Closely Related Mycobacterial Strains Demonstrate Contrasting Levels of Efficacy as Antitumor Vaccines and Are Processed for Major Histocompatibility Complex Class I Presentation by Multiple Routes in Dendritic Cells

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Mycobacteria expressing recombinant antigens are already being developed as vaccines against both infections and tumors. Little is known about how dendritic cells might process such antigens. Two different mycobacterial species, the fast-growing Mycobacterium smegmatis and the slow-growing M. bovis BCG, were engineered to express a model tumor antigen, the K^b-restricted dominant cytotoxic T-lymphocyte epitope OVA257-264. Recombinant M. bovis BCG but not recombinant M. smegmatis conferred protection to mice challenged with the B16-OVA tumor cell line. We went on to investigate whether the contrast in antitumor efficacy could be due to differences in how dendritic cells process antigen from the two mycobacterial strains for class I presentation. Both strains of mycobacteria caused phenotypic maturation of dendritic cells, but recombinant M. smegmatis infection led to a greater degree of dendritic cell maturation than recombinant M. bovis BCG infection. Antigen from recombinant M. smegmatis was processed and presented as OVA257-264 on K^{b} molecules by the dendritic cell line DC2.4 but not by bone marrow-derived dendritic cells (BMDC) or splenic dendritic cells. In contrast, antigen from recombinant M. bovis BCG was presented by all three dendritic cell types as long as the mycobacteria were viable. Such presentation was dependent on proteasome function and nascent major histocompatibility complex (MHC) class I molecules in DC2.4 cells but independent of the proteasome and transporter associated with antigen processings (TAP) in BMDC and splenic dendritic cells. These data demonstrate for the first time that antigen vectored by the slow-growing M. bovis BCG but not that vectored by fast-growing, readily destroyed *M. smegmatis* is processed and presented on MHC class I by in vitro-generated dendritic cells, which has implications for recombinant microbial vaccine development.

Mycobacterium bovis BCG, the only vaccine available against tuberculosis (22), has also been used successfully as a vaccine against leprosy (29) and, for over 20 years, as a local immunotherapy against superficial bladder cancer (2). *Mycobacterium smegmatis* is a related organism which has potential advantages as a recombinant vaccine over *M. bovis* BCG, as it is nonpathogenic and frequently commensal in humans (36, 42). Indeed, emulsified *M. smegmatis* and *M. bovis* BCG have comparable antitumor activity in a tumor immunotherapy model (56). The recent realization that mycobacteria are promising vehicles for a new generation of recombinant vaccines lends urgency to investigating the relative value of different strains in that role.

Despite their close relationship, there are several key differences between *M. bovis* BCG and *M. smegmatis. M. bovis* BCG is derived from the intracellular pathogen *M. bovis* and survives in host cells for months (31). In contrast, *M. smegmatis* is rapidly destroyed within infected cells and is nonpathogenic. Highly pathogenic mycobacteria such as *M. tuberculosis* can survive and replicate within cells because they inhibit phagosome maturation and hence their own degradation (11). Although the exact mechanism is not yet fully understood (47), it is known that such mycobacteria prevent acidification of the phagosome (13) and subsequent formation of a phagolysosome. *M. bovis* BCG shares many characteristics with *M. tuberculosis* and induces biochemical markers of maturation arrest (53, 54). On the other hand, *M. smegmatis* does not arrest phagosome maturation and is degraded rapidly by phagolysosomal proteases, which probably accounts for its lack of pathogenicity (28, 54).

Since the development of mycobacterium-*Escherichia coli* shuttle plasmids (23), there has been much interest in the generation of recombinant mycobacteria as vaccines against infectious disease, such as tuberculosis and human immunodeficiency virus. Furthermore, a recent study has also shown that recombinant *M. bovis* BCG can also protect against tumor challenge in model systems (15). The authors demonstrated that if they vaccinated mice with 10^2 or 10^4 *M. bovis* BCG secreting OVA₂₃₀₋₃₅₉ and then challenged at day 30, 120, or 150 with B16-OVA, a significant delay in tumor growth was observed.

It is generally accepted that the generation of effective antitumor and anti-infective immune responses requires antigenpresenting cells to prime cytotoxic T lymphocytes and T-helper cells by processing antigen and presenting it on MHC class I and class II molecules, respectively. Mice immunized with recombinant *M. bovis* BCG expressing human immunodeficiency virus antigens (1) and β -galactosidase (49) generate specific cytotoxic T lymphocytes, but the mechanisms underlying anti-

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gen-presenting cell processing and presentation of antigen from recombinant mycobacteria have not been elucidated.

Macrophages are well recognized as antigen-presenting cells which host mycobacteria (43). Several lines of evidence suggest that dendritic cells might be equally important processors of mycobacterial antigens. Dendritic cells are a major host cell for mycobacteria and undergo levels of infection similar to those of macrophages (25). Dendritic cells are highly specialized antigen-presenting cells and uniquely potent primers of naïve T-cell responses (4, 20). However, exactly how antigens secreted from mycobacteria reach MHC class I presentation pathways by dendritic cells has not been defined.

Antigen-presenting cell processing pathways for MHC class I and class II presentation were long thought to be distinct (24). The "classical" MHC class I pathway involves degradation of cytosolic proteins by the proteasome into MHC class I binding epitopes, whereas antigens from exogenous sources were presented on MHC class II after phagocytosis. These peptides are transported into the endoplasmic reticulum, where they bind MHC class I molecules by the transporter associated with antigen processing (TAP) (38). However, it has recently been discovered that phagocytic cells can present antigens from exogenous sources, such as extracellular bacteria (e.g., *Escherichia coli*) not only on MHC class II molecules, but also on MHC class I.

The amino acid position 257 to 264 epitope (SIINFEKL) of the model antigen OVA, expressed as a fusion protein in *E. coli* and *Salmonella enterica serovar Typhimurium*, is presented on cell surface MHC class I in macrophages (41), dendritic cells (45, 50) and neutrophils (44). In macrophages, processing of SIINFEKL from *E. coli* occurs via both the classical pathway and a TAP-independent, proteasome-independent pathway, i.e., without requiring nascent MHC class I molecules but involving the cysteine proteases which are more often employed for MHC class II processing (7). Neutrophils also appear to use a proteasome-independent pathway (44). In contrast, dendritic cells process SIINFEKL from *E. coli* via a proteasome-dependent, TAP-dependent pathway (50).

Two alternative pathways have thus been proposed for nonclassical MHC class I presentation in which antigen is either degraded in vacuolar compartments such as the phagolyosome and bound to trafficking MHC class I molecules or regurgitated from the cell whereby it competes for binding to MHC class I molecules at the cell surface (6, 10, 19, 46). The intracellular bacterium *Listeria monocytogenes* escapes the phagosome and enters the cytosol by secretion of the pore-forming protein listeriolysin O (LLO), which allows its proteins access to the classical MHC I pathway (6).

We chose to study the capacity of two very different mycobacteria, *M. bovis* BCG and *M. smegmatis*, to generate a CD8restricted T-cell response to recombinant antigen and the routes by which this antigen is presented to CD8⁺ T cells. The B16-OVA tumor model was employed to assess CD8 T-cell responses after vaccination, as this also enabled the use of the well-characterized MHC class I binding epitope OVA₂₅₇₋₂₆₄ to study antigen presentation (27, 35). Our results demonstrate for the first time that dendritic cells present recombinant antigen from slow-growing mycobacteria on MHC class I and that the nonpathogenic bacterium *M. smegmatis* is a poor candidate

A



FIG. 1. *M. bovis* BCG and *M. smegmatis* were engineered to secrete a polyepitope, OVApet. A polyepitope, OVApet, was constructed to express the OVA₂₅₇₋₂₆₄ H2-K^b binding epitope downstream of the alpha antigen secretory signal from *M. bovis* BCG, which ensured it was secreted out of the bacteria (A). Lysates of wild-type *M. bovis* BCG (1), *M. bovis* BCG-OVApet (2), *M. smegmatis* (3) and *M. smegmatis*-OVApet (4) were run on a sodium dodecyl sulfate–16% reducing polyacrylamide gel, blotted onto a nitrocellulose membrane, and probed with an anti-PK monoclonal antibody followed by a biotinylated secondary antibody and then streptavidin-horseradish peroxidase. Expression of OVApet was detected by enhanced chemiluminescence (Fig. 1B). α-Ag, alpha antigen.

for a CD8-restricted vaccine, reflecting its inability to gain access to the MHC class I processing pathway in dendritic cells.

MATERIALS AND METHODS

Generation of recombinant mycobacteria. Mycobacterium bovis BCG Pasteur strain (American Type Culture Collection) and M. smegmatis (American Type Culture Collection) were grown in Middlebrook 7H9 (Difco) supplemented with 0.1% Tween 80, 0.2% glycerol, and 10% oleic acid dextrose catalase (Difco) or on solid 7H11 (Difco) supplemented with 0.5% glycerol and 10% oleic acid dextrose catalase. The mycobacterial-Escherichia coli shuttle vector pMOD8 was a gift from M. O'Donnell (Department of Urology, University of Iowa, Iowa City, Iowa). The polyepitope OVApet was constructed by PCR amplification of the oligonucleotide (Cancer Research UK peptide synthesis laboratory) 5'-CG GGATCCGCCACCATGAGTATAATCAACTTTGAAAAACTGAACACCG ATGGGAGTACCGACTACGGAATCCTACAGATCAACAGCCGCGGCA AGCCCATCCCCAACCCCCTGCTGGGCCTGGACTCCACCTCTAGATA GGAATTCCG-3'; 20-bp primers were synthesized complementary to the 5' and 3' ends of the oligonucleotide and OVApet was amplified by PCR with Deep vent DNA polymerase (New England Biolabs). The cDNA was cloned into pMOD8 by BamHI and EcoRI restriction sites downstream of the alpha antigen secretory sequence under the control of the constitutive HSP60 promoter. The 148-bp polyepitope contains the H2-K^b OVA₂₅₇₋₂₆₄ epitope (SIINFEKL) followed by an I-Ak46-61 epitope from the irrelevant hen egg lysozyme and a PK antibody tag (Fig. 1).

pMOD8-OVApet plasmid DNA was sequenced and electroporated into *M. bovis* BCG and *M. smegmatis* (39). Recombinant clones were selected for by growth on 7H11 agar (30 µg/ml kanamycin). Single colonies were picked into 7H9 medium as above containing 30 µg/ml kanamycin and grown by rolling for 3 to 4 weeks (*M. bovis* BCG) or by shaking at 100 rpm for 2 to 3 days (*M. smegmatis*) until an optical density at 600 nm was approximately equal to 1. Clones were stored at -80° C in phosphate-buffered saline (PBS)–15% glycerol and tested for expression of OVApet by Western blotting with an anti-PK monoclonal antibody (Serotec). CFU of the frozen clones were determined by

plating serial dilutions on 7H11 plates containing kanamycin and counting resultant colonies.

Mice and immunizations. Female C57BL/6 mice aged 6 to 8 weeks old were obtained from Harlan-UK Ltd., Oxon, United Kingdom. $TAP1^{-/-}$ mice were a gift from Caetano Reis e Sousa (Cancer Research UK London Institute. London, United Kingdom).

Mycobacteria were thawed, diluted in PBS as appropriate, and 100 µl (*M. bovis* BCG-OVApet) or 200 µl (*M. smegmatis*-OVApet) was injected subcutaneously into the left inguinal region on days 0, 14, and 28. Eight mice in each group were injected with 1.5×10^8 , 1.5×10^7 or 1.5×10^6 *M. bovis* BCG-OVApet or 1.5×10^7 wild-type *M. bovis* BCG. Five mice in each group were injected with 10^8 , 10^7 or 10^6 *M. smegmatis*. On day 42, 2×10^5 B16-OVA cells (a gift from Edith Lord, University of Rochester, Rochester, N.Y.) were injected subcutaneously in PBS into the contralateral flank. Tumor growth was measured every 2 days with calipers, and mice were sacrificed when tumor size reached 1.44 cm². Kaplan-Meier curves of survival were constructed with SAS, and statistical significance was determined by log rank tests.

Maintenance of cell lines. The dendritic cell line DC2.4 was a gift from K. Rock (Dana-Farber Cancer Institute, Boston, Mass.) and was maintained in RPMI 1640 (Gibco-BRL) supplemented with 10% heat-inactivated fetal calf serum (Harlan Sera-labs), 2 mM L-glutamine, and 0.05 mM 2-mercaptoethanol. The B3Z hybridoma, which secretes interleukin (IL)-2 upon ligation of its Kb-SIINFEKL-specific receptor (27), was a gift from N. Shastri (Department of Molecular and Cell Biology, University of California, Berkeley). It was maintained in RPMI 1640 supplemented with 10% heat-inactivated fetal calf serum, 2 mM L-glutamine, 1 mM sodium pyruvate, and nonessential amino acids (Gibco-BRL). Cells were screened for mycoplasma contamination by two assays, extranuclear DNA determination with Vero African green monkey indicator cells and a stab culture technique for atypical bacterial colonies on mycoplasma agar, and were universally negative.

Generation of bone marrow-derived dendritic cells and splenic dendritic cells. Bone marrow was removed from the femur and tibia of C57BL/6 and TAP1females by flushing into RPMI. To enrich for dendritic cell precursors, cells were incubated with biotinylated antibodies to B220, I-Ab, Gr1, CD3E, and CD4 (BD-Pharmingen), washed, incubated with streptavidin microbeads, and passed through a magnetic cell separation column (Miltenvi-Biotech). Eluted cells were cultured at 37°C in RPMI 1640 supplemented with 10% fetal calf serum, 2 mM L-glutamine, 0.05 mM 2-mercaptoethanol, penicillin and streptomycin, and 10 ng/ml murine recombinant granulocyte-macrophage colony-stimulating factor. and 100 U/ml murine recombinant IL-4 (both from Biosource International). Two thirds of the medium was replaced on day 2, and on day 4 the medium was replaced with fresh antibiotic-free medium. Bone marrow-derived dendritic cells (BMDC) were used for experiments on days 6 to 8, at which point about 70% were CD11c⁺ by flow cytometry. Single-cell splenocyte suspensions were obtained by digestion of spleens from C57BL/6 and TAP1-/- females in 66.67 Wünsch U/ml Liberase CI (Roche) and 1% (wt/vol) DNase I (Roche) for 30 min at 37°C. Dendritic cells were then purified by positive magnetic selection of single-cell suspensions with CD11c⁺ microbeads (Miltenyi Biotech), which produced a CD11c⁺ purity by fluorescence-activated cell sorting analysis of about 80% CD11c. Splenic dendritic cells were used immediately.

Phenotype of bacterially infected dendritic cells. DC2.4 and BMDC (10⁶) were plated in 24-well plates and infected with *M. bovis* BCG-OVApet or *M. smegmatis*-OVApet for 2 h at an multiplicity of infection of 10. Amikacin was added to a final concentration of 200 μ g/ml for a further 2 h to kill any remaining extracellular bacteria. The medium was then changed for antibiotic-free medium, and the dendritic cells were incubated for 16 h overnight at 37°C.

Dendritic cells (10^5) were washed in PBS containing 1% fetal calf serum and 0.1% sodium azide, incubated for 15 min with a CD16/CD32 Fc γ III/II receptor antibody (BD-Pharmingen) to block nonspecific antibody binding and then incubated with phycoerythrin-conjugated antibodies to CD40, CD54, CD80, CD86, H2-K^b and I-A^b (BD-Pharmingen) or the relevant isotype control for 30 min at 4°C. Cells were washed and 10^4 events were acquired on a FACscalibur flow cytometer. (Becton Dickinson). Analysis was performed with Cell Quest Pro software (Becton Dickinson).

Antigen presentation assays. We plated 10^5 dendritic cells in $100 \ \mu$ l in 96-well plates. Dendritic cells were infected with an equal volume of mycobacteria at multiplicities of infection of 1, 10, 50, and 100 for 2 h and treated with amikacin as above. After several washes in PBS, 10^5 B3Z were added to the wells in 200 $\ \mu$ l of medium and incubated at 37°C for 16 h. Supernatants were stored at 4°C or -20° C until IL-2 was measured. In inhibitor experiments, dendritic cells were treated for 3 h with 10 $\ \mu$ M lactacystin (Calbiochem) in T25 flasks at a cell density of $10^6/m$. Cells treated with brefeldin A were incubated with 2 $\ \mu$ g/ml brefeldin A in a 100- $\ \mu$ l volume in 96-well plates for 30 min and then diluted to 1 $\ \mu$ g/ml by

addition of an equal volume of medium containing mycobacteria. As brefeldin A is reversible after 5 h, dendritic cells were fixed in 4% paraformaldehyde for 10 min at room temperature, washed in PBS, quenched in 0.1 M glycine, and washed twice in PBS before addition of B3Z to prevent further processing in the absence of brefeldin A. Statistical analysis of the effects of TAP, lactacystin, and brefeldin A on antigen presentation were performed with the paired Student *t* test.

Measurement of IL-2 in supernatants by ELISA. We coated 96-well flatbottomed Maxisorb immunoplates (Nunc) with 100 ng of anti-mouse IL-2 capture antibody (BD-Pharmingen) in 0.1 M NaHCO₃ at 4°C overnight, washed in PBS with 0.05% Tween 20, and blocked with 10% heat-inactivated fetal calf serum in PBS. Supernatants from dendritic cells, B3Z antigen presentation assays were added at a 1:5 dilution in medium in a final volume of 100 μ l. Recombinant murine IL-2 (R&D Systems) was used as a standard.

Detection of OVApet protein in bacterial supernatants. *M. bovis* BCG, *M. bovis* BCG-OVApet, *M. smegmatis*, and *M. smegmatis* OVApet were thawed from -80° C and resuspended at 8×10^{7} bacteria per ml. *M. bovis* BCG-OVApet and *M. smegmatis* OVApet were also heat killed at 80° C for 20 min; 8×10^{7} bacteria were incubated at 37° C for 4 h, and 200 µl of supernatant was added to 96-well flat-bottomed Maxisorb immunoplates. One in two serial dilutions were made of 100 µl of supernatant into 100 µl of 0.1 M NaHCO₃ coating buffer and 100 µl of coating buffer was then added to the undiluted supernatants.

OVApet peptide was used as a standard at 1 in 2 consecutive serial dilutions in coating buffer from a concentration of 1 μ M to 0.2 pM. All conditions were performed in triplicate. Samples were incubated at 4°C overnight and then blocked with 10% heat-inactivated fetal calf serum in PBS and incubated with a 1:1,000 anti-PK monoclonal antibody in blocking buffer (Serotec) for 1 h at room temperature. Following further washes, plates were incubated with a 1:500 dilution of anti-mouse immunoglobulin G-horseradish peroxidase (Serotec) and then incubated in the dark with tetramethylbenzidine peroxidase E1A substrate for approximately 5 to 10 min at room temperature. The colorimetric reaction was stopped by the addition of 50 μ l of 1 M H₂SO₄ (BDH). The absorbance at 450 nm was then read, and a standard curve was constructed with the OVApet peptide standards to determine sample OVApet concentrations.

RESULTS

Differential tumor protection mediated by *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet. A polyepitope containing the H2-K^b epitope SIINFEKL downstream of the alpha antigen secretory signal of *M. bovis* BCG under the control of the *hsp60* promoter was constructed (Fig. 1A). A PK tag was fused to the distal portion of the protein for identification. The SIINFEKL epitope was flanked on its N terminus by the antiantigen signal sequence and on its C terminus by an irrelevant I-A^k binding epitope so to generate a K^b-binding epitope, precise cleavage must occur at both the N and C termini of SIINFEKL.

Recombinant clones of *M. bovis* BCG and *M. smegmatis* were examined by nonquantitative Western blotting and confirmed to contain the OVApet polyepitope by Western blot (Fig. 1B). Secretion of OVApet peptide from *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet during a 4-h period was quantified by enzyme-linked immunosorbent assay (ELISA) and *M. bovis* BCG-OVApet was shown to secrete 25 to 150 pM OVApet per 10^7 bacteria, whereas less than 0.2 pM was detected from *M. smegmatis*-OVApet.

Mice were vaccinated with either *M. bovis* BCG-OVApet or *M. smegmatis*-OVApet and challenged with B16-OVA (Fig. 2). All doses of *M. bovis* BCG-OVApet significantly delayed onset of tumor growth compared to wild-type *M. bovis* BCG [P =0.0001 (1.5 × 10⁸), P = 0.008 (1.5 × 10⁷), or P = 0.0028 (1.5 × 10⁶)] (Fig. 2A). Median survival of controls was 18 days (95% confidence interval, 18 to 20 days) compared to 32 days (95% confidence interval, 25 to 38 days)/20 days (95% confidence interval, 20 to 32 days)/20 days (95% confidence inter-



FIG. 2. M. bovis BCG-OVApet protects against B16-OVA challenge but M. smegmatis-OVApet does not. C57BL/6 mice were vaccinated with 1.5×10^8 (\blacksquare), 1.5×10^7 (\blacktriangle), or 1.5×10^6 (\times) *M. bovis* BCG-OVApet or 1.5×10^7 wild-type *M. bovis* BCG (O) in PBS subcutaneously in the inguinal region on day 0, 14, and 28. At day 42, mice were challenged with 2×10^5 B16-OVA tumor cells subcutaneously in the opposite flank, and progression in tumor growth was measured every 2 days. Tumor growth curves for each individual mouse are shown (A). Kaplan-Meier curves for eight mice per group are shown (B). C57BL/6 mice were vaccinated with 10^8 (\blacksquare), 10^7 (\blacktriangle), or 10^6 (\times) *M.* smegmatis-OVApet or 10^7 wild-type *M.* smegmatis (O) in PBS subcutaneously in the inguinal region on days 0, 14, and 28. At day 42, mice were challenged with 2×10^5 B16-OVA tumor cells subcutaneously in the opposite flank, and progression in tumor growth was measured every 2 days. Kaplan-Meier curves for five mice per group are shown (C).

val, 18 to 27 days) for mice vaccinated with 1.5×10^8 , 1.5×10^7 or 1.5×10^6 CFU of *M. bovis* BCG-OVApet, respectively (Fig. 2B). There was no significant difference between the three doses of *M. bovis* BCG-OVApet (P = 0.249), but there was a significant increase in survival time for these three doses of *M. bovis* BCG-OVApet over wild-type *M. bovis* BCG (P = 0.0001). In contrast, *M. smegmatis*-OVApet did not afford pro-

tection against tumor challenge irrespective of dose (Fig. 2C) (P = 0.36).

Mice vaccinated with recombinant mycobacteria were also examined 2 weeks after the last vaccination for antigen-specific T-cell responses. Following a week of restimulation with irradiated EG70VA cells, specific lysis of SIINFEKL-pulsed RMA-S cells was observed in two of five, five of five, and one of five mice given 1.5×10^8 , 1.5×10^7 or 1.5×10^6 CFU of *M. bovis* BCG-OVApet, respectively, whereas no lysis was seen in mice given 10^8 , 10^7 , or 10^6 CFU of *M. smegmatis*-OVApet or wild-type bacteria (data not shown).

Subsequent experiments addressed the effects of the respective mycobacteria on dendritic cells and how their secreted antigen gains access to the MHC class I of dendritic cells.

Both recombinant M. bovis BCG and M. smegmatis efficiently mature dendritic cells. Dendritic cells took up either mycobacterial strain after 2 h of exposure (data not shown). Infection of both the bone marrow dendritic cell-derived cell line DC2.4 and BMDC with M. smegmatis-OVApet led to maturation of the dendritic cells as determined by upregulated surface expression of CD40, CD54, CD80, CD86, and MHC class II molecules (Fig. 3). However, although M. bovis BCG-OVApet led to maturation of BMDC, only a small increase in maturation markers on DC2.4 cells were observed following M. bovis BCG-OVApet infection. Maturation of BMDC was also less than that observed with M. smegmatis- OVApet. Maturation of splenic dendritic cells was not observed. However, splenic dendritic cells are very short lived, and 20 h after infection, 80 to 90% of the dendritic cells were apoptotic (data not shown). Both strains of mycobacteria upregulated MHC and costimulatory molecules in peritoneal macrophages, although expression levels were considerably lower than seen with dendritic cells (data not shown). Therefore, recombinant *M. bovis* BCG and M. smegmatis are capable of inducing maturation of murine dendritic cells, but M. smegmatis is a more potent inducer of maturation of dendritic cells than M. bovis BCG.

Differential processing of MHC class I-restricted epitope when vectored by M. bovis BCG and M. smegmatis. Having confirmed that dendritic cells undergo maturation when infected with M. bovis BCG or M. smegmatis, we investigated antigen processing of the K^b-binding epitope secreted by recombinant mycobacteria-OVApet. Control experiments where dendritic cells were pulsed with SIINFEKL peptide and used to stimulate the B3Z hybridoma showed that the three types of dendritic cells bound and presented SIINFEKL peptide with IL-2 secretion from the hybridoma, plateauing at 10 µM to 100 nM peptide. However, splenic dendritic cells were the most potent presenters of bound peptides, with half maximal stimulation of B3Z being obtained with only 10 pM peptide. BMDC required 500 pM peptide to stimulate half the B3Z cells and DC2.4 cells 1 nM peptide (Fig. 4.A). Comparisons were then made between the peptide and the presentation of SIINFEKL following infection of dendritic cells with recombinant mycobacteria.

The DC2.4 cell line processed SIINFEKL from both *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet, although SIINFEKL was processed somewhat more efficiently from *M. bovis* BCG (Fig. 4B). BMDC also processed SIINFEKL from *M. bovis* BCG-OVApet, though much less efficiently than DC2.4 cells (Fig. 4C). Processing of SIINFEKL by BMDC





FIG. 3. *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet induce a mature dendritic cell phenotype. We infected 10⁶ dendritic cells at a multiplicity of infection of 10 with *M. bovis* BCG-OVApet (grey hashed line) or *M. smegmatis*-OVApet (bold line) or left them uninfected (thin line) for 2 h. The remaining extracellular bacteria were killed with antibiotic treatment, and the dendritic cells were cultured for 16 h. Dendritic cells were incubated with phycoerythrin-conjugated antibodies to CD40, CD54, CD80, CD86, H2-K^b and I-A^b and analyzed by flow cytometry. DC2.4 cell surface expression (A) and BMDC cell surface expression (B) are shown.

following *M. smegmatis*-OVApet was only observed infection in 3/8 experiments. Splenic dendritic cells (CD11c⁺) processed SIINFEKL vectored by *M. bovis* BCG-OVApet (Fig. 4D). However, unlike DC2.4 and BMDC, splenic dendritic cells did not process and present SIINFEKL from *M. smegmatis*-OVApet.

Although SIINFEKL was processed and presented from *M. smegmatis*-OVApet, presentation was not always observed in dendritic cells generated directly from bone marrow precursor cells in vitro and never by splenic dendritic cells. Further controls which included comparison of live and nonviable *M. bovis*

BCG-OVApet (Fig. 5A and C) and *M. smegmatis*-OVApet (Fig. 5B and D) showed that only epitopes secreted from live mycobacteria were processed and presented on MHC class I in both the DC2.4 cell line (Fig. 5A and B) and BMDC (Fig. 5C and D). There was no presentation of SIINFEKL from *M. bovis* BCG-OVApet or *M. smegmatis*-OVApet by peritoneal macrophages (thioglycolate elicited) or by the peritoneal macrophage cell line IC-21 (data not shown).

Presentation of SIINFEKL is only partly proteasome and TAP dependent. We next sought to determine the extent to which the classical MHC class I antigen-processing pathway



FIG. 4. Antigen presentation from recombinant mycobacteria by dendritic cells. DC2.4 (\Box), BMDC (\blacksquare), and splenic dendritic cells (\blacktriangle) were pulsed with increasing concentrations of SIINFEKL peptide for 2 h, washed, and then cultured with B3Z for 16 h. IL-2 production in supernatants was measured (A). Various dendritic cells were infected with increasing amounts of *M. bovis* BCG (\blacklozenge), *M. bovis* BCG-OVApet (\blacksquare), *M. smegmatis* (\bigstar), or *M. smegmatis*-OVApet (\Box) and cultured with the B3Z hybridoma for 16 h. Presentation of SIINFEKL on H2-K^b was measured by IL-2 secretion. The dendritic cell line DC2.4 (B), bone marrow-derived dendritic cells (C), and CD11c⁺ splenic dendritic cells (D) are shown.

was involved for processing antigen from *M. bovis* BCG and *M. smegmatis* by various dendritic cell types. To dissect the pathway by which the SIINFEKL epitope from recombinant mycobacteria is delivered to MHC class I, we used lactacystin, an irreversible proteasome inhibitor; brefeldin A, which blocks the export of newly synthesized MHC class I molecules from the *trans*-Golgi network; and *TAP1*-deficient ($TAP1^{-/-}$) mice

lacking a functional TAP transporter. After establishing that treatment with lactacystin did not diminish presentation of SIINFEKL, presentation from recombinant mycobacteria in dendritic cells treated with lactacystin was studied. Lactacystin reduced presentation of SIINFEKL by DC2.4 cells by 60% (*M. bovis* BCG-OVApet) and by 30% (*M. smegmatis*-OVApet) (Fig. 6A). A more pronounced reduction in SIINFEKL presentation was achieved by blocking export of nascent MHC class I molecules for *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet (Fig. 6B).

Lactacystin did not inhibit presentation of SIINFEKL from *M. bovis* BCG-OVApet by infected BMDC (Fig. 7A). Indeed, presentation increased following lactacystin treatment (two of four experiments) and was unaffected in the remainder. BMDC from $TAP1^{-/-}$ mice retained some ability to process and present SIINFEKL from *M. bovis* BCG-OVApet but with a significant decrease, but not abolishment, in presentation in two of four experiments (Fig. 7B). However, in two further experiments no effect on presentation was observed in BMDC from $TAP1^{-/-}$ mice (Fig. 7C).

Control experiments indicated that SIINFEKL peptide was presented with nearly equal efficiency on BMDC that had been ptreated with lactacystin (Fig. 8A) or derived from $TAP1^{-/-}$ mice (Fig. 8B) compared to C57BL/6 mice. In splenic dendritic cells, presentation of SIINFEKL by *M. bovis* BCG-OVApet-infected cells was also independent of the TAP transporter complex. Indeed, presentation appeared greater on $TAP1^{-/-}$ CD11c⁺ splenic dendritic cells than on splenic dendritic cells from wild-type C57BL/6 mice (Fig. 7D). Control experiments in which splenic dendritic cells were pulsed with SIINFEKL peptide indicate that $TAP1^{-/-}$ dendritic cells may in fact have a higher affinity for SIINFEKL peptide than C57BL/6 dendritic cells (Fig. 8C).

DISCUSSION

We have shown that antigen delivered to professional antigen-presenting cells by recombinant M. bovis BCG is processed and presented on MHC class I molecules, whereas antigen from recombinant M. smegmatis is only processed and presented by dendritic cells sometimes and at much lower levels than that from recombinant M. bovis BCG. We have also shown that in a tumor challenge model, recombinant M. bovis BCG afforded protection whereas M. smegmatis did not. Additionally, SIINFEKL-specific cytotoxic T lymphocytes could be detected following in vitro restimulation after vaccination with M. bovis BCG-OVApet but not M. smegmatis-OVApet. M. bovis BCG-OVApet appears to secrete as much as 750-fold more OVA-pet than M. smegmatis-OVApet, which may impact significantly on the efficacy of the vaccine. We believe this is primarily through the resultant failure of dendritic cells, which are either infected with M. smegmatis-OVApet or have taken up OVApet from apoptotic infected macrophages and dendritic cells, to present peptide antigen on MHC class I and generate an antigen-specific CD8⁺ T-cell response.

Live wild-type *M. smegmatis* has potent nonspecific immunostimulatory properties and is as effective as live *M. bovis* BCG in a preclinical model of nonspecific cancer immunotherapy (data not shown). Indeed, we have shown that *M. smegmatis* induces a higher expression of costimulatory molecules



FIG. 5. SIINFEKL from heat-killed *M. bovis* BCG-OVApet and *M. smegmatis*-OVApet is not presented on MHC class I. We infected 10^5 DC2.4 (A and B) or BMDC (C and D) in 96-well plates with live (solid squares) or heat-killed (open circles) *M. bovis* BCG-OVApet (A and C) or *M. smegmatis*-OVApet (B and D) at a multiplicity of infection of 1, 10, 50, or 100 for 2 h at 37°C. The remaining extracellular bacteria were killed by a 2-h treatment with 200 µg of amikacin per ml at 37°C. Dendritic cells were then washed in PBS and cocultured with 10^5 B3Z cells for 16 h at 37°C. Supernatants were assayed for IL-2 by solid-phase ELISA.

on dendritic cells than does M. bovis BCG. It is possible that the increased maturation of dendritic cells observed with M. smegmatis is due to structural differences in the cell wall of fastand slow-growing mycobacteria which may be related to pathogenicity (9, 14, 30). Although all three "types" of dendritic cells studied processed SIINFEKL from M. bovis BCG-OVApet, only the DC2.4 cell line processed antigen from recombinant M. smegmatis efficiently. Other differences between the three dendritic cell types were found with the DC2.4 cells processing SIINFEKL in a proteasome-dependent manner requiring newly synthesized MHC class I molecules, whereas processing in BMDC and splenic dendritic cells was independent of the proteasome and partly independent of TAP. These results highlight differences or limitations in the use of antigen-presenting cell lines for the study of antigen processing by dendritic cells.

Processing and presentation of SIINFEKL from *M. bovis* BCG required viable bacilli. This is in agreement with previous studies showing concurrent loading of macrophages with live *M. tuberculosis* and OVA protein which resulted in TAP-dependent processing of SIINFEKL (32, 37). This may suggest that *M. tuberculosis* secretes factors that facilitate antigen access to the cytosol. Slow-growing mycobacteria survive within cells, and pore forming in the phagosomal membrane may be an essential mechanism whereby nutrients from the cytosol are obtained or toxins released, as in the case of *Listeria monocy-togenes* (40, 48). This may be one mechanism whereby antigen from *M. bovis* BCG-OVApet is presented more efficiently than from *M. smegmatis*-OVApet, as a fast-growing bacterium, rapidly destroyed in the phagolysosome, would not require access to the cytosol.

One possible reason for the lack of presentation of antigen from M. smegmatis in BMDC and splenic dendritic cells appears to be that no detectable secretion of OVApet is observed by ELISA compared to that from M. bovis BCG, as secreted antigen from mycobacteria has been shown to be the most immunogenic form (18, 21). However, it cannot be ruled out that M. smegmatis may upregulate its secretion of OVApet after uptake by dendritic cells, particularly by DC2.4 cells. The OVApet protein is under the control of the HSP60 promoter, and stresses within the phagosome may upregulate transcription and translation of OVApet and hence secretion, as shown for Mycobacterium avium LacZ (5). As proteasome-dependent processing is observed from M. smegmatis OVApet in DC2.4 cells, this indicates that either this route of presentation is very efficient in the cell line and extremely low levels of secreted OVApet are required for generation of SIINFEKL or that OVApet peptide does not need to be secreted from the bacteria for degradation to SIINFEKL by the proteasome.

Antigen processing of SIINFEKL from OVApet appeared



FIG. 6. Presentation of SIINFEKL from recombinant bacteria in DC2.4 cells is reduced by the proteasome inhibitor lactacystin and the inhibitor of retrograde transport brefeldin A. We infected 10^5 DC2.4 in 96-well plates with *M. bovis* BCG-OVApet or *M. smegmatis*-OVApet at a multiplicity of infection of 1, 10, 50, or 100 for 2 h at 37°C or pulsed them with SIINFEKL peptide. The remaining extracellular bacteria were killed by a 2-h treatment with 200 µg of amikacin per ml at 37°C. Dendritic cells were then washed in PBS and cocultured with 10^5 B32 cells for 16 h at 37°C. Supernatants were assayed for IL-2 by solid-phase ELISA. In some instances dendritic cells were treated with 10 µM lactacystin for 3 h (A) or 1 or 5 µg of brefeldin A per ml for 30 min (B). Dendritic cells were fixed in 1% paraformaldehyde and washed extensively before addition of B32 (B). A statistically significant decrease in presentation is indicated by * (P < 0.05, Student *t* test).

to be partly independent of TAP in BMDC. This is in agreement with Neyrolles et al., who showed that processing by macrophages of the *M. tuberculosis* 19-kDa antigen from *M. vaccae*, *M. smegmatis*, and *M. bovis* BCG was independent of TAP (37). Presentation of a class I-restricted NP epitope, fused to the 19-kDa antigen, in fast-growing *M. vaccae* and *M. smegmatis* was found to be more efficient than in *M. bovis* BCG and correlated with enhanced phagosome acidification. Where our findings contrast with those of Neyrolles et al. is our presentation of SIINFEKL is greater by dendritic cells infected with recombinant *M. bovis* BCG and much less in those infected by recombinant *M. smegmatis*.

As an additional control, presentation from *E. coli* LLO OVA (45) was studied as a control, as the presence of LLO enables OVA to gain access to the cytosol. As expected, presentation from *E. coli* LLO OVA was abolished in $TAP1^{-/-}$ dendritic cells (data not shown). However, incomplete inhibition of presentation was seen following treatment with lactacystin in BMDC, although presentation was abolished in DC2.4 cells (data not shown). Although this suggests that 10 μ M lactacystin was insufficient to completely inhibit proteasome function in BMDC, a reduction in presentation from *E.*

coli LLO OVA was always seen. This is in contrast to presentation from *M. bovis* BCG-OVApet, which was increased in two of four experiments and unaffected in the remainder following lactacystin treatment of dendritic cells.

The presentation seen in BMDC and splenic dendritic cells is likely to be via alternative pathways (26) where antigen is degraded in the phagosome and binds to recycling MHC class I molecules in post-Golgi compartments or is regurgitated from the cell (46). In support of this Canaday et al. found that processing of *M. tuberculosis* antigens to $CD8^+$ T cells was insensitive to brefeldin A (8). Indeed, it has recently been shown that parts of the phagosomal membrane appears to be derived from the membrane of the endoplasmic reticulum which may explain the presence of MHC class I molecules within the phagosome and alternative antigen processing (16). In contrast to the work presented here, presentation of SIINFEKL from *E. coli* and *S. enterica serovar Typhimurium* expressing the fusion protein Crl-OVA is TAP independent in macrophages but TAP dependent in dendritic cells (51, 55).

The fusion protein Crl-OVA contains the OVA₂₅₇₋₂₆₄ epitope internal to the fusion protein, but if recombinant *E. coli* expressing full-length native OVA were used to infect



FIG. 7. MHC class I presentation of SIINFEKL is not inhibited by lactacystin and is partly TAP independent in BMDC and splenic dendritic cells. We treated 10^5 BMDC for 3 h with 10 μ M lactacystin or left them untreated and then infected them in 96-well plates with *M. bovis* BCG-OVApet at a multiplicity of infection of 1, 10, 50, or 100 for 2 h at 37°C or pulsed them with SIINFEKL peptide. The remaining extracellular bacteria were killed by a 2-h treatment with 200 μ g of amikacin per ml at 37°C. Dendritic cells were then washed in PBS and cocultured with 10^5 B3Z cells for 16 h at 37°C. Supernatants were assayed for IL-2 by solid-phase ELISA (A). We infected 10^5 BMDC (B and C) or splenic dendritic cells (D) from C57BL/6 or *TAP1^{-/-}* mice with *M. bovis* BCG-OVApet or pulsed them with SIINFEKL peptide, and presentation of SIINFEKL to B3Z cells was determined as described above. A statistically significant decrease or increase in presentation following lactacystin treatment or the use of *TAP1^{-/-}* dendritic cells is indicated by * (P < 0.05, Student *t* test).

macrophages, TAP-dependent processing was seen (55). The polyepitope in our study contained the $OVA_{257-264}$ K^b-binding epitope in the absence of the native N and C termini, which normally flank this epitope. As the proteasome is largely responsible for correct cleavage of C-terminal amino acids from MHC class I epitopes (12, 33), substitution of flanking residues may alter the necessity for proteasomal cleavage and the resultant pathway and efficiency of presentation (34). However, as only the N-terminal flanking residue differs between OVApet and Crl-OVA, this indicates that both N- and C-terminal residues may affect proteasome-dependent processing in dendritic cells and macrophages. Alternatively, the different mechanisms that the bacteria use to inhibit phagoly-

sosome fusion and gain access to the cytoplasm may affect presentation pathways (3).

The final part of our work explored the use of recombinant mycobacteria as in vivo antitumor vaccines. One study with recombinant mycobacteria as an antitumor vaccine warrants consideration in the context of our work. Recently, Dudani et al. (15) studied recombinant *M. bovis* BCG and showed a similar delay in tumor growth. Both their and our studies used a secretary signal to ensure secretion of the SIINFEKL-containing polypeptide into the phagosome, as this has been described as one of the best forms of antigen display for generation of immune responses (18, 21). However they used the intravenous route of *M. bovis* BCG administration, which, al-



FIG. 8. Lactacystin-treated and $TAP1^{-/-}$ dendritic cells are able to present SIINFEKL peptide with similar efficiencies. We pulsed 10⁵ C57BL/6 BMDC that had been treated with 10 uM lactacystin (**II**) or not (×) with increasing concentrations of SIINFEKL for 2 h and washed and cocultured them with 10⁵ B3Z for 16 h at 37°C. IL-2 secretion from B3Z in supernatants is shown (A). C57BL/6 BMDC (**II**) and $TAP1^{-/-}$ BMDC (**A**) were pulsed with increasing concentrations of SIINFEKL, washed, and cocultured with B3Z cells for 16 h. IL-2 secretion from B3Z in supernatants is shown (B). C57BL/6 splenic dendritic cells (**II**) and $TAP1^{-/-}$ splenic dendritic cells (**A**) were pulsed with increasing concentrations of SIINFEKL, washed, and cocultured with B3Z cells for 16 h. IL-2 secretion from B3Z in supernatants is shown (C).

though it gives direct access of vaccine to the periphery, may not be appropriate in immunocompromised patients, as a tuberculoid type disease may result (17, 52). Subcutaneous and intradermal routes of vaccination, as used in our model, may represent a safer route for use of recombinant BCG as a tumor vaccine.

Dudani et al. also showed that a large proportion of the cytotoxic T lymphocyte response to SIINFEKL was lost in CD4-deficient mice vaccinated with M. bovis BCG-OVA. However, although the M. bovis BCG-OVA construct consisted of OVA₂₃₀₋₃₅₉, encompassing both the K^b-binding and the I-A^b epitope 323 to 339, tumor protection in CD4-deficient mice was not studied. Although we were able to demonstrate only small increases in SIINFEKL-specific lysis in 5 to 10% in mice vaccinated with M. bovis BCG-OVApet but no increase in mice vaccinated with M. smegmatis-OVApet, no SIINFEKL-tetramer-specific CD8⁺ T cells were observed following restimulation. However, the inclusion of IL-2 in the restimulation culture, as others have done, may have increased the expansion and detection of SIINFEKL-specific cytotoxic T lymphocytes (15). However, it must be noted that in contrast to the work of Dudani et al., our M. bovis BCG-OVApet was specifically designed to lack specific MHC class II epitopes (from OVA or

tumor), thus excluding a role for vaccine-induced specific CD4⁺ reactivity.

In summary, it appears that antigen presentation from recombinant mycobacteria onto MHC class I by dendritic cells occurs by multiple pathways which may be both intra- and extracellular and which are most efficient when antigen is secreted from the bacteria. Fast-growing, nonpathogenic mycobacteria, like *M. smegmatis*, may not be suitable for antigenspecific anticancer vaccines in vivo as they are unable to present sufficient antigen on MHC I to stimulate an antigenspecific CD8-restricted T-cell response. Slow-growing mycobacteria such as *M. bovis* BCG, attenuated to be nonpathogenic yet retaining many of the features of pathogenic organisms, are attractive vaccine candidates as they can present their recombinant antigen to CD8⁺ T cells by several routes within dendritic cells.

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