

Experimental *Mycobacterium leprae* Infection in BALB/c Mice: Effect of BCG Administration on TNF- α Production and Granuloma Development¹

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Leprosy is a chronic mycobacterial infection that presents a spectrum of clinical manifestations in which resistance and pathogenesis are mediated by the cell-mediated immune response of the host (²²). At one extreme of the spectrum, patients with tuberculoid leprosy demonstrate a resistance response characterized by the formation of epithelioid-cell granulomas that restrict the growth of *Mycobacterium leprae*. At the opposite extreme of the spectrum, patients with lepromatous leprosy lack the T-cell-mediated immune response, and have diffuse lesions containing macrophages with large numbers of mycobacteria.

Many attempts to protect against *M. leprae* infection have been made with *M. bovis* BCG. Several studies have shown that BCG vaccination in humans induced a protective immune response against *M. leprae* infection (^{5,7}). BCG was also shown to induce a strong nonspecific immune stimulatory action against several pathogens and tumor

cells in bladder cancer (^{26,27}). In addition, it has been demonstrated that immunotherapy carried out with viable BCG and heat-killed *M. leprae* led to clinical and bacteriological improvement of lepromatous leprosy and induction of granuloma formation (^{19,20}). Moreover, changes in the immune response that alter the clinical status in leprosy patients, during reactional states are often observed during the natural course of leprosy. However, this acquired immune response against mycobacterial antigens can also be associated with immunopathologically mediated tissue damage with rapid and severe neuritis. The occurrence of reactions indicates that immunomodulatory agents could be an important therapeutic tool in the regulation of cell-mediated immunity in leprosy, although the relationship between immunopathological response and protective bactericidal immune response has not been completely elucidated.

Intracellular mycobacterial elimination and granuloma formation require the differentiation of macrophages into epithelioid cells, the contribution of CD4 and CD8 T lymphocytes, and the production of cytokines such as interferon gamma (IFN- γ) and tumor necrosis factor-alpha (TNF- α) (⁶). Accordingly, Kindler, *et al.* (¹¹) have demonstrated that administration of anti-TNF- α antibody in BCG-infected mice prevented the development of epithelioid granulomas that resulted in mycobacteria spreading. In addition, previous studies at our laboratory demonstrated that administration of thalidomide, a selective inhibitor of TNF- α synthesis (²⁵), reduced the number and size of granulomas during BCG infection in mice (¹).

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The process of apoptosis seems to correlate with the elimination of bacteria in phagocytes (15). Apoptosis is an important additional mechanism for regulating the immune response and is partly modulated by TNF- α (16, 29). Interestingly, morphological findings consistent with apoptosis have been seen in the epithelioid cells of tubercloid leprosy, sarcoidosis, and tuberculosis granulomas (9).

The experimental model of *M. leprae* infection in mouse foot pads developed by Shepard (28) has been used to investigate several aspects of the pathogenesis, immunology, and therapeutics of leprosy in BALB/c mice. *M. leprae* only grows in the foot pad in limited amounts (10^6 – 10^7), from 6 to 7 months subsequent to infection. The formation of granulomas *in situ* resembling those found in human lepromatous leprosy is then observed, followed by a complete clearance of the bacteria 12–13 months thereafter (21, 28). In a recent study, inhibition of *M. leprae* multiplication in the foot pad was demonstrated after immunizing mice with the ribosomal fraction from ruptured BCG and its culture filtrate (14).

In this study, the same classical experimental model for *M. leprae* infection was used to investigate the effect of BCG administration on TNF- α production and granuloma development in mice previously infected with *M. leprae*. It was observed that co-infection with BCG interfered with TNF- α production locally and systemically, induced the formation of epithelioid granulomas and increased both the number of apoptotic cells at the infection site as well as the rate of *M. leprae* clearance in the mouse foot pad.

MATERIALS AND METHODS

Animals. BALB/c female mice, 6 to 8 weeks of age, were used. The animals were divided into three different groups: Group I = Mice whose foot pads were infected with *M. leprae*; Group II = Mice intravenously infected with BCG; Group III = Mice infected with *M. leprae* that were intravenously administered BCG at the peak of *M. leprae* infection (month 7). The mice were housed in plastic cages and had unlimited access to feed and tap water.

Necropsy procedure. Mice were ether anesthetized, killed by cervical displace-

ment, and autopsied at predetermined intervals during the course of *M. leprae* or BCG infection. The feet of *M. leprae*-infected mice were removed and the foot pads dissected. The left foot pads were fixed in 10% formaldehyde for histopathologic analysis or frozen in liquid nitrogen for immunohistochemistry to determine the presence of TNF- α protein at the site of *M. leprae* infection. The right foot pads were used for determination of *M. leprae* growth during the course of infection. The abdominal cavities were opened aseptically, and the liver and spleen were dissected. Livers were fixed in 10% formaldehyde for histopathologic analysis. Spleens were used for BCG colony forming units (BCG-CFU) counts or for reverse transcription polymerase chain reaction (RT-PCR) analysis.

***M. leprae* infection.** The experimental *M. leprae* infection in mice was performed by the methods described by Shepard (28). Human leproma-derived *M. leprae* were used as the inoculum. A 6-mm punch biopsy was obtained from lepromatous leprosy patients and used for *M. leprae* isolation. Only patients who had given their informed consent were enrolled. Skin biopsies were minced with scissors and homogenized in Hanks' balanced salt solution (HBSS). *M. leprae* recovered from each biopsy was stained by the Ziehl-Nielsen method and counted microscopically under a 1250 \times magnification with oil immersion in 60 microscopic fields. A suspension containing 5×10^3 acid-fast bacilli (AFB) in 0.03 ml of sterile saline was injected into each mouse foot pad. *M. leprae* growth was determined at monthly intervals after infection in four animals per time point in two different experiments. The mouse foot pad tissues were minced with sharp scissors, placed in the cup of a Mickle tissue disintegrator containing glass beads, and 2 ml of sterile isotonic saline solution was added. One min of vibration was used with an amplitude of 5 mm. The remaining clumps of tissue were removed by allowing the preparation to settle in a test tube. Ten μ l of each suspension was transferred to the circles of a Reich counting slide (Bellco Glass Inc., Vineland, New Jersey, U.S.A.). Each sample was spread with a platinum wire over the surface of a circle. The slides were air dried, fixed by exposure to forma-

lin vapor, and Ziehl-Nielsen stained. AFB counts were performed as described (1250× magnification with oil immersion).

BCG infection. Mice were intravenously injected in the dorsal tail vein with 1×10^6 colony forming units (CFU) of viable BCG strain Pasteur 10/11 suspended in 0.2 ml of sterile isotonic saline solution. At weekly intervals after BCG infection, BCG-CFU enumeration in the spleen was performed (four animals per time point). Immediately after autopsy, the spleens were homogenized separately in 1 ml sterile saline solution in a teflon-coated tissue homogenizer (Glas-Col Apparatus Co., Terre Haute, Indiana, U.S.A.). Serial 10-fold dilutions were made in saline and plated onto Middlebrook 7H10 Bacto Agar (Difco Laboratories, Detroit, Michigan, U.S.A.) plates. Appropriate dilutions were tested in sextuple. Plates were sealed in plastic bags and incubated at 37°C. The BCG-CFU were counted visually after 14 days of incubation. Group III mice, previously infected with *M. leprae*, were administered BCG 7 months after *M. leprae* inoculation. In this study, *M. leprae* took an average of 7 months to reach the peak of *M. leprae* infection associated with the presence of inflammatory infiltrate in the mouse foot pads.

Histopathology and immunostaining. Paraffin sections of formalin-fixed, left mouse foot pads and liver were stained by hematoxylin and eosin (H&E) for histopathological analysis. The number of *M. leprae*-induced granulomas in the mouse foot pads and BCG-induced granulomas in the liver were counted on 100 microscope fields at 100× magnification. The extension of the inflamed tissue area and the number of TNF- α positively stained cells in the granuloma in the mouse foot pad were investigated in *M. leprae*-infected mice (N = 4) and *M. leprae* + BCG mice (N = 4) 21 days after BCG inoculation (peak of BCG infection). Two paraffin sections (5- μ m thick) of formalin-fixed tissue were obtained at varying depths from each tissue block. All microscopic fields presenting inflammatory infiltrate were selected, and quantitative measurements (%) of the inflamed area in the mouse foot pads were performed with a Mini-Mop analysis system (Kontron, Germany) during microscopic examination (100× magnification) in

four mice from each group. Quantitative measurements were expressed as a percent of the inflamed tissue area by tissue section. The presence of TNF- α protein in the *M. leprae*-induced granulomas in the mouse foot pad was investigated by immuno-histochemistry. Frozen sections (5- μ m thick) of foot pad tissues were fixed in acetone. The avidin-biotin-peroxidase complex (ABC) procedure was performed. In brief, the sections were incubated with polyclonal rabbit anti-TNF- α antibody (Genzyme, Boston, Massachusetts, U.S.A.) followed by a goat anti-rabbit IgG biotinylated antibody (Dakopatts, Copenhagen, Denmark). Controls for the ABC procedure were performed by replacing anti-TNF- α antibody with normal rabbit serum, or by omitting the anti-TNF- α antibody. Quantitative measurements were expressed as a percent of TNF- α positively stained cells in the granuloma. The sections were analyzed by light microscopy, and photomicrographs were taken with a Nikon Microphot system.

In situ detection of apoptosis. Paraffin sections (5- μ m thick) of the mouse foot pad tissues were placed on siliconized slides (Perkins-Elmer Cetus, Emeryville, California, U.S.A.). Detection of apoptotic cells was investigated by direct immunoperoxidase of digoxigenin-labeled genomic DNA in the foot pads of *M. leprae*-infected mice (N = 4) and *M. leprae* + BCG mice (N = 4) 21 days after BCG inoculation. The reaction procedure was performed according to the Apoptag Plus *in situ* detection kit (Oncor, Inc., Gaithersburg, Maryland, U.S.A.). Briefly, residues of digoxigenin-nucleotide were added to DNA by terminal deoxynucleotidyl transferase (TdT). The incorporated nucleotides formed a random heteropolymer, in a ratio that had been optimized for anti-digoxigenin antibody conjugated with peroxidase binding. Diluted 30% H₂O₂ 1:10 in methanol was used to perform the inactivation of endogenous peroxidase. Diaminobenzidine (DAB) reacted with the labeled cells to generate a colored substrate at the site of DNA fragmentation. Counterstaining with methyl green was performed and the morphology as well as DAB staining were used to interpret the results. Control of the procedure was obtained by using histological sections of rat mammary glands (positive control) or

by omitting the anti-digoxigenin antibody (negative control). Quantitative measurements were expressed as a percent of apoptotic cells in the granuloma.

TNF- α measurement. Serum TNF- α levels were investigated in the different groups of mice at the peak of *M. leprae* infection (month 7), and weekly after BCG infection. For serum collection, mice were ether anesthetized and bled by orbital puncture. Blood was allowed to clot at room temperature, and serum was aliquoted and stored at -70°C until use. The TNF- α concentration in the serum samples was determined by using a TNF- α -specific ELISA (Endogen Inc., Boston, Massachusetts, U.S.A.) as recommended by the manufacturer. Each sample was tested in duplicate, and the detection limit of the assay was 4 pg/ml.

RT-PCR analysis. TNF- α mRNA expression in the spleen was investigated in different groups of mice at the peak of *M. leprae* infection (month 7) and at 7, 14, 21, and 28 days after BCG infection. Healthy mice were used as controls in the experiment. Spleens of mice were collected and stored in liquid nitrogen until used. Fragments (25 mg) of spleen were homogenized in 3 ml Trizol (GIBCO BRL, Gaithersburg, Maryland, U.S.A.) and RNA was extracted and purified according to the manufacturer's instructions. Purified RNA was analyzed by 1.2% formaldehyde-agarose electrophoresis gel for the evaluation of RNA integrity. Following quantitation, 1 μg of total RNA was reverse transcribed into cDNA as previously described⁽²³⁾. Cytokine-specific oligonucleotide primer pairs for mouse TNF- α were kindly donated by Dr. G. M. B. Pereira (Leprosy Laboratory, Oswaldo Cruz Foundation, Brazil) and those for β -actin were purchased (Stratagene Cloning System, La Jolla, California, U.S.A.). Primer sequences (5' and 3') were as follows: TNF- α , AGAAAAGCAAGCAGCCAACCAGGCA, GGGGGCTCTGAGGAGTAGACAATAA; β -actin, TGTGGCCGCTCTAGGCACCA, CGGTTGGCCTTAGGGTTCAGGGGGG. PCR conditions were performed as indicated⁽²³⁾, and the samples amplified in a DNA thermocycler 480 (Perkin-Elmer) for 35 cycles of denaturation at 94°C for 45 sec, annealing at 60°C for 45 sec, and extension at 72°C for 1.5 min. Following

electrophoresis on 1.7% agarose gels, PCR products were transferred to Hybond-N nylon membranes (Amersham Corp., Arlington Heights, Illinois, U.S.A.), and hybridized with a radioactive oligonucleotide probe complementary to sequences internal to those recognized by the specific primers. Sequences of the probes were: TNF- α , AAACCCTGGTATGAGCCCA; β -actin, GGTGGGAATGGGTCAGAAGG. The primers were RNA specific in that both the 5' sense and 3' antisense primers spanned the junctions of two exons, thus precluding amplification of genomic DNA. Densitometer analysis was performed by scanning the images from autoradiographs using an imaging densitometer (Bio-Rad Laboratories, Hercules, California, U.S.A.). Densitometric values for each band were established by Molecular Analyst Software 1.2 (Bio-Rad Laboratories). The relative amount of PCR product present in an individual sample was expressed as a percentage relative to the most intense band, that band being assigned the value of 100. To make sure that the samples contained identical quantities of cDNA, β -actin was used to normalize the exact levels of input cDNA present among the different samples tested.

Statistical analysis. Data on *M. leprae* counts in mouse foot pads, BCG-CFU in the spleens, and TNF- α serum levels were compared using the Mann-Whitney two-sample rank sum test. The number of granulomas in the foot pads, TNF- α positively stained cells (%), apoptotic cells (%), and quantitative measurements (%) of the inflamed tissue area were compared using the Student's *t* test. The significance level adopted was $p < 0.05$. Both S.D. (standard deviation) and S.E.M. (standard error of the mean) were used to express variance.

RESULTS

Induction of *M. leprae* clearance by BCG infection. The number of *M. leprae* (AFB staining) in the mouse foot pads was investigated during the course of infection. Similar to what was observed in Shepard's landmark study (1960), the growth of *M. leprae* in these animals (BALB/c mice, Group I) was very slow. The resulting limited infection after inoculation of 5×10^3

microorganisms was preceded by a lag phase of approximately 60 days, followed by a logarithmic growth (exponential phase) which, after 6–7 months, yielded 10^6 – 10^7 *M. leprae* per foot pad (peak of infection). This was followed by a stationary phase (plateau phase) in which the number of bacteria remained more or less constant (between 7 to 10 months of infection) until the total clearance of *M. leprae* observed from 12 to 13 months later (Fig. 1). Interestingly, in Group III mice BCG inoculation at the peak of *M. leprae* infection led to an accelerated clearance of *M. leprae* in the mouse foot pad. The plateau phase was not observed in these animals, and complete clearance of *M. leprae* took place at month 11.

Effect of BCG infection on *M. leprae*-induced granulomas. At the peak of *M. leprae* growth (Group I), a diffuse mononuclear cell infiltrate associated with granulomas could be seen in the foot pads. Two histological types of granulomas were then observed. Small granulomas were formed by the accumulation of macrophages and lymphocytes (Fig. 2A); whereas epithelioid granulomas presented a well-developed structure in which central mononuclear cells showed high levels of epithelioid-cell differentiation surrounded by immature macrophages and cuffs of lymphocytes. Necrosis areas were not observed. Following *M. leprae* clearance (month 13), no histological lesion was noted (not shown). Interestingly, BCG administration in the *M. leprae*-infected mice (Group III) altered both the extension of the inflamed tissue and the number of epithelioid granulomas in the foot pad. Quantitative measurements (%) of the inflamed tissue area showed that mice from Group III (day 21 after BCG inoculation) presented a significantly ($p < 0.05$) larger area of inflammation (mean \pm S.D. = $16.5 \pm 0.5\%$) than did Group I mice (mean \pm S.D. = $8.4 \pm 2.7\%$). Furthermore, around 70% of the granulomas in Group III animals were classified as epithelioid granulomas (Fig. 2B) as compared to 30% in Group I mice ($p < 0.05$). Accordingly, analysis of the expression of TNF- α protein detected at the site of granuloma development (Fig. 2C) showed a higher number ($p < 0.05$) of TNF- α positively stained cells in epithelioid granulomas (mean \pm S.D. = $50 \pm 5\%$) than in non-epithelioid granulomas

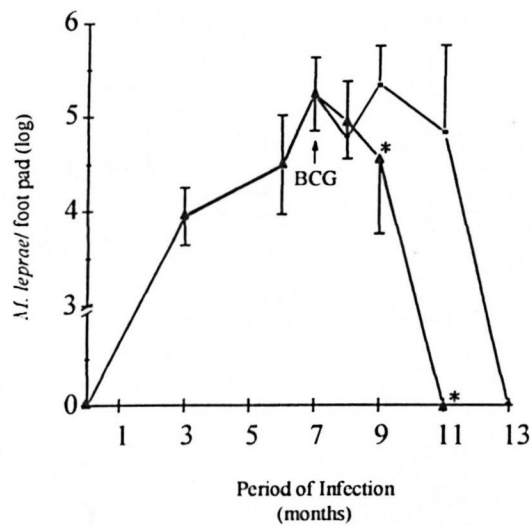


FIG. 1. Effect of BCG administration on *M. leprae* growth curve in BALB/c mice. Animals were infected in the foot pad with *M. leprae* alone (■ = Group I mice) or co-infected with BCG intravenously (▲ = Group III) at the peak of *M. leprae* growth (month 7, arrow). Bacterial growth was then periodically evaluated during *M. leprae* infection. At the times indicated, foot pads were recovered and bacterial counts performed as described in the Materials and Methods section. Results are reported as mean AFB counts (number of *M. leprae* per foot pad) \pm S.D. of two individual experiments (four animals were assayed per data point in each experiment). * Differences in *M. leprae* counts were found to be significant when compared to Group I mice.

(mean \pm S.D. = $20 \pm 2.5\%$). Therefore, a higher number of TNF- α -positive cells was detected in the foot pads of Group III mice.

Staining for fragmented DNA (3'-OH DNA ends), a hallmark of apoptosis, revealed the presence of apoptotic cells in the mouse foot pad granuloma (Fig. 2D). Most of the apoptotic nuclei were pyknotic and roughly rounded or oval in shape. Epithelioid granulomas presented a greater number ($p < 0.05$) of apoptotic cells (mean \pm S.D. = $30 \pm 3\%$) than non-epithelioid granulomas (mean \pm S.D. = $15 \pm 1\%$). Moreover, two different patterns of distribution for apoptotic cells were noted. Cuffs of apoptotic cells were often detected in epithelioid granulomas in contrast to the diffuse distribution found in the non-epithelioid granulomas.

Evolution of BCG infection in mice previously infected with *M. leprae*. The

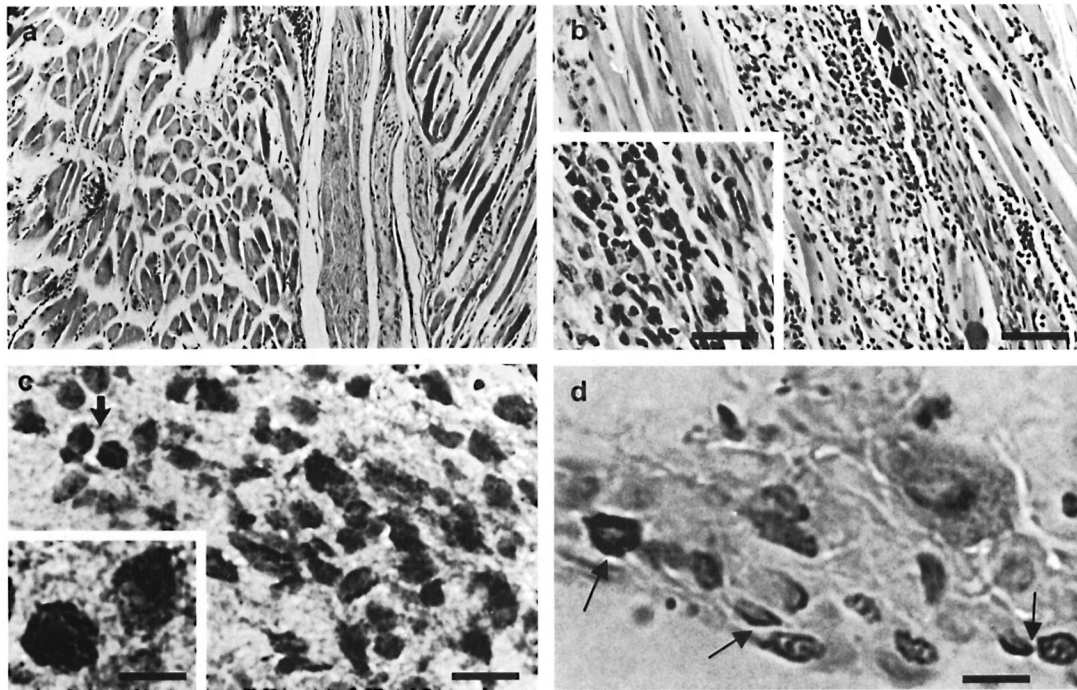


FIG. 2. Histological comparison of foot pads from *M. leprae*-infected mice (Group I) and *M. leprae* + BCG-infected mice (Group III). **A** = presence of diffuse granulomatous inflammation in mouse foot pads (Group I) (H&E; bar = 20 μ m). **B** = Epithelioid granuloma in mouse foot pads from Group III mice (H&E; bar = 20 μ m). **Inset** = Presence of epithelioid cells (H&E; bar = 10 μ m). **C** = Positive staining of TNF- α into an epithelioid granuloma in the mouse foot pad (Group III) (Bar = 10 μ m). **Inset** = Higher magnification of TNF- α -positive cells (Bar = 5 μ m). **D** = Detection of inflammatory apoptotic cells in an epithelioid granuloma in the foot pads of Group III mice (arrows = apoptotic cells; bar = 5 μ m).

number of BCG-CFU in the spleen was investigated during the course of BCG infection. In the present study, the evolution of BCG infection (Group II animals) was similar to that previously described by other investigators (¹⁷). Thus, the progressive increase in BCG-CFU clearly perceived as soon as 7 days after BCG inoculation persisted up to 21 days and began decreasing after 28 days (Fig. 3). In contrast, the number of live BCG recovered from the spleen of *M. leprae*/BCG-infected mice (Group III) was always very low. At 7, 14, 21, and 28 days after BCG inoculation, the number of BCG-CFU in the spleens of Group III mice was significantly lower ($p < 0.05$) than that observed in Group II mice (Fig. 3). As regards the liver, animals from Group III presented a higher number of BCG-induced granulomas ($p < 0.05$) than the Group II animals (data not shown).

Detection of TNF- α in sera and cytokine mRNA expression in spleen.

TNF- α levels in the serum of infected animals were investigated by ELISA. The kinetic of TNF- α secretion in BCG-negative and BCG-positive *M. leprae*-infected mice is shown in Figure 4. Mice infected only with BCG (Group II, 4 animals per data point) did not present detectable TNF- α in the serum until day 21 of infection (mean \pm S.E.M. = 7.0 ± 5.0 pg/ml). Moreover, low TNF- α levels (mean \pm S.E.M. = 20.8 ± 8.2 pg/ml) were also detected in most mice infected only with *M. leprae* (Group I, N = 9); whereas higher TNF- α levels were seen in animals infected with *M. leprae* + BCG (Group III, N = 4 per data point). A significant difference ($p < 0.05$) in the mean TNF- α values was noted 14 and 21 days after BCG infection when Groups II and III were compared. Mean TNF- α \pm S.E.M. were 0 versus 29 ± 4 pg/ml and 7 ± 5 versus 121 ± 42 pg/ml, respectively.

Analysis of TNF- α mRNA expression in the spleen from infected mice was also per-

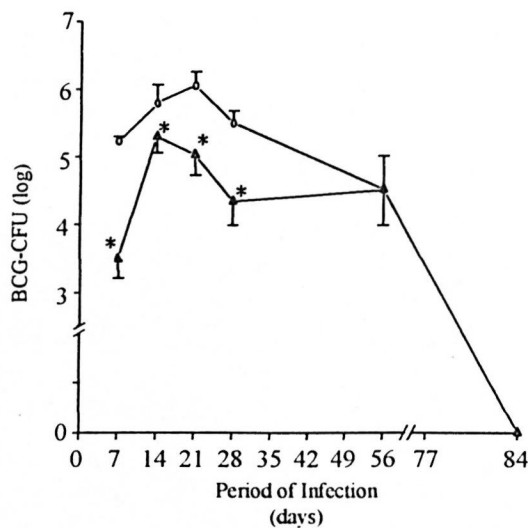


FIG. 3. Evaluation of BCG-CFU during the evolution of BCG infection in BCG- and *M. leprae*/BCG-infected mice. Co-infected animals (▲) are those infected with BCG intravenously 7 months after *M. leprae* injection in the foot pad (at the peak of *M. leprae* growth). Another group of animals was infected with BCG alone (○), and bacterial growth evaluated weekly during BCG infection. At the indicated times, spleens were recovered and bacterial counts (BCG-CFU) performed, as previously described. Data represent mean \pm S.D. of BCG-CFU enumeration (four animals per data point) from two separate experiments. * = Significant differences ($p < 0.05$) were noted in BCG-CFU at 7, 14, 21, and 28 days after BCG inoculation in co-infected mice (Group III) when compared to BCG-infected mice (Group II).

formed. The kinetic of cytokine gene expression was evaluated in the same groups (Groups II and III) at 7, 14, 21 and 28 days following BCG injection. In addition, a control group (healthy animals) and a *M. leprae*-infected group (animals analyzed at the peak of *M. leprae* growth) were also assayed. As demonstrated in Figure 5, a differentiated pattern of TNF- α mRNA expression was observed after BCG infection when mice from Groups II and III were compared. The graphic representation of the relative amounts of the TNF- α gene indicates that, in the BCG group (Group II), TNF- α gene expression in the spleen was enhanced during the infection period and showed maximal expression at day 21, thereby confirming previous observations⁽¹⁾. Surprisingly, in the BCG + *M. leprae*

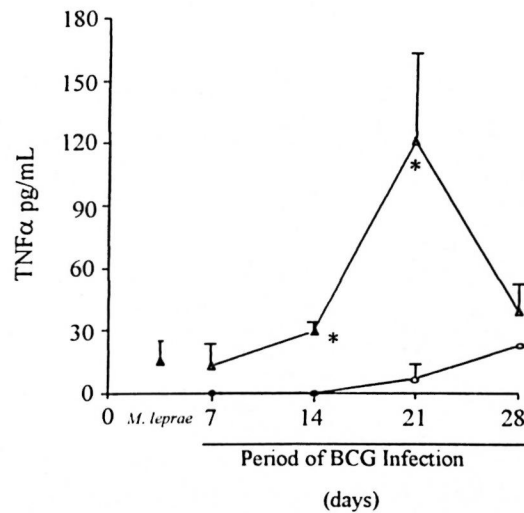


FIG. 4. Levels of TNF- α in the serum of mice infected with mycobacteria. BALB/c were infected with either BCG alone (○, Group II mice), *M. leprae* (▲, Group I), or BCG + *M. leprae* (△, Group III) as described. Serum samples were collected from these animals at month 7 of *M. leprae* infection and 7, 14, 21, and 28 days after BCG inoculation. Kinetic TNF data are mean TNF (pg/ml) \pm S.E.M. from quadruplicates per data point of one representative experiment. * = Significant difference ($p < 0.05$) in mean TNF values of Group III mice when compared to Group II animals. Group I, N = 9.

group (Group III) an early expression of the TNF- α gene was observed at day 7 of BCG infection, which became enhanced again at day 21, and at day 28 returned to background levels.

DISCUSSION

Several studies have demonstrated that BCG vaccination and immunotherapy with heat-killed *M. leprae* plus BCG, used as an adjunctive to multidrug therapy in leprosy, is responsible for both a histopathological shift of polarity toward the tuberculoid end of the spectrum and a clinical upgrading (reversal) reaction in multibacillary leprosy patients^(4, 5, 19, 20). It has been demonstrated that upon BCG vaccination, household contacts of leprosy patients presented positive, proliferative, T-cell responses as well as high interleukin-2 (IL-2) and IFN- γ production to several mycobacterial antigens *in vitro*⁽⁵⁾. A characteristic Th1-type of response found to be associated with a self-healing case of tuberculoid leprosy precipitated by BCG vac-

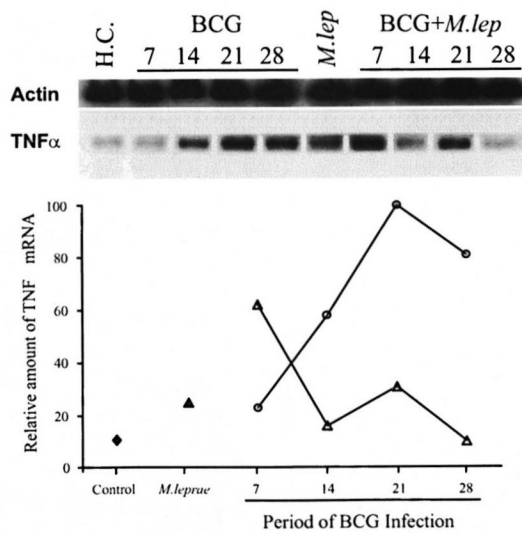


FIG. 5. Semi-quantitative RT-PCR analysis of TNF- α mRNA expression in spleens of mycobacteria-infected mice. TNF- α gene expression was investigated in different groups of mice at the peak of *M. leprae* infection (month 7 for Group I) and at 7, 14, 21, and 28 days after BCG infection (Groups II and III). Lane 1 = Healthy control mice (\blacklozenge); lane 6 = Group I mice (\blacktriangle); lanes 2–5 = Group II mice (\circ); lanes 7–10 = Group III mice (\triangle). Total RNA obtained from the spleen was reverse transcribed and the cDNA normalized to yield equivalent β -actin products. The relative amount of TNF- α mRNA was compared among samples from each group of animals and assessed as a percentage of the most intense band. Data from one representative experiment for each group is shown. Groups I, II, and III were defined as mentioned above. H.C. = Healthy control, *M. leprae* = *M. leprae*-infected mice (Group I).

cination (Sarno, E. N.; personal communication) has also been observed.

In this study, it has been demonstrated that systemic BCG administration at the peak of *M. leprae* infection modified the evolution of the immune inflammatory response initially induced by *M. leprae* in the mouse foot pad, and that co-infected mice presented a more effective clearance of *M. leprae*. Additionally, histopathological analyses have shown that BCG administration influenced the granulomatous response to *M. leprae* by converting mononuclear cell granulomas into epithelioid-cell granulomas with an enhanced number of apoptotic cells and extension of the inflamed tissue area in the foot pads.

In vivo studies of mycobacterial-induced granulomas have suggested that TNF- α plays a crucial role in granuloma formation (6, 10, 11). *M. leprae*, BCG, and several of their bacterial components have been shown to induce TNF- α secretion by monocytes *in vitro* (3, 24). In the present study, a series of observations indicated that the increased TNF- α production in this co-infection model correlated with a high degree of granulomatous inflammation (expressed by an extensive inflamed tissue area and number of epithelioid granulomas) in the foot pads of the infected mice: a) the increased number of TNF- α -positive stained cells observed in epithelioid as compared to non-epithelioid granulomas; b) enhanced serum TNF- α levels were detected only in Group III mice; and c) the difference in the kinetic expression of TNF- α mRNA in the spleen of co-infected animals.

Likewise, it has been demonstrated that BCG administration in BCG-susceptible mice (C57BL/6, BALB/c and B10.A) generated an efficient protective response to the challenge with homologous BCG and heterologous pathogens, such as *Lysteria monocytogenes* (17). BCG-induced restriction of *M. avium* proliferation and enhanced expression of TNF- α and IFN- γ mRNA in spleen cells have been described as well (2), suggesting that these cytokines can act in an additive or synergistic fashion in the induction of bacteriostasis. We have herein shown similar results. Interestingly, the accelerated process of bacterial elimination in co-infected mice (Group III) was also expressed by the reduction of BCG-CFU in the spleen (Fig. 3). Furthermore, our preliminary data indicated that BCG administration to these animals induced the early expression of IFN- γ mRNA (data not shown) in addition to TNF- α .

In Group III mice, an early expression of the TNF- α gene was observed at day 7 of BCG infection. It is possible that this early expression of TNF- α mRNA in the spleens is associated with the higher detection of TNF- α protein in the serum at day 21 when compared with Group I mice. In addition, BCG-infected mice (Group II) presented progressive increases of TNF- α mRNA and TNF- α serum levels. Interestingly, Groups II and III mice showed an early expression of TNF- α mRNA when compared with

TNF- α serum levels. The present data also suggest that there is a correlation between TNF- α serum levels or TNF- α mRNA in the spleen with the augmented TNF- α production and development of epithelioid granulomas in the foot pads of Group III mice. However, further studies are needed to more fully investigate these findings.

A higher number of apoptotic cells was noted in the epithelioid granulomas from the co-infected animals. It is possible that the enhanced rate of apoptosis in the epithelioid granulomas is associated with the higher expression of TNF- α *in situ* and to a more effective bacterial elimination in the foot pads of the Group III mice. Cree, *et al.* (8,9) have suggested that apoptosis may exceed mitosis as a cause for the rapid turnover of granulomas, which is probably associated with the fact that mycobacteria are generally not observed in epithelioid granulomas. However, the ability of BCG to induce or prevent apoptosis in mononuclear phagocytic cells remains controversial. It was recently demonstrated that BCG infection of resting human monocytes prevented apoptosis, and that this effect was accompanied by the induction of the A1 anti-apoptotic gene expression (13). On the other hand, it has also been observed that *in vitro* and *in vivo* *M. tuberculosis*-induced apoptosis of mononuclear phagocytes could be mediated by downregulation of the bcl-2 gene (12).

These data suggest that BCG administration in *M. leprae*-infected mice modulates the synthesis of TNF- α *in situ* and leads to the induction of protective granuloma formation and subsequent mycobacteria clearance. It may also be possible that apoptosis plays a role in the relationship between the immunopathological response and the development of protective granulomas containing bactericidal macrophages differentiated into epithelioid cells.

SUMMARY

In the present study, the experimental model of *Mycobacterium leprae* infection in the foot pads of BALB/c mice was used to investigate the effects of BCG administration on tumor necrosis factor-alpha (TNF- α) production and granuloma development. It was observed that mice intravenously infected with BCG 7 months after

M. leprae inoculation into the foot pads presented a more effective mycobacteria clearance, revealed by a significant reduction of BCG-colony forming units in the spleen and by the reduction of acid-fast bacilli (AFB) in the foot pads. BCG infection at the peak of *M. leprae* infection also modulated the granulomatous response to *M. leprae* by converting mononuclear granulomas into an epithelioid-cell granuloma. Furthermore, lower TNF- α serum levels were detected in *M. leprae*-infected mice when compared to mice infected with *M. leprae* + BCG. An analysis of the TNF- α gene expression in the spleen by semiquantitative reverse transcription-polymerase chain reactions (RT-PCR) demonstrated that co-infection with BCG induced an earlier expression of TNF- α mRNA than in *M. leprae*-infected mice. The numbers of TNF- α -positive cells and apoptotic cells were also enhanced in epithelioid versus non-epithelioid granulomas. As a whole, the data suggest that co-infection of *M. leprae*-infected mice with BCG modulates TNF- α synthesis which, in turn, leads to induction of protective epithelioid granuloma formation in the foot pads and subsequent mycobacterial clearance. Macrophage differentiation into epithelioid cells, in association with the enhancement of TNF- α production at the granuloma site, may represent a triggering signal that induced apoptosis in these cells, leading to mycobacterial elimination. Moreover, the rate of apoptosis in epithelioid granulomas may well be related to the extent of immunopathologically mediated tissue damage.

RESUMEN

Se utilizó el modelo de la infección de la almohadilla plantar del ratón BALB/c con *Mycobacterium leprae* para investigar el efecto del BCG sobre la producción de factor de necrosis tumoral alfa (TNF- α) y el desarrollo del granuloma. Se observó que los ratones infectados intravenosamente con BCG, siete meses antes de la inoculación de *M. leprae* en la almohadilla plantar, presentaron una eficiente depuración de *M. leprae*, una reducción significativa en el número de unidades formadoras de colonias de BCG en el bazo, y una reducción en el número de bacilos ácido-resistentes (BAAR) en la almohadilla plantar. La infección con BCG en el pico de infección con *M. leprae* también moduló la respuesta granulomatosa a *M. leprae* al convertir los granulomas mononucleares en granulomas epitelioides. Además, los niveles de TNF- α

en suero fueron más bajos en los animales infectados con *M. leprae* que en los animales inoculados con BCG y *M. leprae*. El análisis por RT-PCR de la expresión del gene TNF- α en el bazo mostró que la expresión de mRNA-TNF- α fue más temprana en los ratones coinfectados con BCG que en los animales infectados sólo con *M. leprae*. En los granulomas epitelioides se encontraron células TNF- α -positivas y células apoptóticas, en números mayores que los encontrados en los granulomas no epitelioides. Los datos sugieren que la coinfección de los ratones con *M. leprae* y BCG modula la síntesis de TNF- α , lo cual, a su vez, promueve la formación de granulomas epitelioides protectores en las almohadillas plantares, y la depuración eficiente de las micobacterias. La diferenciación de los macrófagos en células epitelioides, asociada con el aumento en la producción de TNF- α en el sitio del granuloma, podría representar un señal inductora de apoptosis en estas células que finalmente también conduciría a la eliminación de las micobacterias. Además, la frecuencia de apoptosis en los granulomas epitelioides podría estar relacionada con el grado de daño tisular mediado por mecanismos inmunopatológicos.

RÉSUMÉ

Dans cette étude, le modèle expérimental de l'infection de la patte de souris de souches BALB/c par *Mycobacterium leprae* fut utilisé pour évaluer les effets de l'administration de BCG sur la production de tumeur nécrosis facteur alpha (TNF- α) et le développement de granulômes. Il a été observé que l'infection par voie intra-veineuse de souris par le BCG, 7 mois après l'inoculation de *M. leprae* dans leurs pattes, favorisait l'élimination des mycobactéries, comme en témoigne une réduction significative du nombre de d'unités de BCG isolées à partir de la rate et capables de former des colonies, ainsi que du nombre de bacilles acido-alcool-résistants (BAAR) dans les pattes. L'infection par le BCG au moment du pic d'infection par *M. leprae* a entraîné la modification de la réponse granulomateuse contre *M. leprae*, de granulômes à cellules mononucléées à des granulômes à cellules de type épithélioïde. De plus des niveaux sériques plus bas en TNF- α furent détectés chez les souris infectées avec *M. leprae* seule, comparés aux souris infectées par *M. leprae* et BCG. Une analyse de l'expression du gène TNF- α dans la rate par transcription inverse-réaction de polymérase en chaîne (RT-PCR) a montré une expression plus précoce de l'ARNm de TNF- α chez les souris co-infectées avec le BCG que chez les souris infectées par *M. leprae* seule. Le nombre de cellules positives pour TNF- α et de cellules en apoptose était aussi augmenté dans les granulômes de type épithélioïdes en comparaison des granulômes non-épithélioïdes. En conclusion, les données suggèrent que la co-infection par le BCG de souris infectées par *M. leprae* module la synthèse de TNF- α , qui, en retour, conduit à l'induction de granulômes épithélioïdes protecteurs et à l'élimination des mycobactéries. La différenciation des macrophages en cellules épithélioïdes, en association

avec l'augmentation de la production de TNF- α au site des granulômes, pourrait représenter un signal clé pour l'induction de l'apoptose de ces derniers, conduisant à l'élimination des mycobactéries. De plus, le taux d'apoptose dans les granulômes épithélioïdes pourrait bien être relié au degré de dommage tissulaire à médiation immunopathologique.

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