

## Cytokine Gene Expression in the Foot Pad and Spleen of BALB/cAJcl Mice Infected with *M. leprae*<sup>1</sup>

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The mechanisms of pathogenesis for *Mycobacterium leprae* infection remains unknown. Mice with cellular immunodeficiency are highly susceptible to *M. leprae* which is an intracellular pathogen (4, 6, 13, 16, 22, 33, 49, 50). In immunocompetent mice, the only susceptible sites to *M. leprae* infection are skin tissues, such as the foot pad and the ear. When *M. leprae* are inoculated into the foot pads of normal mice, the bacilli can multiply in the site but the infection is limited to  $10^7$  bacilli (32, 37-39). Even though *M. leprae* were inoculated into normal mice by intravenous injection, they were not able to multiply in the viscera such as the liver and the spleen (33). To elucidate the reasons for the restricted growth of *M. leprae* in the foot pads of immunocompetent mice, we examined and compared the expression of cytokine mRNAs in the foot pads and spleens of BALB/cAJcl mice infected with *M. leprae*.

### MATERIALS AND METHODS

**Mice.** BALB/cAJcl mice were bred in the Central Institute for Experimental Animals, Kanagawa, Japan. Twenty female mice aged 6 weeks were used in the experi-

ment. They were housed in the Animal Care Facility of our laboratory after inoculation with *M. leprae*.

***M. leprae*.** Leprosy bacilli, Thai-53 strain, derived from foot pad passage of nude mice were used. The suspensions of viable (37) or heat-killed *M. leprae* (20) were prepared as described by references indicated.

**Inoculation.** Mice were infected with viable or heat-killed *M. leprae* each in both hind foot pads ( $2 \times 10^7$ ) combined with intravenous inoculation at a dose of  $4 \times 10^7$ . Two mice each were sacrificed on days 1, 10, 30 and 150 of inoculation, and the results were compared with those of the mice prior to infection (24 hr before infection; day 0).

### Cytokine genes detection by RT-PCR.

The expression of cytokine genes in the foot pads and spleens was examined by the reverse transcriptase-polymerase chain reaction (RT-PCR) method. RT-PCR of cytokine mRNAs from mouse tissues uninoculated (day 0) or inoculated with *M. leprae* at varying periods of time as described above was performed as described (47, 48). In brief, total RNA was extracted by the acid guanidinium method from 2 foot pads and 2 spleens which were isolated from 2 infected mice, pooled and frozen, and cDNA was synthesized using Moloney murine leukemia virus (M-MLV) reverse transcriptase and oligo (dT). Reactions were incubated in a thermal cycler (Astec High Voltage, Ashland, Massachusetts, U.S.A.; PC-800) for 35 cycles. The cDNA concentrations were normalized to yield equivalent  $\beta$ -Actin PCR products. After PCR, 10  $\mu$ l of the DNA from each tube were loaded onto 1.5% agarose gels in TAE buffer. Products were visualized by ethidium bromide staining. The sense and antisense primers used were as follows: Interleukin-1 alpha (IL-1-

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$\alpha$ ), 5'-CTCTAGAGCACCATGCTACAG-AC-3' and 5'-TGGAATCCAGGGGAAA-CACTG-3'. IL-2, 5'-ATGTACAGCATG-CAGCTCGCATC-3' and 5'-GGCTTGTT-GAGATGATGCTTTGACA-3'. IL-4, 5'-ATGGGTCTCAACCCCAAGCTAGT-3' and 5'-GCTCTTTAGGCTTTCCAGGAAGTC-3'. IL-6, 5'-ATGAAGTTCCTCTCTGCA-AGAGACT-3' and 5'-CACTAGGTTTGC-CGAGTAGATCTC-3'. IL-10, 5'-TACC-TGGTAGAAGTGATGCC-3' and 5'-CA-TCATGTATGCTTCTATGC-3'. IL-12 (p40), 5'-CAGAAGCTAACCATCTGGTTTG-3' and 5'-TCCGGAGTAATTTGGTGCTTC-ACAC-3'. Gamma interferon (IFN- $\gamma$ ), 5'-TGAACGCTACACACTGCA TCTTGG-3' and 5'-CGACTCCTTTTCCGCTTCCT-GAG-3'. Tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), 5'-GGCAGGTCTACTTTGG-AGTCATTGC-3' and 5'-ACATTTCGAG-GCTCCAGTGAATTCGG-3'. TNF- $\beta$ , 5'-TGGCTGGGAACAGGGGAAGGTTGAC-3' and 5'-CGTGCTTT CTTCTAG AAC-CCCTTGG-3'. CD4, 5'-TGTGCCGAGC-CATCTCTCTTAGG-3' and 5'-GCACTG-AGAGTGTCATGCCGAAC-3'. CD8, 5'-ATGCAGCCATGGCTCTGGCTGG-3' and 5'-GCATGTGAGGCCCTTCTGGG-TC-3'.  $\beta$ -Actin, 5'-TGGAAATCCTGTGG-CATCCATGAAAC-3' and 5'-TAAAACG-CAGCTCAGTAACAGTCCG-3'.

**Immunohistologic staining of mouse foot pad tissues.** Lymphocytes bearing CD4 and CD8 surface expression in the foot pad prior to the inoculation and 30 days after viable *M. leprae* inoculation were determined by immunohistochemistry. The tissue with OCT medium (Miles, Inc., Elkhart, Indiana, U.S.A.) was cut for 6- $\mu$ m thickness, fixed with acetone and chloroform, and then blocked with normal rabbit serum (Dako Corp., Carpinteria, California, U.S.A.; X902) before undergoing incubations with the monoclonal antibodies (anti-CD4 and anti-CD8; Serotec, Kidlington, U.K.; KT174 and KT15) overnight at 4°C followed by biotinylated rabbit anti-rat immunoglobulins (Dako E0468) for 30 min. Slides were washed with phosphate buffered saline (PBS) between incubations. Primary antibodies were visualized by using the LSAB kit (Dako). Slides were counterstained with hematoxylin and mounted in Ukitt (O. Kinder, Germany).

**ELISA.** The mouse serum immunoglobulin of each group was measured by an ELISA method using goat anti-mouse IgG and IgM. The serum antibodies to *M. leprae* phenolic glycolipid-I (PGL-I), lipoarabinomannan-B (LAM-B) and heat-shock protein 65-kDa (hsp65 kDa) were measured by the ELISA method of Cho, *et al.* (5). LAM-B was prepared in our laboratory from a *M. leprae* cell-wall fraction obtained from armadillo tissues (11); PGL-I also was prepared from *M. leprae*-infected armadillo tissues using the method by Hunter, *et al.* (10, 12). *M. leprae*-hsp65 was also prepared in our laboratory from an *Escherichia coli* strain and affinity-purified with monoclonal antibody to hsp65 (26). Mouse serum was diluted at 1:100 with 2% bovine serum albumin-phosphate buffered saline-azide (BSA-PSA-azide) for antibody titration. The microplates (Nunc, Roskilde, Denmark) were coated with 50  $\mu$ l/well of PGL-I at 2  $\mu$ g/ml, LAM-B at 1  $\mu$ g/ml or hsp65 at 2  $\mu$ g/ml in 0.05 ml carbonate buffer (pH 9.6); the other half of each plate was coated with a coating buffer alone as a control. The peroxidase-conjugated goat anti-mouse IgG F(ab')<sub>2</sub> and IgM F(ab')<sub>2</sub> fragment (diluted 1:1000) was used for secondary antibody. Serum antibody level (OD<sub>490-650 nm</sub>) was calculated by subtracting the mean optical density (OD) of the control wells from the mean OD of the antigen-coated wells.

## RESULTS

**Cytokine mRNA expression in foot pads of *M. leprae*-inoculated mice.** The changes in the cytokine mRNAs expression over time in the foot pads of uninoculated (24 hr before infection; day 0) and intravenously inoculated mice combined with the foot pad injection with viable or heat-killed *M. leprae* are shown in Figure 1. Physiologic expression of IL-1 $\alpha$  and TNF- $\beta$  mRNAs were observed in the foot pads of the uninoculated mice. In addition, in mice inoculated with viable *M. leprae*, the usual appearance of IL-1 $\alpha$  and TNF- $\beta$  mRNA was intensified; TNF- $\alpha$  mRNA was induced on day 1 of inoculation. The expression of these mRNAs increased with the progression of the infection until day 10, and then decreased on day 30 and increased again on day 150. The same pattern of mRNA expression was also observed in

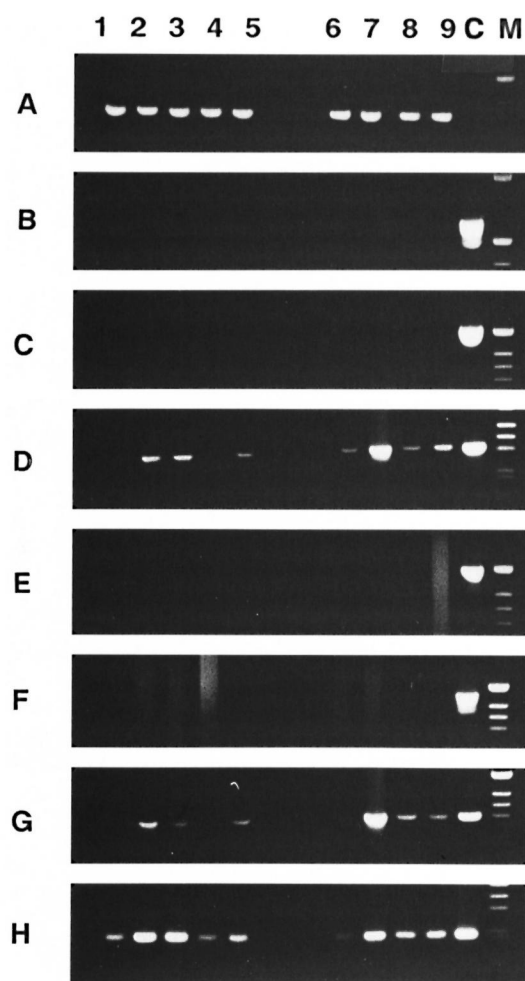


FIG. 1. Cytokine mRNA expression in foot pads of *M. leprae*-inoculated mice. PCR analysis of cDNA from two foot pad tissues pooled from 2 infected mice was carried out at varying times by: lane 1, 24 hr before infection (day 0); lane 2, on day 1; lane 3, on day 10; lane 4, on day 30; lane 5, on day 150 after infection with viable *M. leprae*. Lane 6, on day 1; lane 7, on day 10; lane 8, on day 30; lane 9, on day 150 after infection with heat-killed *M. leprae*. Lane C was the positive control from ConA-stimulated spleen cells. Specific primers used for the sequence of cytokines, CD4 and CD8 were: A =  $\beta$ -Actin, B = CD4, C = CD8, D = IL-1- $\alpha$ , E = IL-2, F = IFN- $\gamma$ , G = TNF- $\alpha$  and H = TNF- $\beta$ . Reactions were incubated in an Astec-800 for 35 cycles as conditions with denaturation 1 min, 94°C; annealing 2 min, 60°C or 65°C; extension 3 min, 72°C.

mice inoculated with heat-killed *M. leprae*; the degree of induction of these mRNAs was lower on day 1 and the expression was slightly stronger after day 10.

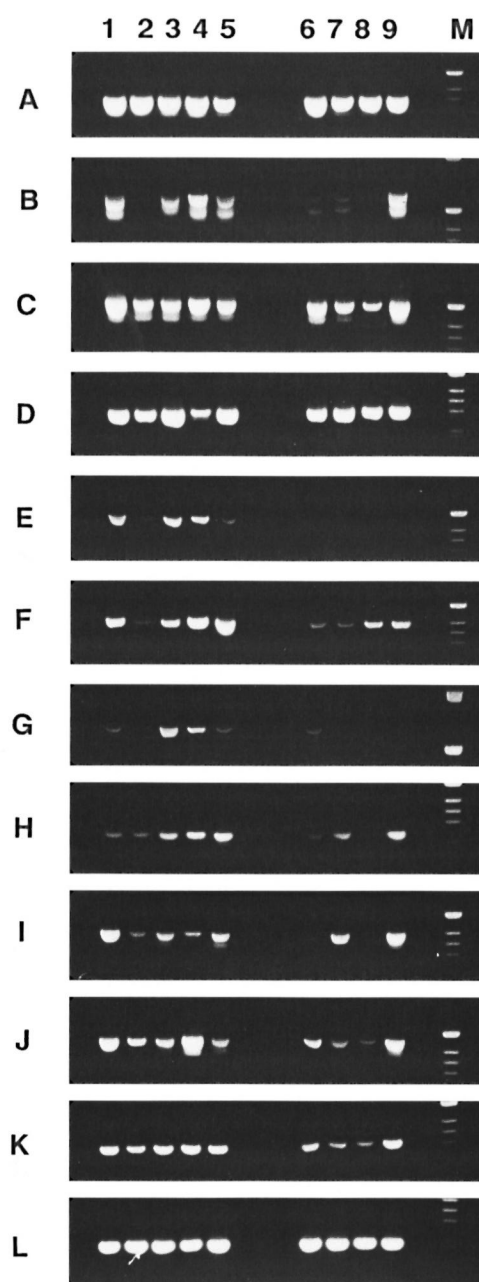


FIG. 2. Cytokine mRNA expression in the spleens of *M. leprae*-inoculated mice. PCR analysis of cDNA from 2 spleen tissues pooled from 2 infected mice were carried out at varying times as well as the foot pad after inoculation with viable or heat-killed *M. leprae*. Lane 1, 24 hr before inoculation (day 0); lane 2, on day 1; lane 3, on day 10; lane 4, on day 30; lane 5, on day 150 after inoculation with viable *M. leprae*. Lane 6, on day 1; lane 7, on day 10; lane 8, on day 30; lane 9, on day 150 after inoculation with heat-killed *M. leprae*. Specific primers used were as follows: A =  $\beta$ -Actin, B = CD4, C = CD8, D = IL-1- $\alpha$ , E = IL-2, F = IL-4, G = IL-6, H = IL-10, I = IL-12(p40), J = IFN- $\gamma$ , K = TNF- $\alpha$ , L = TNF- $\beta$ .



**Cytokine mRNA expression in spleens of *M. leprae*-inoculated mice.** The changes of cytokine mRNA expression in the spleen over time in uninoculated mice (day 0) and mice inoculated with viable or heat-killed *M. leprae* at varying periods of 1, 10, 30 and 150 days are shown in Figure 2. All of the cytokine mRNAs were examined; IL-1 $\alpha$ , -2, -4, -6, -10, -12, IFN- $\gamma$ , TNF- $\alpha$  and TNF- $\beta$  were expressed physiologically in the spleens of uninoculated mice 24 hr before infection. CD4 and CD8 mRNAs also were expressed physiologically. In mice inoculated with viable *M. leprae*, CD4 mRNAs were undetectable on day 1 of inoculation, but the expression was recovered over time. In mice inoculated with heat-killed *M. leprae*, CD4 mRNA expression in the spleen was decreased up to day 30 of inoculation, and IL-2 and IL-6 mRNAs (except for a weak expression on day 1) were not expressed. An interesting observation was that the loss of CD4 mRNA expression on day 1 of inoculation of live *M. leprae* was accompanied by a diminished expression of IL-2, -4, -6 and IL-12 mRNAs, and the fluctuation in cytokine mRNA expression followed the pattern of fluctuation of CD4 mRNA expression. An increase in IL-4 and IL-10 mRNAs expression was observed during the course of infection in mice inoculated with viable *M. leprae*. Fluctuation of IL-1- $\alpha$  mRNA expression was observed during the course of infection in mice inoculated with live bacteria but not in those inoculated with killed bacteria. TNF- $\beta$  mRNA expression remained constant at the pre-inoculation level in both groups inoculated with viable and heat-killed *M. leprae*.

**Immunohistological examinations of CD4+ and CD8+ lymphocytes in foot pads of *M. leprae*-infected mice.** On day 30 of inoculation the foot pads received viable *M. leprae* and the foot pads prior to the inoculation were stained immunohistologically to examine for T-cell induction due to *M. leprae* infection. CD4+ and CD8+ lymphocytes were detected in the subcutaneous tissue and in the intermuscular layer as shown in Figure 3 but were not observed prior to the inoculation of *M. leprae* (data not shown).

**ELISA analyses of sera from *M. leprae*-inoculated mice.** Increases in the

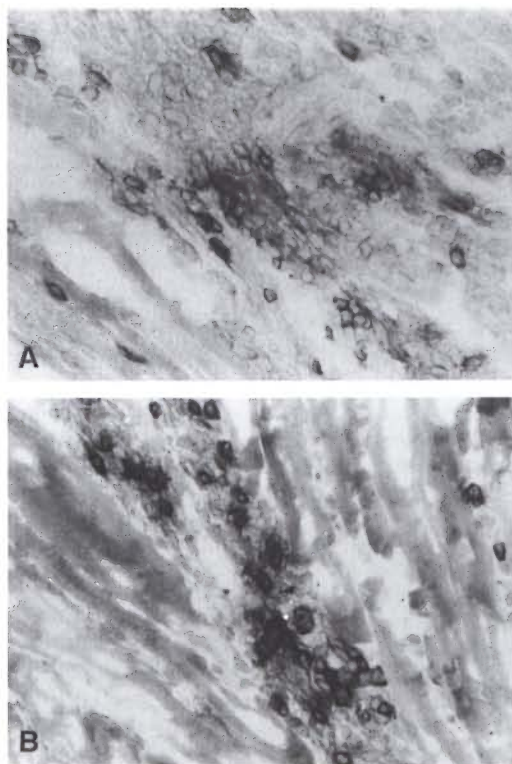


FIG. 3. Immunohistological stains of CD4 and CD8 lymphocytes in foot pads of a mouse on day 30 of infection with viable *M. leprae* showing CD4+ (A) and CD8+ (B) lymphocyte infiltration into the intermuscular layer ( $\times 400$ ).

IgM and IgG levels in the serum, due presumably to *M. leprae* inoculation, were observed 10 days after inoculation of *M. leprae* (The Table). There were no differences in increase of the antibody response to recombinant hsp65 between the groups infected with viable and heat-killed *M. leprae*. The antibody level to LAM-B antigen was evidently higher in the group inoculated with heat-killed *M. leprae* after 150 days of inoculation. No significant increases in antibody responses to PGL-I were observed in either group infected with viable or killed bacteria after 150 days of inoculation.

## DISCUSSION

Since Hansen first described *M. leprae* as the pathogenic organism of leprosy, numerous animal experiments have been conducted (<sup>21, 29</sup>). Shepard developed the foot



THE TABLE. ELISA analyses of sera from viable or heat-killed *M. leprae* inoculated mice.<sup>a</sup>

	IgM		IgG		65-kDa	LAM-B	PGL-I
Day	(μg/ml)				ELISA (OD)		
0	18.13 ±	1.63	89.69 ±	4.56	0.115 ± 0.015	0.172 ± 0.004	0.004 ± 0.001
Viable <i>M. leprae</i>							
1	16.22 ±	4.31	89.23 ±	15.80	0.134 ± 0.013	0.226 ± 0.030	−0.002 ± 0
10	27.63 ±	1.12	142.07 ±	10.57	0.109 ± 0.006	0.209 ± 0.008	−0.006 ± 0.003
30	38.30 ±	2.59	175.95 ±	10.08	0.221 ± 0.045	0.268 ± 0.001	−0.011 ± 0.013
150	343.14 ±	80.42	562.82 ±	113.0	0.269 ± 0.049	0.201 ± 0.026	0.007 ± 0
Heat-killed <i>M. leprae</i>							
1	20.15 ±	2.79	87.06 ±	11.96	0.129 ± 0.002	0.230 ± 0.009	0.010 ± 0.014
10	25.32 ±	0.82	125.29 ±	0.67	0.119 ± 0.007	0.218 ± 0.006	0.008 ± 0.004
30	42.39 ±	2.98	131.50 ±	8.57	0.139 ± 0.013	0.246 ± 0.028	−0.018 ± 0.021
150	145.46 ±	4.51	541.98 ±	38.60	0.308 ± 0.011	0.739 ± 0.031	0.033 ± 0.007

<sup>a</sup> Sera diluted at 1:1000 for IgM and IgG, 1:100 for hsp65-kDa, LAM-B, and PGL-I. Data shown as mean ± S.D. of two mice.

pad inoculation method in mice and described that the low temperature in the foot pad has been an important factor for *M. leprae* growth even though the growth was limited (<sup>39, 40</sup>). We injected *M. leprae* into BALB/cAJcl mice intravenously and through foot pad roots combination, and studied the patterns of cytokine mRNA expression in the spleen and the foot pad of the mice with *M. leprae* by the RT-PCR method, extracting total RNA from these tissues, and synthesizing them into cDNA using a M-MLV reverse transcriptase. The patterns of cytokine mRNAs expressed in the foot pad tissue which permits the restricted growth of *M. leprae* were markedly different from that found in the spleen which does not allow *M. leprae* multiplication.

Kita, *et al.* (<sup>15</sup>) reported on the physiological expression of cytokine mRNAs in various organs of the BALB/cAJcl mouse. We could demonstrate a presence of patterns similar to Kita's in the spleen of mice prior to the inoculation (day 0), namely, all of the cytokine genes examined were found (Fig. 2). In the foot pads of mice, only the physiological presence of IL-1- $\alpha$  and TNF- $\beta$  mRNA alone have been expressed (Fig. 1). Following the inoculation of viable or heat-killed *M. leprae*, the usual appearance of IL-1- $\alpha$  and TNF- $\beta$  mRNA was intensified in the foot pad on day 1, and also the new expression of TNF- $\alpha$  mRNA was observed. In contrast to the fact that TNF- $\alpha$  is pro-

duced from activated various cells such as macrophages, TNF- $\beta$  is considered to be produced from activated lymphocytes (<sup>28</sup>). In our results, the expression of mRNA for TNF- $\beta$  but not TNF- $\alpha$  was detected in the foot pad prior to the inoculation of *M. leprae* without detectable mRNA for CD4+ and CD8+ cells. However, as seen in Figure 3, on day 30 following the foot pad inoculation of *M. leprae*, the infiltration of CD4+ and CD8+ lymphocytes was confirmed by immunohistological staining. Here it is difficult to explain fully how a molecule can be expressed without a detectable mRNA, but it is considered that the number of infiltrated cells for CD4+ and CD8+/total number of cells in the foot pad may not be enough for the detection of the mRNA by the RT-PCR method. Further studies are necessary to elucidate the reasons.

The TNF- $\alpha$  was reported as a factor which induced hemorrhagic necrosis in the site of tumorigenesis (<sup>3</sup>), and it has begun to be construed as a cytokine closely related to defenses against pathogens (<sup>27, 46</sup>) and tumors (<sup>30, 44</sup>). In addition, IL-1- $\alpha$ , TNF- $\alpha$  and TNF- $\beta$  possess similar biological activities and are known to exert synergism (<sup>8</sup>). In our experiments, the expressions of mRNA for these cytokines were observed to be stronger in the foot pads of mice following the inoculation of heat-killed *M. leprae* on day 10 than that seen in the foot pads following viable *M. leprae* inoculation, while

the appearance of these cytokines was not changed in the spleens of mice inoculated with either viable or heat-killed *M. leprae* (Figs. 1 and 2).

The changes in the serum antibody titers of infected BALB/cAJcl mice were found from day 10 following the infection of *M. leprae* for the amount of IgM and IgG, and both of the specific antibodies against hsp65 kDa and LAM-B were found to increase when compared with titers prior to the infection. The titers of LAM-B were higher in the mice inoculated with heat-killed *M. leprae* than in those receiving viable bacteria on day 150 as shown in The Table. Although the elevation of the antibody level against PGL-I was not observed, the presence of PGL-I was confirmed by immunohistological stain in the tissues of viable *M. leprae* infection (data not shown). PGL-I<sup>(18, 23, 25, 45)</sup> and LAM-B<sup>(41, 42)</sup> are reported to be immunosuppressive antigens for cellular immunity, and hsp65<sup>(24)</sup> is reported to be a highly immunogenic antigen for humoral and cellular immunity. When the foot pads of the BALB/cAJcl (euthymic) mice were inoculated with a large amount of *M. leprae*, the bacilli remain for a period of time in the inoculation site of the foot pad, demonstrating the establishment of persistent and resistant infection, even though there is no increase in the number of bacilli as reported by Rees<sup>(32)</sup>. In the foot pad, cytokine production considered to eliminate the leprosy bacilli may be suppressed by PGL-I which is producing and releasing into tissues by viable *M. leprae*. In fact, PGL-I was detectable in the foot pad inoculated with *M. leprae* by immunohistological examination as mentioned above, and the antibody levels to LAM-B were evidently higher in the serum of the infected mice than those prior to the inoculation.

In contrast to the foot pad, all of the cytokine genes tested appeared in the spleen, except the disappearance of CD4 mRNA on day 1 of *M. leprae* infection accompanying the reduced expression of IL-2, -4, -6, and IL-12 mRNAs. The CD4 mRNA expression was recovered on day 10 of infection with viable *M. leprae*, with a corresponding increase in lymphokine mRNA expression which had been decreased on day 1. In other words, following the infection with

*M. leprae*, the expression of CD4 mRNA is lost and reduced immediately; at the same time the reduction of the appearance of lymphokine genes occurs.

The results of the present study suggest that *M. leprae* exert a direct effect on CD4 lymphocytes which is related to inhibiting the expression of cytokine genes for the elimination of bacilli from the host temporarily. The temporary loss of expression of CD4 mRNA and the loss of accompanying cytokine mRNA expressions in the spleen were stronger by the killed *M. leprae* inoculation than by the viable bacilli in the spleen. In the case of the killed bacilli inoculation, they never recovered up to 30 days of the post-inoculation, and the expression for IL-2 and IL-6 mRNAs never recovered, even the end of this experiment (Fig. 2). Based on these facts for cytokine patterns, in addition to the low temperature of the foot pad, as described by Shepard<sup>(39, 40)</sup>, it is considered that a long-term continuation specific to *M. leprae* infection is established in the foot pad.

Mitsuyama, *et al.* explained that killed bacteria have a weak ability to induce IL-1 production in macrophages so that T-cell activation cannot proceed, leading to a failure in eliciting delayed-type hypersensitivity (DTH)<sup>(19)</sup>. In *M. leprae* infection, however, IL-1- $\alpha$  mRNA was detected in both the foot pads and the spleens of mice inoculated with killed *M. leprae* the same as viable *M. leprae* in our results shown in Figures 1 and 2. On day 10 after infection, the expression of IL-1- $\alpha$  mRNA was stronger in the foot pad by the killed *M. leprae* infection. It has been reported that the DTH reactions have been induced in mice pre-inoculated with killed<sup>(17)</sup> or viable *M. leprae*<sup>(26)</sup> by challenge with *M. leprae* lysate.

Recently, the analysis for the experimental results using TNF- $\alpha$  knockout mice clarified the role of this cytokine against intracellular parasites. When TNF- $\alpha$  p55 knockout mice were inoculated intravenously with *Lysteria monocytogenes*, all of the knockout mice died within 6 days after the inoculation while all of the normal control mice were alive<sup>(31, 36)</sup>. However, these knockout mice were normal against virus infection, showing normal function for cytotoxic T lymphocytes. There is a report that the administration of IL-1- $\alpha$  into mice

inhibit the growth of intracellular parasites (<sup>7</sup>). In the intravenous inoculation of *Lysteria*, 90% of them are distributed in the liver, then killed and removed by Kupffer's cells. Thus, recent studies for the understanding of cellular parasitic infections indicated that nonspecific immunoregulation of macrophages is more important than the antigen-specific immune system by T lymphocytes (<sup>9,35</sup>). In the case of *M. leprae* which cannot multiply in the visceral organs, such as the spleen, similar removal mechanisms may also be established by the estimation of TNF- $\alpha$  mRNA patterns in our results (Figs. 1 and 2).

The cytokine genes, IL-1- $\alpha$ , TNF- $\alpha$  and TNF- $\beta$  mRNAs, which express in the foot pad of immunocompetent BALB/cAJcl mice infected with *M. leprae* were also observed in the foot pad of SCID and nude mice which are highly susceptible to *M. leprae* (<sup>51</sup>). It has been reported that IL-1 has the ability to promote the growth of pathogenic *Escherichia coli* (<sup>34</sup>). Beutler, *et al.* (<sup>2</sup>) have reported that a high degree of homology exists between the TNF and macrophage-secreted factor cachectin, and Kawakami and Hayata (<sup>14</sup>) have reported that TNF may be a potent cachexia or autoimmune inducing agent for the harmful excess of TNF production and, also, Amiri, *et al.* (<sup>1</sup>) have reported that schistosome utilizes TNF to maintain its progeny after oviposition. TNF- $\beta$  also has been shown to be an important factor in the construction of lymph nodes (<sup>43</sup>). In the mechanisms of *M. leprae* infection and immunity, whether these cytokines play only a role in the elimination of *M. leprae* remains to be elucidated. In immunocompetent mice, the only sites that are susceptible to *M. leprae* infection are skin tissues, including the foot pad. Although the significance of the expression of IL-1- $\alpha$ , TNF- $\alpha$  and TNF- $\beta$  mRNAs in the foot pads of these mice cannot be ascertained without further studies, our present study suggests that tissue-specific, local, immunologic characteristics are important in *M. leprae* growth, and the overall immunologic competence of the infected host exerts further influence on its multiplication.

### SUMMARY

The cytokine mRNAs expressed in the foot pads and spleens of BALB/cAJcl mice

infected with *Mycobacterium leprae* were studied by the reverse transcriptase-polymerase chain reaction (RT-PCR) method using cytokine-specific primers for interleukin-1 alpha (IL-1 $\alpha$ ), -2, -4, -6, -10, -12 (p40), gamma interferon (IFN- $\gamma$ ), tumor necrosis factor-alpha (TNF- $\alpha$ ), and TNF- $\beta$ , and then for CD4 and CD8 markers. The pattern of cytokine gene expression in the foot pad which supports *M. leprae* growth was different from the expression in the spleen which does not permit *M. leprae* multiplication in mice. Before BALB/cAJcl mice were infected with *M. leprae*, IL-1 $\alpha$  and TNF- $\beta$  mRNAs were expressed physiologically in the foot pad while all of the cytokine genes examined were expressed in the spleen. In the foot pads of mice inoculated with *M. leprae*, in addition to the physiological appearance of IL-1 $\alpha$  and TNF- $\beta$  mRNAs, these signals were intensified. TNF- $\alpha$  expression was induced by the infection. On the other hand, in the spleens of mice inoculated with *M. leprae*, CD4 mRNA expression disappeared on day 1 of the infection, which was accompanied by the reduced expression of IL-2, -4, -6, and -12 mRNAs. The recovery of CD4 mRNA expression at a latter stage was accompanied by a corresponding increase of the cytokine mRNA expression. It was suspected that these results might permit restricted growth of *M. leprae* in the foot pads of normal mice. Furthermore, our study suggests that tissue-specific, local, immunologic characteristics are important in *M. leprae* growth.

### RESUMEN

Utilizando la reacción en cadena de la polimerasa-transcriptasa reversa (RT-PCR) se estudiaron los mRNAs para citocinas expresados en las almohadillas plantares de ratones BALB/cAJcl infectados con *Mycobacterium leprae*. Se utilizaron sondas (iniciadores) para interleucina-1 alfa (IL- $\alpha$ ), IL-2, IL-4, IL-6, IL-10, IL-12 (p40), interferón gamma (IFN- $\gamma$ ), factor de necrosis tumoral alfa (TNF $\alpha$ ), TNF $\beta$ , y para los marcadores CD4 y CD8. El patrón de expresión de los genes para citocinas en la almohadilla plantar (que permite el crecimiento de *M. leprae*) fue diferente del patrón de expresión de citocinas en el bazo (el cual no permite la multiplicación de *M. leprae* en el ratón). Mientras que sólo los mRNAs para IL-1 $\alpha$  y TNF $\beta$  se expresaron fisiológicamente en las almohadillas plantares de los ratones BALB/cAJcl antes de su infección con *M. leprae*, todos los genes para citocinas examinados fueron expresados en los bazos. En las



almohadillas plantares de los ratones inoculados con *M. leprae*, además de la aparición fisiológica de los mRNAs para IL-1 $\alpha$  y TNF $\beta$ , estas señales fueron intensificadas. La expresión de TNF $\alpha$  fue inducida por la infección. Por otro lado, en los bazos de los animales inoculados con *M. leprae*, la expresión del mRNA para CD4 desapareció hacia el día 1 de la infección y esto se acompañó de la expresión reducida de los mRNAs para IL-2, IL-4, IL-6, e IL-12. La recuperación de la expresión del mRNA para CD4 en los estadios más avanzados de la infección, se acompañó del correspondiente incremento en la expresión de los mRNAs para las otras citocinas. Estos resultados podrían estar relacionados con el crecimiento restringido de *M. leprae* en las almohadillas plantares de los ratones normales. Además, nuestro estudio sugiere que las características locales y específicas de tejido influyen importante-mente en el crecimiento de *M. leprae*.

### RÉSUMÉ

Les ARNm de cytokine exprimés dans les coussinets plantaires et les rates de souris BALB/cAJcl infectées par *Mycobacterium leprae* ont été étudiés par la méthode de réaction réverse de transcriptase-polymerase en chaîne (RT-PCR), en utilisant des marqueurs spécifiques de cytokines pour l'interleukine-1 alpha (IL-1 $\alpha$ ), -2, -4, -6, -10, -12(p40), gamma interféron (IFN- $\gamma$ ), le facteur de nécrose tumorale alpha (FNT- $\alpha$ ), et le FNT- $\beta$ , et pour les marqueurs CD4 et CD8. Le mode d'expression du gène de cytokine dans le coussinet plantaire qui supporte la croissance de *M. leprae* était différent de l'expression dans la rate qui ne permet pas la multiplication de *M. leprae* chez la souris. Avant que les souris BALB/cAJcl ne soient infectées par *M. leprae*, les ARNm d'IL-1 $\alpha$  et FNT- $\beta$  ont été exprimés physiologiquement dans le coussinet plantaire alors que tous les gènes de cytokines examinés ont été exprimés dans la rate. Dans les coussinets plantaires des souris inoculées avec *M. leprae*, en plus de l'apparition physiologique des ARNm d'IL-1 $\alpha$  et FNT- $\beta$ , ces signaux ont été intensifiés. L'expression de FNT- $\alpha$  était induite par l'infection. D'autre part, dans les rates de souris infectées par *M. leprae*, l'expression de l'ARNm de CD4 disparut au premier jour de l'infection, qui était accompagnée d'une réduction de l'expression des ARNm d'IL-2, -4, -6 et -12. Le rétablissement de l'expression de l'ARNm de CD4 à un stade plus tardif était accompagné d'une augmentation correspondante de l'expression de l'ARNm de cytokine. On a soupçonné que ces résultats pourraient permettre une croissance limitée de *M. leprae* dans les coussinets plantaires de souris normales. De plus, notre étude suggère que les caractéristiques locales, spécifiques par tissu, sont importantes pour la croissance de *M. leprae*.

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