protect cattle against brucellosis and tuberculosis simultaneously. In order for this vaccine to have practical application in the field, it not only has to protect against the two diseases, it should not induce positive tuberculin skin reactions in the vaccinated animals. If positive tuberculin reactions would be induced, the practical use of the vaccine would be limited.

In order to assess if RB51TB would induce positive tuberculin reactions after vaccination, eight tuberculin negative calves were immunised with RB51TB and booster vaccinated 4 months after the first vaccine application. A tuberculin test was carried out 4 months after the booster (unpublished). All cattle remained tuberculin negative suggesting that vaccination of cattle with RB51TB will not introduce problems related to tuberculosis and brucellosis diagnosis. One major challenge still remains, that is, to demonstrate that RB51TB protects cattle against tuberculosis. Such protection experiments, including safety

and stability experiments, will be very costly and time consuming but will be necessary before approval to use this promising vaccine is granted.

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