

Mycobacterium avium subsp. *paratuberculosis* Infection Causes Suppression of RANTES, Monocyte Chemoattractant Protein 1, and Tumor Necrosis Factor Alpha Expression in Peripheral Blood of Experimentally Infected Cattle

Joram J. Buza,¹ Yasuyuki Mori,² Abusaleh M. Bari,¹ Hirokazu Hikono,¹ Aodon-geril,¹ Sachiyo Hirayama,¹ Yujing Shu,¹ and Eiichi Momotani^{1*}

Paratuberculosis and Inflammatory Bowel Disease Research Team¹ and Immune System Section,² National Institute of Animal Health, 3-1-5 Kan-nondai, Tsukuba 305-0856, Japan

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Blood from cattle with subclinical *Mycobacterium avium* subsp. *paratuberculosis* infection was stimulated with *M. avium* subsp. *paratuberculosis* antigens, and expression of interleukin-1 β (IL-1 β), tumor necrosis factor alpha (TNF- α), RANTES, monocyte chemoattractant protein 1 (MCP-1), and IL-8 was measured. Expression of TNF- α , RANTES, and MCP-1 was lower in infected than in uninfected cattle. The reduced response may weaken protective immunity and perpetuate infection.

Information is now accumulating on cellular and cytokine changes associated with *Mycobacterium avium* subsp. *paratuberculosis* infection in cattle (8, 17–20), but little or no information is available on chemokines, which are involved in granuloma formation. Chemokines are chemotactic cytokines with small molecular size (8 to 14 kDa) that mediate the recruitment of leukocytes from blood to tissues in maintenance of health or in response to inflammation (4, 11). Four subfamilies of chemokines, namely CC, CXC, C, and CX₃C, have been designated based on the presence of a cysteine-containing signature motif at the amino-terminal end (11). The response of chemokines to *Mycobacterium tuberculosis* infection is induced by inflammatory cytokines among other factors (8, 10). The chemokines monocyte chemoattractant protein 1 (MCP-1 [CCL2]), RANTES (CCL5), and interleukin-8 (IL-8 [CXCL8]) are suggested to play a role in formation of granuloma to *M. tuberculosis* (6, 9, 18); however, no information is available on their responses to *M. avium* subsp. *paratuberculosis* infection. In the present study, we evaluated mRNA expression of the chemokines RANTES, MCP-1, and IL-8 and the proinflammatory cytokines IL-1 β and tumor necrosis factor alpha (TNF- α) during *M. avium* subsp. *paratuberculosis* infection in cattle.

The experimental animals included five 25-month-old castrated male Holstein calves and five age-matched uninfected control animals. The infected animals were orally inoculated with 20.7×10^8 CFU of *M. avium* subsp. *paratuberculosis* at 1 week of age. At the beginning of this study, the animals were positive for *M. avium* subsp. *paratuberculosis* infection in the gamma interferon (IFN- γ) enzyme-linked immunosorbent assay (2) but were negative for clinical signs, *M. avium* subsp. *paratuberculosis*-specific antibodies, and *M. avium* subsp. *paratuberculosis* shedding in feces by PCR and culture. Jugular blood was dispensed at 1 ml per well of a 24-well tissue culture

plate and stimulated with different stimulants to a final concentration of 10 μ g of lipopolysaccharide (LPS) per ml from *Escherichia coli* (Sigma), 10 μ g of *M. avium* subsp. *paratuberculosis* lysate per ml, 0.5 μ g of *M. avium* subsp. *paratuberculosis* PPD per ml or 60×10^6 CFU of live *M. avium* subsp. *paratuberculosis*. After 0, 1, 3, 6, 12, and 24 h of incubation in a humidified incubator at 37°C and 5% CO₂ in air, plasma was separated from the blood cell fraction and tested for IFN- γ with the BioX ELISA kit (Marche-en-Famenne, Belgium). The blood cell fraction was used for RNA isolation with the Trizol reagent (Life Technologies, Rockville, Md.) according to manufacturer's instructions. Real-time reverse transcription-PCR (RT-PCR) was done with the SYBR Green RT-PCR kit (Qiagen, Tokyo, Japan). Primers were designed from the

TABLE 1. Primers used for real-time RT-PCR

Primer	Sequence (5' to 3')	Product size (bp)	GenBank accession no.
IL-1 β			
Sense	ATCTTCGAAACGTCCTCCGAC	187	M37211
Antisense	CCTCTCCTTGACAAAGCTCA		
TNF- α			
Sense	CCATCAACAGCCCTCTGGTT	138	AF011926
Antisense	CCATGAGGGCATTGGCATA		
RANTES			
Sense	GCCCTGCTGCTTTCCTATAT	190	BTA7043
Antisense	TCCACCCTAGCTCAACTCAA		
MCP-1			
Sense	CCTCCTGTGCCTGCTACTCA	236	L32659
Antisense	CTGGACCCATTCTGCCTGG		
IL-8			
Sense	CCACACCTTTCTACCCCAA	142	AF061524
Antisense	CCTTCTGCACCCACTTTTCCT		
GAPDH			
Sense	GGCGTGAACCACGAGAAGTATAA	118	U85042
Antisense	CCTCCACGATGCCAAAGTG		

* Corresponding author. Mailing address: Paratuberculosis and Inflammatory Bowel Disease Research Team, NIAH, 3-1-5 Kan-nondai, Tsukuba 305-0856, Japan. Phone: 81-29-838-7781. Fax: 81-29-838-7781. E-mail: momotani@affrc.go.jp.

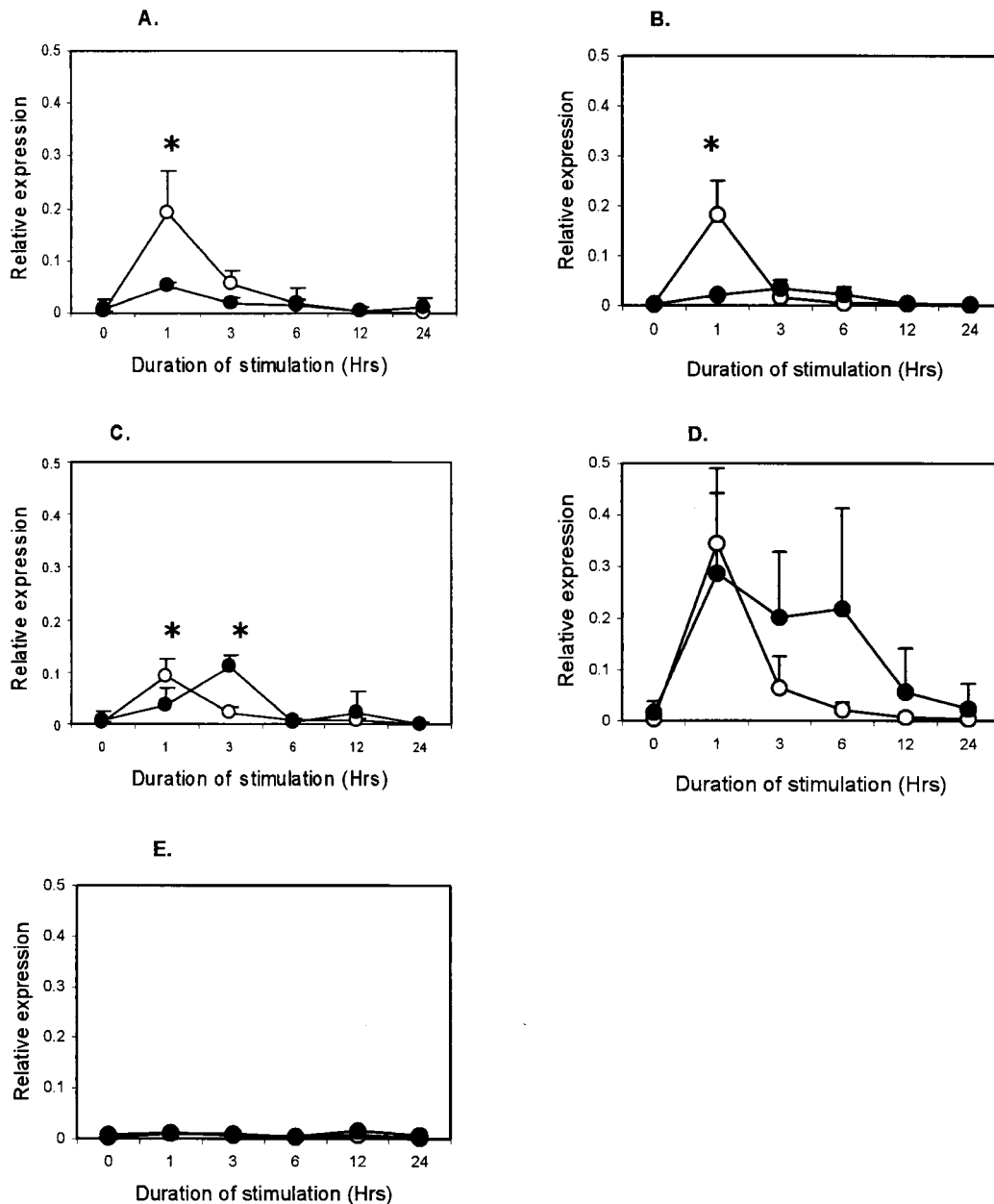


FIG. 1. Kinetics of TNF- α in whole-blood samples from *M. avium* subsp. *paratuberculosis*-infected (●) and uninfected (○) cattle after in vitro stimulation with 0.5 μ g of *M. avium* subsp. *paratuberculosis* PPD per ml (A), 10 μ g of *M. avium* subsp. *paratuberculosis* lysate per ml (B), 60×10^6 live *M. avium* subsp. *paratuberculosis* CFU (C), 10 μ g of *E. coli* LPS per ml (D), and medium (E). Each point represents the mean \pm standard error of five animals. Differences between groups were considered significant when $P < 0.05$ (*).

published bovine genome by using Primer Express version 1.5 software (Applied Biosystems, Foster City, Calif.) and are described in Table 1. The 25- μ l one-step RT-PCR mixture consisted of 12.5 μ l of master mix, 0.5 μ l each of a forward and reverse primer, 0.25 μ l of RT mix, 10 μ l of RNase-free water, and 1.25 μ l of template RNA (containing 500 to 1,500 ng). The real-time cycler conditions included 50°C for 30 min and 95°C for 15 min followed by 40 cycles of 94°C for 15 s, 60°C for 30 s, and 72°C for 30 s. Relative quantities for different cytokines were extrapolated from the standard curve generated from 10-fold dilutions of the reference mRNA sample (6-h LPS-

stimulated whole blood) and then normalized for GAPDH expression as previously described (<http://www.ambion.com/techlib/tn/85/857.html>). Data were analyzed by analysis of variance with repeated measures. P values of < 0.05 were considered significant.

M. avium subsp. *paratuberculosis* antigens induced IFN- γ production in plasma from infected animals ($P < 0.001$), but not from uninfected animals (data not shown), confirming the infection status (2, 16). To the contrary, mycobacterial antigens induced lower TNF- α , RANTES, and MCP-1 responses in the infected animals. Responses for IL-1 β and IL-8 were not dif-

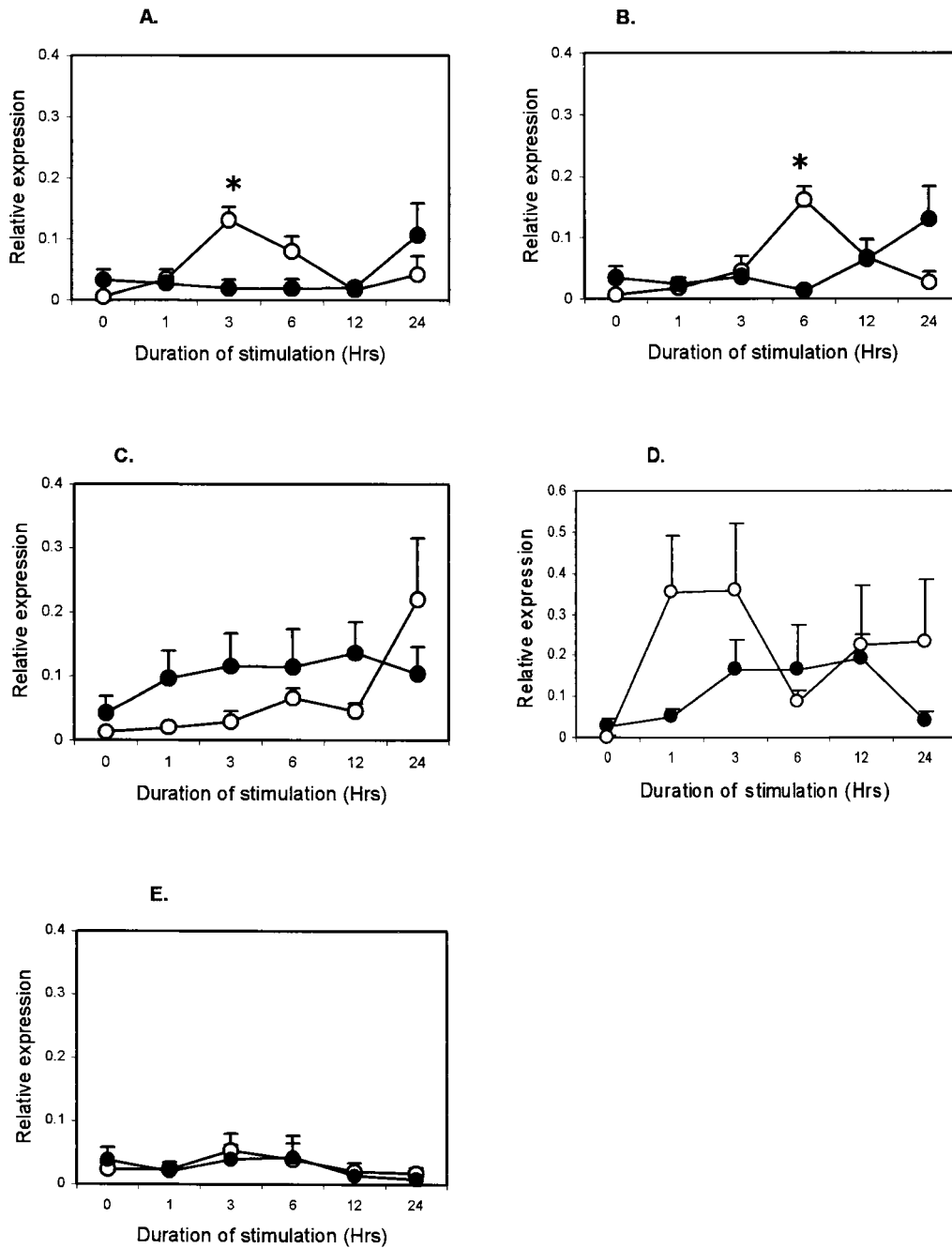


FIG. 2. Kinetics of RANTES in whole-blood samples from *M. avium* subsp. *paratuberculosis*-infected (●) and uninfected (○) cattle after in vitro stimulation with 0.5 μg of *M. avium* subsp. *paratuberculosis* PPD per ml (A), 10 μg of *M. avium* subsp. *paratuberculosis* lysate per ml (B), 60 × 10⁶ live *M. avium* subsp. *paratuberculosis* CFU (C), 10 μg of *E. coli* LPS per ml (D), and medium (E). Each point represents the mean ± standard error of five animals. Differences between groups were considered significant when $P < 0.05$ (*).

ferent between groups (data not shown). The TNF-α expression in infected cattle was lower ($P < 0.001$) at 1 h after purified protein derivative (PPD) (Fig. 1A), *M. avium* subsp. *paratuberculosis* lysate (Fig. 1B), and live *M. avium* subsp. *paratuberculosis* stimulation (Fig. 1C). Live *M. avium* subsp. *paratuberculosis*, however, induced higher ($P < 0.001$) TNF-α expression in infected animals at 3 h. Expression of RANTES in the infected group was lower ($P < 0.01$) at 3 h after PPD stimulation (Fig. 2A) and at 6 h after *M. avium* subsp. *paratu-*

erculosis lysate stimulation ($P < 0.001$) (Fig. 2B), but live *M. avium* subsp. *paratuberculosis* did not confer differential expression (Fig. 2C). MCP-1 expression was lower ($P < 0.01$) in infected animals at 3 h after stimulation with PPD (Fig. 3A) but was much lower ($P < 0.001$) at 6 and 12 h after stimulation with *M. avium* subsp. *paratuberculosis* lysate (Fig. 3B) and live *M. avium* subsp. *paratuberculosis* (Fig. 3C). Stimulation with LPS, a non-*M. avium* subsp. *paratuberculosis* antigen, failed to induce differential expression between groups (Fig. 1D, 2D,

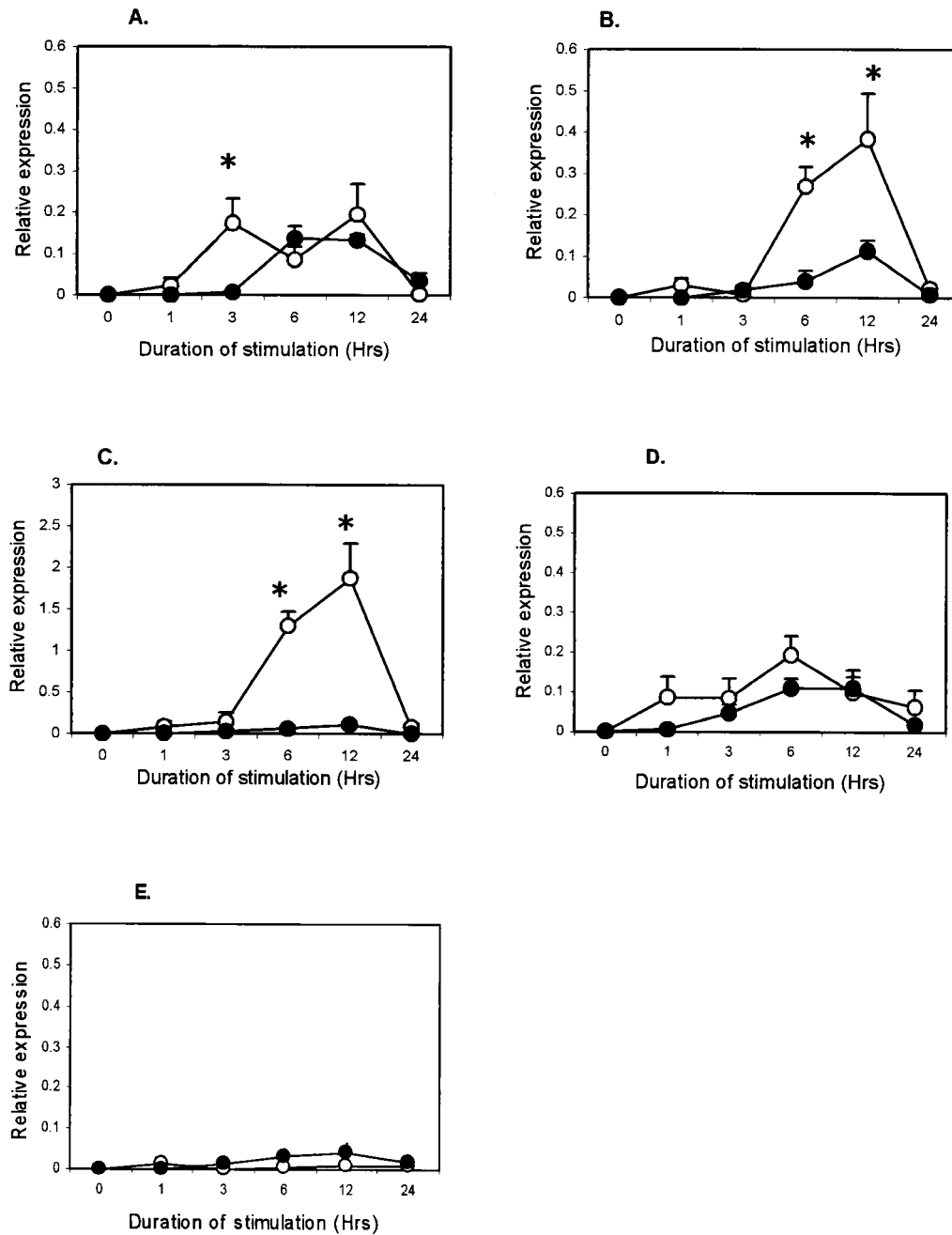


FIG. 3. Kinetics of MCP-1 in whole-blood samples from *M. avium* subsp. *paratuberculosis*-infected (●) and uninfected (○) cattle after in vitro stimulation with 0.5 µg of *M. avium* subsp. *paratuberculosis* PPD per ml (A), 10 µg of *M. avium* subsp. *paratuberculosis* lysate per ml (B), 60×10^6 live *M. avium* subsp. *paratuberculosis* CFU (C), 10 µg of *E. coli* LPS per ml (D), and medium (E). Each point represents the mean \pm standard error of five animals. Differences between groups were considered significant when $P < 0.05$ (*).

and 3D), and unstimulated samples did not show change in expression (Fig. 1E, 2E, and 3E).

The reduced TNF- α , RANTES, and MCP-1 responses observed may have a negative effect on granuloma formation and function. TNF- α plays an important role in the host defense against mycobacterium infections, including the initiation and formation of granulomas and activation of antimycobacterial killing mechanisms (3, 5, 7, 13). RANTES and MCP-1 contrib-

ute to granuloma formation through chemotactic effects on T cells, monocytes, and macrophages (1, 9, 14, 15, 21). Our results concur with observations that fewer if any granulomas are found in cattle during the subclinical stage of *M. avium* subsp. *paratuberculosis* infection (3, 5, 22). On the other hand, increased MCP-1 expression during *M. avium* subsp. *paratuberculosis* infection in mice was associated with high numbers of granulomas in liver, indicating its direct association with

granuloma formation (E. Momotani, M. Miyama, T. L. To, K. Yoshihara, and H. Gotoh, 14th European Immunology Meeting, Poznan, Poland, Abstr. Immunol. Lett., abstr. 490, 2000). Interestingly, the reduced levels of TNF- α , RANTES, and MCP-1 expression occurred after stimulation with *M. avium* subsp. *paratuberculosis* antigens rather than with a non-*M. avium* subsp. *paratuberculosis* antigen LPS, suggesting the presence of suppressive substances emanating from the specific response. In this regard, it has been suggested that IL-10 secreted by *M. avium* subsp. *paratuberculosis*-infected macrophages suppresses TNF- α expression (20). At the same time, induction of RANTES (11) and MCP-1 (12, 13) by mycobacterial antigens is mediated by TNF- α . It is possible therefore that *M. avium* subsp. *paratuberculosis* antigens induced IL-10, which in turn suppressed TNF- α production with subsequent reduction in RANTES and MCP-1 expression. Despite an increased cell-mediated immune response during the subclinical stage of *M. avium* subsp. *paratuberculosis* infection in cattle, the infection persists in a number of cases, ending with precipitation of the clinical disease (18). The suppression of TNF- α , RANTES, and MCP-1, among other factors, may compromise this response, allowing *M. avium* subsp. *paratuberculosis* to escape destruction. Further studies are still needed to directly correlate peripheral chemokine responses with intestinal granuloma formation.

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