Phenotypic changes in T cell populations during the reactivation of tuberculosis in mice

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(Accepted for publication 11 October 1997)

SUMMARY

The phenotypic changes of T lymphocytes during the reactivation of latent *Mycobacterium tuberculosis* infection by activation of the hypothalamic–pituitary–adrenal (HPA) axis was monitored using flow cytometric analysis. Subsets of $CD4^+$ and $CD8^+$ lymphocyte populations from the lung, spleen and draining lymph nodes of infected mice were identified based on their differential expression of the cell surface antigens CD44 and CD45RB. Latent infection was characterized by an accumulation of both naive, activated and memory CD4 and CD8 T lymphocytes in the lung and mediastinal lymph nodes. No changes were observed in the spleen of mice with latent infection when compared with uninfected mice. Immediately following the activation of the HPA axis, a reduction in all $CD4^+$ and $CD8^+$ T cells in the lung and mediastinal lymph nodes was observed. This correlated with the reactivation of mycobacterial growth. The decrease was transient for memory and naive CD4 and CD8 T lymphocyte populations in the lung. However, the number of naive CD4 and CD8 T lymphocyte populations in the mediastinal lymph node following reactivation was less than that found in mice with latent infection. These data provide the first characterization of T lymphocyte populations which may be functionally involved in the immunological response to HPA axis-induced reactivation of *M. tuberculosis* infection.

Keywords tuberculosis T cells stress hypothalamic–pituitary–adrenal axis

INTRODUCTION

Infection of individuals with *Mycobacterium tuberculosis* results in the establishment of immunity to the bacterium and the control of Mycobacterial growth [1,2]. The immunologically mediated inhibition of bacterial growth is thought to be mediated by antigenspecific T cells that result in macrophage activation [3–6]. The development of cell-mediated immunity results in the establishment of a latent infection; that is, viable bacteria are present but they do not increase in number. The latent state can last for the life of an infec\ted individual, who stands a 10% life time chance that the growth of the bacteria will reactivate. Reactivation leads to the growth of large numbers of bacteria, which disseminate into the airways and spread to the general population via droplet aerosol. The reactivation of the growth of the Tubercule bacillus is probably the result of a temporary suppression of the immune response, thus permitting the bacteria to grow [2,7–10].

Studies in our laboratory have shown that the growth of *M. tuberculosis* can be reactivated by activation of the hypothalamic– pituitary–adrenal (HPA) axis [11–13]. A steady state infection was

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established following systemic infection with low numbers of Mycobacteria. Reactivation resulted in the re-initiation of growth of the bacterium, especially in the lungs of mice. We show now that intranasal infection of mice, as described by Saunders & Cheers [14], also results in a steady state infection that was initially localized to the lung. Activation of the HPA axis resulted in the reactivation of the growth of the Tubercule bacillus. We monitored the changes in T cells following the establishment of the steady state infection and during reactivation by multi-parameter flow cytometry. Reactivation of mycobacterial growth by HPA axis activation resulted in changes in both CD4 and CD8 memory populations in the lung as well as the mediastinal lymph nodes. The results suggest that CD8 T cells may play an important role in the maintenance of latency.

MATERIALS AND METHODS

Mice

Specific pathogen-free male BALB/c mice were obtained from Charles River Labs (Wilmington, MA). Mice were 6–8 weeks of age at the start of each experiment. The animals were housed under sterile conditions in micro isolator cages (Lab Products,

Maywood, NJ) in a BSL-3 facility and given food and water *ad libitum.*

Bacterial infection

Mycobacterium tuberculosis (Erdman) was obtained from the American Type Culture Collection (ATCC 35801, TMC 107; Rockville, MD) and maintained as previously described by us [11]. Mice were infected intranasally as described by Saunders & Cheers [14]. In preliminary experiments we determined that instillation of $10³$ colony-forming units (CFU) resulted in the appearance of lung colonies 2 weeks after inoculation. Lower doses did not yield productive infections in all inoculated mice, while higher doses resulted in the progressive growth of the bacillus. The mice were inoculated, following anaesthesia with a ketamine/xylacine mixture, with 25μ l of the bacterial suspension containing 1000 CFU placed on the external nares with a micropipette. The number of microorganisms was confirmed by plate count. Bacterial growth was monitored by determining the number of CFU in the lung and spleen at various times after inoculation. A steady state infection was achieved by week 12 after the initial infection. This is similar to that which we had previously observed following systemic infection using 100 CFU [13].

HPA axis activation

The HPA axis was activated as previously described by us using a restraint paradigm [11–13]. Briefly, mice were placed into well ventilated 50-ml conical centrifuge tubes overnight. The mice were restrained for five daily 15-h cycles, rested for 2 days and then restrained again for five 15-h cycles. The mice were placed in conventional housing at the end of each period of HPA activation. Control mice, with latent infection, were housed without access to food and water.

Lymphocyte preparation

Mice were killed at selected time points following reactivation and the spleen and mediastinal lymph nodes were removed and placed separately into 5 ml of ice cold Dulbecco's modified Eagles' medium (DMEM; GIBCO BRL, Grand Island, NY) supplemented with 20% heat-inactivated fetal bovine serum (FBS; Hyclone, Logan, UT) containing glutamine and penicillin/streptomycin. Single-cell suspensions were prepared as previously described [11], resuspended in PBS and stored on ice until analysed.

The lungs were placed into Hanks' balanced salt solution (HBSS) containing 1% bovine serum albumin (BSA; Sigma, St Louis, MO), aspirated to remove blood and then cut into fine pieces. The tissue was digested for 45 min with a pre-screened lot of collagenase (150 U/ml; Worthington Biochemical, Cleveland, OH). The digested tissue was passed through 40 G stainless steel sieves and then through 18 G and 21 G gauge needles. The lymphocytes were purified from the lung cell suspension by centrifugation through a Ficoll–Paque (Pharmacia, Uppsala, Sweden) gradient. Following centrifugation the purified lymphocyte population was washed twice with PBS.

Flow cytometric analysis

Flow cytometric analysis was used to determine the expression of CD3, CD4, CD8, CD44 and CD45 by the T cell populations. The following conjugated antibodies were obtained from Pharmingen (San Diego, CA): anti-CD3, cytochrome-conjugated; anti-CD8, PE-conjugated; anti-CD45RB, FITC-conjugated and biotinylated anti-CD44. Anti-CD4 conjugated to RED 613 was obtained from

GIBCO BRL and streptavidin conjugated to AMCA was from Vector Technology (Burlingame, CA). Staining was performed by incubating $1-2 \times 10^6$ cells with antibody (1 μ g/ml) for 25 min at 4°C. After washing twice the strepavidin-conjugated AMCA was added and the samples incubated for an additional 20 min before washing with ice-cold PBS. Five-colour multi-parameter analysis was performed on a Coulter EPICS Elite flow cytometer (Coulter Corp., Hialeah, FL). Optical laser alignment calibration of the flow cytometer was performed using Coulter's DNA-Check EPICS alignment fluoro-sphere beads (Coulter) with coefficient of variations routinely < 2%. Standard Brite (Coulter) fluoro-sphere reference channel beads were used to standardize photomultiplier tube fluorescent intensities. Approximately 100 000 lymphocytes at a rate of 500 events/s were gated from monocytes and debris using linear forward light scatter *versus* 90° light scatter characteristics. CD3–Cychrome, CD4–PE, CD8–Red 613, and CD45RB–FITC were excited with a 488-nm 15-mW air-cooled Argon laser, and fluorescent light emission was collected through 675-nm, 575-nm, 610-nm and 525-nm bandpass filters, respectively. CD44–AMCA was excited with multiline UV (351–364 nm) 50-mW watercooled Argon laser and collected with a 460-nm bandpass filter at $40-\mu s$ gated amplifier delay. All fluorescent signals were measured in logarithmic mode. Data files were stored in list mode format and extended analyses was performed using Winlist Software (Verity Software House, Topsham, ME). Each single colour-stained control was used to establish an off line compensation curve using Winlist Software. The T cells were identified by gating of the $CD3^+$ cells. Those cells were then analysed for CD4 and CD8 expression and those populations were further analysed for the expression of CD45RB and CD44. The MoAbs to CD44 and CD45RB revealed bimodal staining patterns which resulted in the identification of cells with bright and dim intensities indicating the presence of two populations**.** The data represent the means of four different experiments using three mice per time point. Data are expressed as cell numbers based on the total yield of cells and the percentage of each population.

Preparation of tissue section for histologic examination and staining

The lungs were perfused via the trachea with HBSS and immersed for at least 24 h with 10% buffered formalin. Specimens were processed, stained with haematoxylin and eosin and sectioned routinely for light microscopic quantitative evaluation.

Statistical analysis

The differences in T lymphocyte populations at each time point was determined by one-way analysis of variance (ANOVA) (Stat-Most, Salt Lake City, UT). The differences between the T lymphocyte populations from uninfected mice and mice with latent infection were evaluated, as was the effect of time on changes in T cells from latently infected mice. The changes in T cell subpopulations that resulted from activation of the HPA axis were also analysed by ANOVA. The results were significant if they equalled or exceeded the 95% confidence level (*P* < 0·05).

RESULTS

Reactivation of M. tuberculosis *growth by activation of the HPA axis*

Activation of the HPA axis resulted in the resumption of growth of *M. tuberculosis*. The results in Fig. 1 show that the number of

q 1998 Blackwell Science Ltd, *Clinical and Experimental Immunology*, **111**:309–315

Fig. 1. *Mycobacterium tuberculosis* growth and reactivation. Mice were infected with $10³$ colony-forming units (CFU) of *M. tuberculosis* (Erdman) intranasally. Growth in the lung was monitored at the times indicated. Once a latent infection was established, the hypothalamic–pituitary–adrenal (HPA) axis was activated by restraining the mice for 10 15-h sessions. The growth of the Mycobacteria was monitored by plate count. The time represents weeks after initial infection. Values represent means \pm s.e.m. of $n = 4$ individual groups of three mice. The effect of HPA activation and of time was significant as determined by ANOVA. \blacktriangle , Control; \blacklozenge , HPA activation.

microorganisms isolated from the lung following intranasal instillation increased to 46 400 CFU at week 6 after infection and then decreased until it reached a level that varied from 14 000 to 28 000 CFU/lung by weeks 12–14. This level remained relatively constant for the duration of the experiment. Following activation of the HPA axis beginning at week 12, the numbers of bacteria increased to 81 600 CFU/lung by week 20 before declining. Bacteria were also detected in the spleen, and the pattern of growth, while similar, remained 10-fold less than that isolated from the lungs (data not shown).

Analysis of T cell populations in mice with latent tuberculosis

The changes in CD45 and CD44 were used as markers for naive, activated and memory cells [15–17]. Thus, cells expressing higher levels of CD45RB and low levels of CD44 were considered to be naive, those expressing low levels of both CD45RB and CD44

were considered to be activated, and those expressing high levels of CD44 but low levels of CD45RB were classified as memory cells. The total cellularity of all organs was significantly decreased following activation of the HPA axis (week 2) and there was a significant increase in total cellularity at 6 weeks only in the lung. At all other time points there was no change in cellularity compared with the number of cells obtained from animals with latent disease. Therefore, changes in T cell populations were due to alterations in the percentage of distinct T cell subsets.

The results in Fig. 2a show that there was an increase in both naive (CD45RB^{hi}), activated (CD45RB^{lo}) and memory (CD44^{hi}) $CD4^+$ and $CD8^+$ cell populations in the lungs of mice with latent disease compared with the T cell populations in the lungs of uninfected mice. The number of CD44^{hi} memory cells isolated from the lungs of mice with latent disease was also greater then the number of memory cells from uninfected mice and greater then CD45RB^{hi} (naive) cells from infected mice. Similarly, there was a significant increase in both naive, activated and memory $CD4^+$ and $CD8^+$ populations in the draining mediastinal lymph nodes (Fig. 2b). In this case, and in contrast to that observed in the lung, high levels of activated (CD45RB^{lo}/CD44^{lo}) and low numbers of memory (CD44^{hi}) CD4⁺ or CD8⁺ cells were found in the mediastinal node of infected mice. No changes in T cell populations were associated with latent disease in the spleen (Fig. 2c).

Changes in naive, activated and memory T cell populations

associated with reactivation of latent M. tuberculosis *infection* HPA activation resulted in a reduction in the number of naive (CD45RB^{hi}), activated (CD45RB^{lo}) and memory (CD44^{hi}) $CD4^+$ and $CD8^+$ lymphocytes in the lungs, spleen and mediastinal lymph nodes. These changes coincided with a decrease in the cellularity of these organs as a result of HPA activation (data not shown). Figure 3 shows the changes in T cell populations in the lung following reactivation. It should be noted that an analysis of T cell populations in animals with latent disease was also done at each of the time points indicated. Since these did not change during the course of these studies as determined by ANOVA, they were treated as one group and the collapsed data are found at time 0, i.e. before activation of the HPA axis.

Fig. 2. Analysis of T lymphocyte populations in uninfected mice and in mice with latent *Mycobacterium tuberculosis* infection. Mice were infected intranasally with 10³ colony-forming units (CFU) of *M. tuberculosis*. Twelve weeks after the initial infection, when latent infection was established (see Fig. 1), the mice were killed and the T cell populations from these mice and from age-matched uninfected control mice were analysed by flow cytometry. The number of T cells was determined by multiplying the total number of viable cells by the percent of $CD3⁺$ T cells in the population isolated from the lung, spleen, and mediastinal lymph nodes. The number of $CD4^+$ and $CD8^+$ cells was calculated by multiplying the percentage of gated $CD4^+$ or $CD8^+$ cells harvested from each organ by the total number of $CD3⁺$ cells. Finally, dual expression of CD45RB and CD44 was calculated by multiplying the percentage of gated populations determined to be either high and low expressors of CD45RB and CD44 by the total number of either CD4⁺ or CD8⁺. (a) Lung. (b) Mediastinal lymph node. (c) Spleen. Values represent means \pm s.e.m. for four groups of $n = 3$ mice. The effect of infection with *M. tuberculosis* on T cell populations in lung and mediastinal lymph node was significantly different as determined by ANOVA. \Box , Uninfected; ,,,,,,,latent infection.

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Fig. 3. Analysis of T lymphocyte populations in the lung following reactivation of latent *Mycobacterium tuberculosis* infection. Mice with latent infection were restrained to activate the hypothalamic–pituitary–adrenal (HPA) axis. Immediately following HPA activation and at the times indicated the T lymphocyte populations in the lung were analysed using multi-parameter flow cytometry. Mice with latent disease served as controls. No differences in the cell populations from mice with latent disease were observed throughout the time course. The data are therefore collapsed and expressed as time 0, i.e. before HPA activation. The number of cells in each subpopulation was calculated as described in the legend for Fig. 2. (a) $CD45^{h}/CD4^{h}$. (b) $CD45^{h}/CD4^{h}$. (c) CD45^{lo}/CD44^{hi}. Values represent means \pm s.e.m. for four groups of $n=3$ mice. The effect of HPA activation and of time was significant on the naive, memory and activated populations as determined by ANOVA. \Box , CD4; \blacksquare , CD8.

Activated (Fig. 3b) (CD45RB^{lo}/CD44^{lo}) CD4 and CD8 populations recovered quickly, as did memory (Fig. 3c) (CD44^{hi}) CD4 and CD8 populations, the lungs reaching a peak in cell number by week 6 following reactivation. These cell numbers declined thereafter, which coincided with a decline in the numbers of Tubercle bacilli isolated from the lung. The recovery of the naive (Fig. 3a) $(CD45RB^{hi})$ populations of T cells in the lung mimicked that observed for memory cells, except the cell numbers that were recovered remained depressed compared with that of the memory population.

The CD4 and CD8 naive, activated and memory cell populations in the mediastinal lymph node were also depressed initially (Fig. 4). However, the CD4 and CD8 naive and activated populations (Fig. 4a,b) remained depressed throughout the 12 weeks of observation. In contrast, the levels of CD8 memory cells recovered to levels found in animals with latent disease and then declined (Fig. 4c). The increase in CD8 memory cells occurred prior to the increase of memory cells in the lung, and their decline was associated with an increase of this T cell population in the lung. The CD4 population of memory cells, however, remained depressed following HPA activation.

The naive, activated and memory CD4 cell populations in the spleen also recovered following HPA activation (Fig. 5). In contrast, while the number of naive and activated CD8 T cells in the spleen recovered, the number of memory CD8 T cells remained depressed. No changes were observed in any population of cells obtained from the superficial cervical lymph node (data not shown).

Reactivation of M. tuberculosis *results in an increase of inflammation in the lung*

The lungs of mice with latent disease had multifocal, mild perivascular and peribronchiolar lymphoid infiltrates and minimal granulomatous pneumonia (Fig. 6a). Immediately following activation of the HPA axis, no inflammation was observed in the lungs, which appeared microscopically normal (Fig. 6b). In contrast, 6 weeks after reactivation the lungs contained many foci of severe granulomatous pneumonia and prominent perivascular and peribronchiolar lymphoid infiltrates (Fig. 6c). There were numerous areas of macrophages containing a significant amount of acid-fast debris that was not found in the lungs during latency or immediately after activation of the HPA axis (not shown).

DISCUSSION

Most people who are infected with *M. tuberculosis* do not develop disease. Infected individuals are assumed to mount an effective immune response to the initial infection that limits proliferation of the bacteria. Thus, infection is likely to be asymptomatic. Infected people can only be identified by a positive skin test reaction. The host factors that are responsible for the development of clinical disease have not been clearly described, but are thought to involve a temporary suppression of the immune response and re-initiation of Mycobacterial growth [1,2]. A key factor in studying the changes that result in clinical disease is the establishment of a 'latent' infection followed by re-initiation of Mycobacterial growth resulting from a temporary suppression of immunity. Previously,

Fig. 4. Analysis of T lymphocyte populations in the mediastinal lymph node following reactivation of latent *Mycobacterium tuberculosis*. See legend for Fig. 2 for an explanation of methodology. (a) CD45^{hi}/CD44^{lo}. (b) CD45^{lo}/CD44^{lo}. (c) CD45^{lo}/CD44^{hi}. Values represent means \pm s.e.m. for four groups of *n* = 3 mice. The effect of hypothalamic–pituitary–adrenal (HPA) activation was significant on all populations tested. The effect of time on the naive and memory populations was also significant. \Box , CD4; \blacksquare , CD8.

Fig. 5. Analysis of T lymphocyte populations in the spleen following reactivation of latent *Mycobacterium tuberculosis*. See legend for Fig. 2 for an explanation of methodology. (a) CD45^{hi}/CD44^{lo}. (b) CD45^{lo}/CD44^{lo}. (c) CD45^{lo}/CD44^{hi}. Values represent means \pm s.e.m. for four groups of *n* = 3 mice. The effect of hypothalamic–pituitary–adrenal (HPA) activation was significant by ANOVA. The effect of time following HPA activation was not significant. \square , CD4; \blacksquare , CD8.

we have shown that systemic infection of mice with a low dose of *M. tuberculosis* (Erdman) resulted in a steady state infection in which the numbers of bacteria isolated from the lung and spleen did not increase for several months [13]. The results of this investigation, using intranasal instillation of low doses of microorganisms, also resulted in the establishment of a steady state infection in the lungs. Our results confirm our previous observations that activation of the HPA axis by restraint resulted in the reinitiation of growth of the Tubercule bacillus [13]. The reactivation was not the result of changes in the nutritional status of the mice, since the control mice, those with latent disease, were housed in cages without food or water during the same time that other mice were being stressed. The restraint stress paradigm, as previously described by us, results in a sustained increase in plasma corticosterone as well as a decreased cellularity of the lymphoid organs and the lung. While this model may not strictly represent clinical disease in man, it does provide an opportunity to monitor the changes in T cell populations that may be responsible for controlling mycobacterial growth during a steady state infection and during an increase in Mycobacterial growth following a temporary suppression of immunity that results from HPA activation [18–23]. To $\frac{1}{2}$ Cell number of Representation in the plane of the New York (1998) and $\frac{1}{2}$ Cell number (x 1040) an

Activation-associated alterations of T cell surface phenotypes have been reported to occur following stimulation with antigen [24,25]. Expression of CD44 increases coincident with the frequency of antigen-specific reactivity. After antigenic challenge, there is also an increase in the expression of VLA-4, LFA-1 and CD54 cell adhesion molecules and a decrease in CD62L and CD45RB. Thus, T cell activation has been shown to correlate with a shift from the CD44^{lo} to CD44^{hi} and from CD45RB^{hi} to the $CD45RB^{lo}$ phenotype.

Based on these phenotypic changes, we found that there was an accumulation of both naive, activated and memory CD4 and CD8 populations in the lung and mediastinal lymph node of mice during latent *M. tuberculosis* infection compared with uninfected control mice. The populations decreased as a result of HPA activation, which resulted in resumption of the growth of *M. tuberculosis*. This suggests that increased levels of corticosterone that occur during activation of the HPA axis [11–13] resulted in the breakdown of immunological control of tuberculosis (TB). Interestingly, the cell populations found in the spleen of mice with latent disease resembled the phenotype of those from uninfected mice. The initial immune response to the Mycobacterium, therefore, was probably limited to the lungs because the mice were infected intranasally. Alterations in the T cell population in the spleen were not observed until Mycobacterial growth was reinitiated following HPA activation.

The changes in T cell populations that followed reactivation also indicate that the immune response was largely localized to the lung and draining lymph nodes. The populations in the lung that changed following HPA activation indicated that both CD4 and CD8 activated and memory populations were stimulated. The response in the mediastinal lymph node was different from that observed in the lung; only the activated population increased in response to the increased numbers of microorganisms. Increases in memory cells were not observed.

Our results are similar to those observed by Griffin & Orme [24] and by Andersen *et al*. [26], who showed that there was a preferential accumulation of CD44^{hi}/CD45RB^{lo} memory subsets in the spleen of mice injected intravenously with *M. tuberculosis*. Coulson & Wilson also reported increases in CD44^{hi} memory CD4 cells in the lungs of mice following immunization with attenuated *Schistosoma mansoni* cercariae [27]. We too found that memory cells accumulated in the lungs of mice infected intranasally. Our observations, while similar, were made after allowing the mice to control the growth of low numbers of microorganisms and then temporarily suppressing the immune response by activation of the HPA axis. This allowed the bacteria to resume growth. Recovery of immunocompetence was associated with a decrease in bacterial load. This is analogous to human disease in the pre-antibiotic era. Latent TB is reactivated due to a temporary suppression of immunity; patients would experience episodes of reactivation followed by latent disease during their lifetime [28].

While it is generally agreed that CD4 cells are important cytokine producers that result in macrophage activation, the role of CD8 T cells in mediating protective immunity to TB is not clearly understood. One study by Saunders & Cheers [14] examined the changes in CD4 and CD8 T lymphocytes in response to intranasal infection with *M. avium*. They found that depletion of $CD8⁺$ T cells had no effect on the growth of *M. avium* in the lung and that CDS^+ cells from *M. avium*-infected mice did not produce interferon-gamma (IFN- γ) *in vitro*. In contrast, mice that were depleted of $CD4⁺$ T cells were more susceptible to the growth of *M. avium*. We showed that there was an increase in both CD8 and CD4 T cells in response to intranasal infection with *M. tuberculosis*. The decrease in both T cell populations as a result of HPA activation correlated with an increase in bacterial growth. These results suggest that $CD8⁺$ cells may be an important effector cell during an infection with *M. tuberculosis*. This possibility has been reinforced by recent studies that show that β_2 -microglobulin knock-out mice, that lack functional MHC class I molecules and consequently are deficient in CD8 T lymphocytes, are more susceptible to the growth of *M. tuberculosis* [29]. Also, both

Fig. 6. Histology of the lung following reactivation of *Mycobacterium tuberculosis.* Haematoxylin and eosin staining under light microscope. (a) Twelve weeks post-infection (latency). (b) Immediately following hypothalamic–pituitary–adrenal (HPA) activation. (c) Six weeks later (week 20).

 $CD4^+$ and $CD8^+$ T lymphocytes confer a high degree of protective immunity against TB in mice after vaccination with the Mycobacterial protein heat shock protein (hsp)65 [30].

The results of this study indicate that memory immune T cells play a significant role in controlling *M. tuberculosis* following reactivation of latent disease. These cells may also be more

resistant to the suppressive effects of corticosteroids [19–23]. Both populations of CD4 and CD8 memory cells may mediate cytolysis of infected macrophages releasing bacteria into the environment [31–38]. The release of cytokines, especially IFN- γ produced by both populations, may result in the activation of macrophages that can control the growth of the Mycobacterium. The stimulation of memory immune T cells following reactivation of Mycobacterial growth coincided with the decrease in Mycobacteria in the lungs. The role of the memory immune populations in re-establishing control of the growth of *M. tuberculosis* is currently under investigation.

ACKNOWLEDGMENTS

Supported by NIMH grants MH45679, MH54966 and by Cancer Centre support grant CA16058.

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