# Plasmid instability when the *hsp* 60 gene promoter is used to express the protective non-toxic fragment B of the diphtheria toxin in recombinant BCG

Dilzamar V. Nascimento<sup>1</sup>, Odir A. Dellagostin<sup>2</sup>, Raphael Hirata Jr.<sup>1</sup>, Geraldo M. B. Pereira<sup>1,3</sup>, Ana Luiza Mattos-Guaraldi<sup>1</sup>, Geraldo R. Garcia Armôa<sup>3</sup>

Email: guaraldi@pq.cnpq.br, guaraldi@uerj.br

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### **ABSTRACT**

The genetic modification of the live attenuated Mycobacterium bovis BCG to deliver a protective Corynebacterium diphtheriae antigen in vivo could be a safer and less costly alternative to the new and more expensive DTP vaccines available today, in particular to third world-countries. The stability of expression of heterologous antigens in BCG, however, is a major challenge to the use of live recombinant bacteria in vaccine development and appears to be dependent to a certain extent, on a genetic compatibility between the expression cassette within the plasmid construct and the mycobacterium host. In the quest for the best recombinant BCG transformant to express the dtb gene of C. diphtheriae we generated two new rBCG strains by transforming the Moreau substrain of BCG with the mycobacterial expression vectors pUS973 and pUS977, each one carrying a different promoter to drive the expression of the target antigen. After transformation recombinant BCG clones were selected on Middlebrook 7H10 kanamycin Agar plates, expanded in Middlebrook 7H9 kanamycin Broth and analyzed by agarose gel electrophoresis and immunoblotting. rBCGs transformed with the construct carrying the weak  $P_{AN}$  promoter from M. paratuberculosis stably expressed the dtb gene. Conversely, rBCGs transformed with the construct carrying the strong mycobacterium hsp60 promoter were unstable and consequently unfit for the expression of the C. diphtheriae gene.

**Keywords:** BCG Vaccine; Recombinant BCG; Fragment B of Diphtheria Toxin; Anti-Diphtheria Vaccine

### 1. INTRODUCTION

Diphtheria, the classical disease caused by C. diphtheriae is an acute communicable infection of the upper respiratory tract that can be fatal [1,2]. Although the global incidence of diphtheria has been reduced dramatically as a result of increased vaccine coverage, infections by different biotypes of C. diphtheriae continue to be reported in many countries [3,4]. The diphtheria toxin (DT) is a polypeptide with a molecular weight of about 58,000 Daltons that is cleaved into two fragments to become active. The fragment A (DTA) with 193 amino acids is an enzyme that catalytically inactivates the ribosomal elongation factor-2 of eukaryotic cells while the 342 amino acids long fragment B (DTB) recognizes and binds to specific receptors on the surface of most eukaryotic cells, facilitating the translocation of the fragment A across the membrane [5]. Despite the existence of a very good toxoid vaccine, diphtheria has not been eradicated vet. On the other hand, the current toxoid based vaccine presents risks to vaccinees associated to the production process. These risks include incomplete inactivation and the presence of reactor and unknown proteins generated during the inactivation process. For this reason the immunization against diphtheria has to be done with three doses to minimize adverse effects that could take place with a full single dose [6,7]. Adverse side reactions of that kind could be easily avoided by purification of the current vaccine. However, the purification of the diphtheria toxoid would increase the cost of production considerably.

The Bacillus of Calmette-Guérin (BCG), a live attenuated substrain of *M. bovis* obtained almost a century ago, is currently the only vaccine licensed against TB world wide. The recombinant BCG approach developed



<sup>&</sup>lt;sup>1</sup>Faculdade de Ciências Médicas, Universidade do Estado do Rio de Janeiro, Rio de Janeiro, Brazil

<sup>&</sup>lt;sup>2</sup>Centro de Biotecnologia, Universidade Federal de Pelotas, Campus Universitário, Pelotas, Brazil

<sup>&</sup>lt;sup>3</sup>Instituto Oswaldo Cruz, Fundação Oswaldo Cruz, Rio de Janeiro, Brazil

on top of it emerged in the late 80's as a creative technology for in vivo delivery of protective antigens and brought a new hope for the development of low cost multivalent vaccines [8,9]. In this context, a genetically modified BCG strain expressing a relevant diphtheria antigen could be a safer alternative to the current classic vaccine used against diphtheria in places where BCG is mandatory. Nevertheless, the stability of the expression of heterologous antigens in BCG is still a major issue and perhaps one of the most important obstacles to the use of this technology in vaccine development. In fact there are several reports in the literature showing that genetic rearrangements lead to the disruption of the expression of heterologous genes or promoters in mycobacteria [10-14]. In this paper, we report our first results over the expression of the C. diphtheriae gene  $dtb_{PW8}$  by the M. bovis BCG substrain Moreau used in Brazil to produce the vaccine against TB, modified by the transformation with distinct plasmid vectors, pUS973 and pUS977, to generate two new recombinant strains in which the expression of the target gene is driven by a strong (hsp60) or a weak ( $P_{AN}$ ) mycobacterium promoter. A comparison of the structural stability of plasmid constructs used to transform BCGs with constructs rescued after transformation and growth of the two rBCGs in culture medium are presented.

### 2. MATERIALS AND METHODS

#### 2.1. Bacterial Strains and Media

The toxigenic strain of *C. diphtheriae* Park Williams 8 (ATCC 13812) was used as the source of genomic DNA, while the *Escherichia coli* DH5α strain was used for amplification of all plasmid vectors. *C. diphtheriae* was grown in Brain Heart Infusion broth and *E. coli* in Luria-Bertani broth or agar plates with or without ampicillin (75 μg/mL). The BCG vaccine (*M. bovis* BCG, Moreau substrain) was provided by the Brazilian National producer Fundação Athaulpho de Paiva in Rio de Janeiro, RJ [15] and was typically grown at 37°C in Middlebrook 7H9 or 7H10 medium supplemented with ADC (Difco Laboratories) plus kanamycin 25 μg/mL, when needed.

### **2.2.** Construction of Mycobacterial Expression Vectors Carrying the *dtb* Gene

The DNA fragment corresponding to the *dtb*<sub>PW8</sub> gene was amplified from the *C. diphtheriae* genomic DNA by Polymerase Chain Reaction (PCR) using primers *dtb* F (5'CGTCTAGAAGGTAGCTCATTGTC3') and *dtb* R (5'GCTCTAGACCCCACTACCTTTC3') designed from the sequence encoding the diphtheria toxin, GenBank accession code K01722 [16]. The PCR reaction was performed in the Gene Amp PCR System 2400 thermo cycler (Perkin Elmer) with 30 cycles of 5 minutes de-

naturation at 94°C, 1.5 minutes annealing at 55°C and 2 minutes extension at 50°C. The PCR product was analyzed in agarose gel and the 1051 bp amplicon band corresponding to the size of the *dtb*<sub>PW8</sub> DNA was purified, digested with *Xba*I and then cloned into the same site of the expression vectors pUS973 and pUS977 [17, 18], giving rise to pUS973*dtb*<sub>PW8</sub> and pUS977*dtb*<sub>PW8</sub> constructs, respectively. The identity of the DNA inserts was confirmed by DNA sequencing.

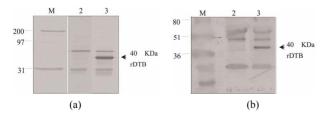
## 2.3. Transformation of BCG, Analysis of rBCG Clones for Expression of $dtb_{\rm PW8}$ and Plasmid Integrity

The Moreau BCG was transformed with pUS973dtb<sub>PW8</sub> or pUS977dtb<sub>PW8</sub>, or with an empty plasmid, by electroporation. In order to do this BCG was grown to an optical density of 600 nm in Middelbrook 7H9 broth, centrifuged twice at 4000× g for 10 minutes and resuspended to 1/20 of its original volume in 10% glycerol. Further, 200 microliters of BCG were mixed with each plasmid DNA construct (1 - 2 µg) in a pre-chilled 0.2 cm electroporation cuvette (Bio-Rad) and transformation was performed using a Gene Pulser apparatus (Bio-Rad) at 2.5 kV, 1000 Ω and 25 μF [19]. Recombinant BCG clones were selected after 14 - 21 days of incubation at 37°C on Middlebrook 7H10 agar plates containing kanamycin, expanded in Middlebrook 7H9 broth with kanamycin at 37°C and 2.0 mL aliquots of each culture were harvested at mid-log growth phase, lysed and subjected to 12% SDS-PAGE. After separation in the SDS gel, proteins were electrotransferred to nitrocellulose membranes (Bio-Rad, Hercules, California) and subjected to immunoblotting analyses to detect the expression of the DTB protein using mice anti-toxoid polyclonal antibodies (1:5000 dilution), according to standard procedures described in our previews studies [16]. Recombinant BCG clones were also analyzed with regards to plasmid integrity to confirm the presence of the  $dtb_{PW8}$ gene by restriction analyses with XbaI of plasmids recovered from rBCG expressing  $dtb_{PW8}$ , or not [20].

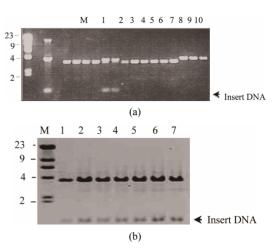
### 3. RESULTS

A 1051 bp long DNA fragment, named  $dtb_{\rm PW8}$ , necessary to build the expression vectors used in this study to modify BCG was successfully amplified by PCR from the genome of the *C. diphtheriae* PW8 vaccine strain, using specific primers designed to do it as referred above. The identity of the  $dtb_{\rm PW8}$  gene was confirmed by DNA sequencing and analysis of similarity against the sequence of DT deposited in the GenBank. The gene dtb was further cloned into the extrachromossomal mycobacterial expression vectors pUS973 and pUS977 generating constructs pUS973 $dtb_{\rm PW8}$  and pUS977 $dtb_{\rm PW8}$ , respec-

tively, and these were used to transform BCG. The two new rBCG strains resulting from transformation were evaluated in terms of their capacity to express rDTB<sub>PW8</sub> in vitro. The only difference between vectors pUS973 and pUS977 is the promoter sequence used to drive the expression of the foreign gene. The pUS973 vector carries the strong constitutive hsp60 promoter from M. tuberculosis, while pUS977 carries the  $P_{AN}$  promoter from M. paratuberculosis. The expression of DTB in rBCG was evaluated by immunoblotting assays using polyclonal antibodies against the diphtheria toxoid. As predicted, a band of the expected size for DTB, 40 kDa, was recognized in lysates of rBCGpUS973dtb<sub>PW8</sub> or rBCGpUS977dtb<sub>PW8</sub> subjected to immunoblot analysis. The level of expression of rDTB<sub>PW8</sub> observed in BCG transformed with pUS973dtb<sub>PW8</sub> was similar to BCG transformed with pUS977 $dtb_{PW8}$  (**Figures 1(a)** and **(b)**). Mycobacteria and corynebacteria have a common ancestry and may as well share antigens. This fact possibly contributed to the fact that the use of polyclonal antibodies in the immunoblotting assays led to the visualization of non-specific rest bands in both non-transformed and transformed BCG. In order to verify the structural stability of the recombinant plasmids, selected BCG transformants were subjected to two successive passages in 7H9 Middlelbrook medium supplemented with kanamycin. Subsequently, plasmid DNA was recovered by electroduction into E. coli [20]. Several colonies were selected and the purified plasmid DNA was digested with XbaI. Electrophoresis of the digested DNA revealed that most BCG clones transformed with pUS973dtb<sub>PW8</sub> had undergone mutations (Figure 2(a)), whereas none of the pUS977dtb<sub>PW8</sub> clones showed alteration in the structure of the plasmid (Figure 2(b)). In fact, a few constructs from rBCGpUS973dtb<sub>PW8</sub> (Figure 2(a), lanes 2-5) lost the XbaI site and part of the plasmid DNA since they exhibited a lower molecular mass when compared to the original vector (lanes 8-10) used as control. Also, other constructs recovered from clones of the same rBCGpUS973dtb<sub>PW8</sub> strain apparently lost part of their



**Figure 1.** Immunoblotting analysis of rBCGpUS973 $dtb_{PW8}$  and rBCGpUS977 $dtb_{PW8}$  cell lysates. (a) lane M: molecular weight markers; lane 2: non-modified BCG lysate; lane 3: rBCG-pUS973 $dtb_{PW8}$  lysate; (b) lane M: molecular weight markers; lane 2: non-modified BCG; lane 3: rBCGpUS977 $dtb_{PW8}$  lysate. The arrows at the right of gels indicate a band of the expected size for rDTB.



**Figure 2.** Agarose gel electrophoresis analysis of pUS973 $dtb_{PW8}$  and pUS977 $dtb_{PW8}$  constructs recovered from rBCGpUS973 $dtb_{PW8}$  and rBCGpUS977- $dtb_{PW8}$  clones after digestion with XbaI. (a) lane M: DNA markers (λ HindIII); lane 1: pUS973 $dtb_{PW8}$ . The top band corresponded to an artifact resulting of the non-digested plasmid applied in excess on the gel. the two lower bands corresponded to the linearized pUS973 plasmid (middle band) and the excised insert (lowest band); lanes 2-7: pUS973 $dtb_{PW8}$  clones; lanes 8-10: empty pUS973; (b) lane M: DNA markers; lane 1: pUS977 $dtb_{PW8}$ ; lanes 2-7: pUS977- $dtb_{PW8}$  clones. the arrows at the right of gels indicate the DNA insert liberated after digestion.

DNA in spite of having released the insert (lanes 6 and 7). Indeed, the instability of constructs based on the hsp60 promoter is not new. It has been shown that mutations may occur as a result of modifications of the promoter or insert sequences [13]. What is new is that it also occurs with the  $dtb_{PW8}$  gene from C. diphthheriae.

### 4. DISCUSSION AND CONCLUSIONS

The genetic modification of the BCG vaccine as an approach to the development of new multivalent vaccines has been actively pursued ever since it was proposed in 1989 [8] and has been the focus of our research over the last years [21-28]. BCG is a powerful candidate to a multivalent vaccine platform. We successfully constructed a new and stable rBCG strain able to express the non-toxic fragment B of the C. diphthteria toxin by genetically modifying the Moreau substrain of BCG used in the production of the vaccine against TB in Brazil. Although a mutant of the DT, CRM197, has been expressed in BCG using the promoter pBlaf [29], this is the first clear demonstration that the safer B fragment of DT can be expressed and well tolerated by BCG. These are definitive conditions for the development of vaccine prototypes based on this technology since not every antigen is necessarily expressed or tolerated by BCG. Our best results came out with the BCG transformed with the pUS977dtb<sub>PW8</sub> construct in which the expression of DTB is driven by the  $P_{AN}$  promoter, a regulatory sequence derived from M. paratuberculosis. The  $P_{AN}$  sequence, a weak mycobacterial promoter [18], has been successfully used by others for the expression of the  $\alpha$ -galactosidade of E. coli [30], the LacZ of E. coli [31], the Gp63 surface antigen of Leishmania [32], the Nef antigen of Simian Immunodeficiency Virus [33], the Nef, Gag, Env antigens of Simian Immunodeficiency Virus [34], the S1 subunit of the Bordetella pertussis toxin [24], the hepatitis B surface antigens [35] and the LipL32 antigen of Leptospira interrogans [28]. In our study all rBCGpUS977dtb<sub>PW8</sub> clones stably expressed significant amounts of DTB while rBCGpUS973dtb<sub>PW8</sub> clones were unstable and, consequently, unsuitable for DTB expression. Both hsp60 and  $P_{AN}$  are constitutive mycobacterial promoters and have been widely used in recombinant BCG studies [18, 35,36]. The reason for the instability of rBCGs transformed with the pUS973 construct designed with the stronger hsp60 promoter is not yet clear and was not the objective of this study. However, it is known that the expression of heterologous antigens in recombinant BCG essentially imposes a metabolic burden with an unpredictable outcome and also, although not frequent in mycobacteria, the modification of the hsp60 promoter may result from a recombination event between host and plasmid sequences [11]. Consequently, the foreign gene expression driven by the hsp60 promoter is not always feasible in BCG [11,37]. The hsp60 promoter has a much stronger activity when compared to the  $P_{AN}$  promoter and this may result in a more stressful metabolic scenario for the host cell, which in turn may select point mutations or deletions in the hsp60 promoter region disrupting the expression of the heterologous protein. In fact, there may be a lethal level of synthesis of the foreign protein followed by selection of mutants with deletion in the promoter region immediately after induction of the hsp60 promoter in the transformed host [13]. Based on the preliminary data presented here we believe that the recombinant BCG expressing the rDTB<sub>PW8</sub> polypeptide under the  $P_{AN}$  promoter can be further exploited for the development of a vaccine prototype as an alternative for the current vaccine against diphtheria. We showed that  $P_{AN}$ promoter was expressed continuously in vitro during intracellular growth of M. bovis BCG, and basically all rBCGpUS977dtb<sub>PW8</sub> clones stably expressed the nontoxic fragment B of the diphtheria toxin. On the other hand, the pUS973, a similar expression vector based on the use of a heat shock protein promoter (hsp60) was highly unstable and inadequate for expression of the  $dtb_{PW8}$  gene.

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