

Importance of murine V δ 1⁺ $\gamma\delta$ T cells expressing interferon- γ and interleukin-17A in innate protection against *Listeria monocytogenes* infection

Satoru Hamada,^{1,2} Masayuki Umemura,^{1,3} Takeru Shiono,^{1,3} Hiromitsu Hara,⁴ Kenji Kishihara,⁵ Kensho Tanaka,¹ Hirokazu Mayuzumi,^{1,3} Takao Ohta² and Goro Matsuzaki^{1,3}

¹Molecular Microbiology Group, Centre of Molecular Biosciences, University of the Ryukyus, Okinawa, Japan, ²Division of Child Health and Welfare, ³Division of Host Defence and Vaccinology, University of the Ryukyus, Okinawa, Japan, ⁴Division of Molecular and Cellular Immunoscience, Department of Biomolecular Sciences, Saga University School of Medicine, Saga, Japan, and ⁵Department of Immunology and Parasitology, Institute of Health Bioscience, University of Tokushima School of Medicine, Tokushima, Japan

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Correspondence: Dr G. Matsuzaki, Molecular Microbiology Group, Centre of Molecular Biosciences, University of the Ryukyus, Nishihara, Okinawa 903-0213, Japan.
Email: matsuzak@comb.u-ryukyu.ac.jp
Senior author: Goro Matsuzaki

Introduction

T cells expressing the T-cell receptor (TCR)- $\gamma\delta$ heterodimer belong to a unique T-cell lineage distinct from conventional T cells expressing the TCR- $\alpha\beta$ heterodimer.¹ A restricted number of TCR V γ and V δ genes is available to encode functional TCR- $\gamma\delta$, suggesting that the $\gamma\delta$ T cells recognize a restricted number of ligands. However, the ligand specificity of the majority of $\gamma\delta$ T cells has not been determined.² $\gamma\delta$ T cells can be divided into functional subpopulations, which are similar to the T helper type 1 (Th1), Th2, Th3, Th17 and cytotoxic T cells of

Summary

Murine $\gamma\delta$ T cells participate in the innate immune response against infection by an intracellular pathogen *Listeria monocytogenes*. V δ 1⁺ $\gamma\delta$ T cells coexpressing V γ 6 are a major $\gamma\delta$ T-cell subpopulation induced at an early stage of *L. monocytogenes* infection in the livers of infected mice. To investigate the protective role of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells against *L. monocytogenes* infection, V δ 1 gene-deficient (V δ 1^{-/-}) mice were analysed because these mice selectively lacked a V γ 6/V δ 1⁺ $\gamma\delta$ T-cell subpopulation in the *L. monocytogenes*-infected liver. The V δ 1^{-/-} mice showed increased bacterial burden in the liver and spleen, and decreased survival rate at an early stage of *L. monocytogenes* infection when compared to wild-type mice. Histological examination showed abscess-like lesions and unorganized distribution of macrophages in the liver of the V δ 1^{-/-} mice but not in the wild-type mice after *L. monocytogenes* infection. The V γ 6/V δ 1⁺ $\gamma\delta$ T cells produced interferon- γ and interleukin-17A. All the results suggest that murine V γ 6/V δ 1⁺ $\gamma\delta$ T cells control the innate protective response against *L. monocytogenes* infection through production of the proinflammatory cytokines interferon- γ and interleukin-17A in the infected liver.

Keywords: interferon- γ ; interleukin-17A; *Listeria monocytogenes*; $\gamma\delta$ T cell

TCR- $\alpha\beta$ T cells,²⁻⁶ while the role of the $\gamma\delta$ T cells in *in vivo* immune responses is not well characterized.

The $\gamma\delta$ T cells have been reported to participate in immune responses against various infections.⁷ *Listeria monocytogenes*, an intracellular bacterium, has been extensively used in analyses of the role of $\gamma\delta$ T cells, and the importance of $\gamma\delta$ T cells in innate protective immunity against this infection is well established.⁸⁻¹⁰ Murine $\gamma\delta$ T cells induced by the *L. monocytogenes* infection can be divided into three subpopulations, V γ 1/V δ 6⁺,¹¹⁻¹³ V γ 4⁺,¹² and V γ 6/V δ 1⁺,^{14,15} $\gamma\delta$ T cells, according to their TCR V γ /V δ expression. The protective role

of the V γ 1⁺ $\gamma\delta$ T cells has been demonstrated,¹² although another report indicated an opposite role for V γ 1⁺ T cells.¹³ V γ 1⁺ $\gamma\delta$ T-cell-mediated killing of activated macrophages has also been demonstrated.¹⁶ Depletion of V γ 4⁺ $\gamma\delta$ T cells has no significant effect on protective immunity against *L. monocytogenes* infection, suggesting that V γ 4 does not have a significant role in protective immunity.¹³

To further clarify the repertoire-specific function of the $\gamma\delta$ T cells in protection against *L. monocytogenes* infection, we analysed V δ 1 gene-deficient (V δ 1^{-/-}) mice.¹⁷ The results demonstrated that the V δ 1^{-/-} mice selectively lacked V γ 6/V δ 1⁺ $\gamma\delta$ T cells but retained V γ 1⁺ and V γ 4⁺ $\gamma\delta$ T cells in the liver after *L. monocytogenes* infection, and that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells had an important role in innate protective immunity against *L. monocytogenes* infection.

Materials and methods

Animals

Wild-type C57BL/6 mice were purchased from Japan SLC (Shizuoka, Japan). V δ 1^{-/-} mice¹⁷ were backcrossed more than eight times to C57BL/6 mice. The mice were maintained in a conventional environment, and were used for experiments at 8–10 weeks of age. Experiments were conducted according to the Institutional Ethical Guidelines for Animal Experiments and the Safety Guideline for Gene Manipulation Experiments of the University of the Ryukyus.

Microorganisms and bacterial infection

Listeria monocytogenes strain EGD was inoculated into C57BL/6 mice, fresh isolates were obtained from infected spleens, grown in tryptic soy broth (Difco, Detroit, MI), resuspended in phosphate-buffered saline, and stored at -80° in small samples until use. Mice were infected by intraperitoneal (i.p.) inoculation of 5 × 10⁴ colony-forming units of *L. monocytogenes* [which corresponds to 1/10 of the 50% lethal dose (LD₅₀) for wild-type C57BL/6 mice] except for the analysis of survival rate, for which 2.5 × 10⁵ colony-forming units of *L. monocytogenes* was inoculated.

Cell preparation

Liver mononuclear cells were prepared as described previously.¹⁸ To enrich TCR- $\gamma\delta$ T cells, the cells were passed through nylon wool columns, then separated using a magnetic cell sorter system (autoMACS, Miltenyi, Bergisch Gladbach, Germany) by using biotin- or fluorescein isothiocyanate (FITC)-conjugated anti-TCR C δ monoclonal antibody (mAb; GL-3, Becton Dickinson, San Jose, CA)

and streptavidin or anti-FITC microbeads (Miltenyi), respectively.

Gene expression analysis by reverse transcription-polymerase chain reaction (RT-PCR)

Using Trizol reagent (Invitrogen, Carlsbad, CA), total RNA was extracted. First-strand cDNA was synthesized, and then amplified by PCR using *Taq* polymerase (Takara Shuzo, Kyoto, Japan) with TCR V γ or V δ sense primers and C γ or C δ antisense primers, as described previously.¹⁹ The following primers were used (nomenclature of V γ genes is according to Heilig and Tonegawa²⁰): C γ , CTT ATG GAG ATT TGT TTC AGC; V γ 1-3, ACA CAG CTA TAC ATT GGT AC; V γ 2, CGG CAA AAA ACA AAT CAA CAG; V γ 4, TGT CCT TGC AAC CCC TAC CC; V γ 5, TGT GCA CTG GTA CCA ACT GA; V γ 6, CTC CAA AGA ATG CTG TGT AG; V γ 7, AAG CTA GAG GGG TCC TCT GC; C δ , CGA ATT CCA CAA TCT TCT TG; V δ 1, ATT CAG AAG GCA ACA ATG AAA G; V δ 2, AGT TCC CTG CAG ATC CAA GC; V δ 3, TTC CTG GCT ATT GCC TCT GAC; V δ 4, CCG CTT CTC TGT GAA CTT CC; V δ 5, CAG ATC CTT CCA GTT CAT CC; V δ 6, CTT AGT GGA GAG ATG GTT TT; V δ 7, CGC AGA GCT GCA GTG TAA CT; and V δ 8, AAG GAA GAT GGA CGA TTC AC. The V γ 1-3 primer amplifies the V γ 1, V γ 2 and V γ 3 genes, whereas the V γ 2 primer amplifies the V γ 2 gene but not the V γ 1 gene. Since $\gamma\delta$ T cells with functional V γ 3 expression have not yet been documented, we consider that the V γ 1-3 primer detects TCR of V γ 1⁺ and V γ 2⁺ $\gamma\delta$ T cells. The PCR products were electrophoresed through 1.8% agarose gel, stained with ethidium bromide, and photographed using the Gel-Documentsation system (BioRad, Hercules, CA).

Flow cytometry (FCM)

Cells were stained with mAbs against leucocyte surface molecules and cytokines, and analysed by FCM. The following mAbs were used to detect surface molecules: FITC-conjugated anti-CD3, anti-Gr1 (Becton Dickinson), and anti-TCR C β (Becton Dickinson) mAbs, phycoerythrin-conjugated anti-TCR C δ , anti-CD11b (Caltag, Burlingame, CA), anti-CD4 (Becton Dickinson), anti-CD8 (Becton Dickinson) and anti-NK1.1 (Becton Dickinson) mAbs. After staining, cells were analysed using a FACSCalibur system (Becton Dickinson), and the proportions of macrophages (Gr1⁻ CD11b⁺), neutrophils (Gr1⁺ CD11b⁺), TCR $\gamma\delta$ T cells (CD3⁺ TCR C δ ⁺), CD4⁺ T cells (CD4⁺ TCR C β ⁺), CD8⁺ T cells (CD8⁺ TCR C β ⁺) and natural killer T cells (NK1.1⁺ TCR C β ⁺) were determined. The absolute number of each fraction was calculated using absolute number of liver mononuclear cells per mouse and the ratio of each cell population. To detect cytokine expression of $\gamma\delta$ T cells, the $\gamma\delta$ T cells were enriched with biotin-conjugated anti-TCR C δ mAb and autoMACS, cultured

with 1 $\mu\text{g/ml}$ calcium ionophore, 25 ng/ml phorbol 12-myristate 13-acetate, and brefeldin A for 4 hr, stained with mAb against surface antigens and then treated with Cytofix/Cytoperm solution (Becton Dickinson) according to the manufacturer's instructions followed by intracellular cytokine staining with phycoerythrin-conjugated anti-interleukin-17A (IL-17A), or phycoerythrin-conjugated anti-interferon- γ (IFN- γ) mAbs. To detect biotin-conjugated anti-TCR C δ mAb, allophycocyanin-conjugated streptavidin was used. FITC-conjugated anti-TCR V γ 1 (clone 2.11²¹), anti-TCR V γ 4 (Becton Dickinson), anti-TCR V γ 5 (Becton Dickinson) and anti-TCR V γ 7 (clone F2.67²²) mAbs were mixed (anti-V γ mAb mixture) and used to detect V γ 6⁺ and V γ 2⁺ $\gamma\delta$ T cells as anti-V γ mAb mixture-negative TCR C δ -positive cells.

Bacteria counts in organs

The *L. monocytogenes*-infected mice were killed on day 5 of the infection, the liver and spleen were homogenized in saline, and the homogenates were plated on nutrient agar plates containing 0.4% glucose to calculate the number of bacteria in each organ.

Histopathology

The livers of *L. monocytogenes*-infected mice were fixed in buffered formalin on day 5 of the infection, and embedded in paraffin. Thin sections were prepared and stained with haematoxylin & eosin. The stained sections were examined under a BX41 microscope (Olympus, Tokyo, Japan) equipped with 4 \times /0.13 and 20 \times /0.50 objectives. Images were acquired with a DP70 digital camera and DP software (Olympus). The liver was also embedded in optimal cutting temperature compound (Sakura, Tokyo, Japan), and frozen in dry ice-acetone. Thin sections were prepared with a cryostat, stained with allophycocyanin-conjugated anti-CD3 mAb and Alexa 488-conjugated anti-CD11b mAb (Becton Dickinson), and analysed using a Radiance 2100 (BioRad) confocal laser scanning microscope (CF-LSM) equipped with a 20 \times /0.70 objective. The images of CF-LSM were acquired with LASERSHARP 2000 software, (BioRad) and merged using ADOBE PHOTOSHOP software (Adobe, San Jose, CA).

Statistics

Data were statistically evaluated using Student's *t*-test and STATWORK software (Cricket Software, Philadelphia, PA). The survival rate was analysed by the Kaplan-Meier method and statistically evaluated by Log rank test using STATCEL 2 software (MOS, Saitama, Japan). A *P* value < 0.05 was considered to indicate statistical significance.

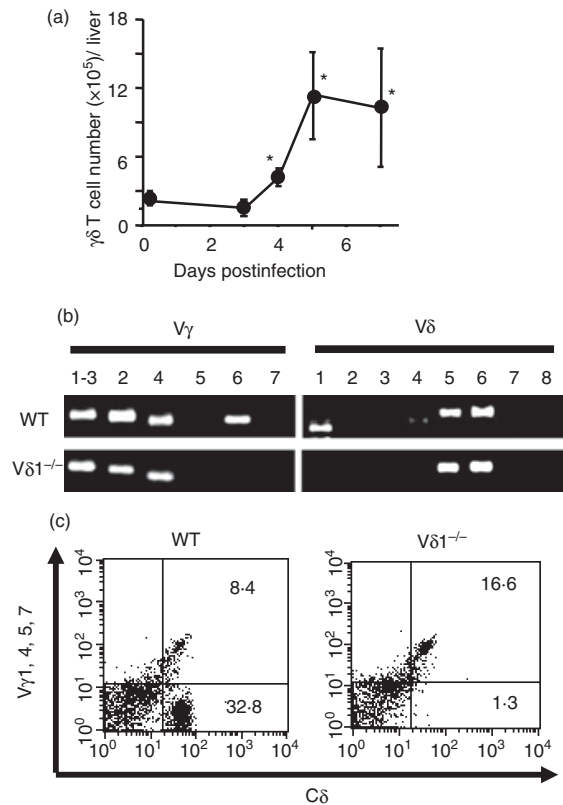


Figure 1. T-cell receptor (TCR) V region repertoire of the $\gamma\delta$ T cells induced in the liver of *Listeria monocytogenes*-infected mice. (a) Wild-type C57BL/6 mice were intraperitoneally inoculated with *L. monocytogenes*, and kinetics of the liver $\gamma\delta$ T-cell number was calculated using total liver mononuclear cell number and the ratio of the $\gamma\delta$ T cells determined by flow cytometry (FCM). The data shown are representative of three independent experiments. (b) V γ and V δ gene expression of the $\gamma\delta$ T cells were analysed by reverse transcription-polymerase chain reaction on the liver mononuclear cells from wild-type (WT) or V δ 1^{-/-} mice on day 5 of *L. monocytogenes* infection. The data are representative of three independent experiments. (c) Liver $\gamma\delta$ T cells were analysed on their expression of V γ gene products by FCM. The liver $\gamma\delta$ T cells were enriched from WT or V δ 1^{-/-} mice on day 5 of *L. monocytogenes* intraperitoneal infection, and were stained with anti-C δ monoclonal antibody (mAb) and anti-V γ mAb mixture as described in the *Materials and methods* section. The cells in the lower right quadrant correspond to the $\gamma\delta$ T cells that lack expression of V γ 1, V γ 4, V γ 5 and V γ 7. The experiments were repeated three to five times, and representative FCM profiles are shown.

Results

Increase of $\gamma\delta$ T cells expressing TCR V γ 6/V δ 1 in the liver of *L. monocytogenes*-infected mice and lack of the $\gamma\delta$ T-cell subpopulation in the V δ 1^{-/-} mice

It has been reported that $\gamma\delta$ T cells increase and participate in innate protective immunity against *L. monocytogenes*.

genes infection.^{8–10} In the liver of wild-type C57BL/6 mice, $\gamma\delta$ T cells increased from day 4 and reached a maximum on day 5 of *L. monocytogenes* infection (Fig. 1a). Analysis of V γ and V δ gene expression by the *L. monocytogenes*-induced liver $\gamma\delta$ T cells using RT-PCR showed that the liver $\gamma\delta$ T cells expressed V γ 1 and/or V γ 2, V γ 4 and V γ 6 as V γ genes, and V δ 1, V δ 5 and V δ 6 as V δ genes (Fig. 1b, upper panels). The data were consistent with a previous report showing the presence of V γ 1/V δ 6⁺, V γ 4⁺, and V γ 6/V δ 1⁺ $\gamma\delta$ T cells in the *L. monocytogenes*-infected mice.^{11–16} When the liver $\gamma\delta$ T cells were stained with an anti-V γ (anti-V γ 1, anti-V γ 4, anti-V γ 5 and anti-V γ 7) mAb mixture, 50–80% of the $\gamma\delta$ T cells were not stained (Fig. 1c, left panel), suggesting the presence of a high percentage of $\gamma\delta$ T cells expressing V γ 6 or V γ 2.

Selective pairing of V δ 1 with V γ 5 and V γ 6 has been reported.²³ Analysis using RT-PCR showed that V γ 6 expression, but not V γ 5 expression, was detected in the liver $\gamma\delta$ T cells from *L. monocytogenes*-infected wild-type mice (Fig. 1b, upper panel). Analysis by FCM using anti-TCR V γ 5 mAb also failed to detect V γ 5⁺ $\gamma\delta$ T cells (data not shown). Therefore, we reasoned that V δ 1^{-/-} mice selectively lack the V γ 6/V δ 1⁺ $\gamma\delta$ T cells in the liver $\gamma\delta$ T cells after *L. monocytogenes* infection. Consistent with this theory – not only the V δ 1 gene but also the V γ 6 genes were undetectable in the liver $\gamma\delta$ T cells of the V δ 1^{-/-} mice while V γ 2 gene expression was not affected (Fig. 1b, lower panels). Analysis by FCM of the liver $\gamma\delta$ T cells in the V δ 1^{-/-} mice showed that the ratio of anti-V γ mAb mixture-negative $\gamma\delta$ T cells decreased to 5–15% of the $\gamma\delta$ T cells (Fig. 1c, right panel), indicating that the majority of the anti-V γ mAb mixture-negative $\gamma\delta$ T cells in the *L. monocytogenes*-infected liver of wild-type mice were V γ 6/V δ 1⁺ $\gamma\delta$ T cells. All the results demonstrated that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells formed a major subpopulation of the $\gamma\delta$ T cells induced in the liver by *L. monocytogenes* infection, and that the V δ 1^{-/-} mice selectively lacked this $\gamma\delta$ T-cell subpopulation.

Protective role of V γ 6/V δ 1⁺ $\gamma\delta$ T cells against *L. monocytogenes* at an early stage of infection

To analyse the protective role of V γ 6/V δ 1⁺ $\gamma\delta$ T cells against *L. monocytogenes* infection, the bacterial count in the liver and spleen was determined on day 5 of infection. As shown in Fig. 2(a), bacterial burden in the liver and spleen significantly increased in the V δ 1^{-/-} mice compared to the wild-type mice. When a higher dose of *L. monocytogenes* (1/2 LD₅₀ for the wild-type C57BL/6 mice) was inoculated, half of the V δ 1^{-/-} mice died within 10 days after the infection while all of the wild-type mice survived (Fig. 2b). These results suggested that V γ 6/V δ 1⁺ $\gamma\delta$ T cells are important in innate protective immunity against *L. monocytogenes* infection.

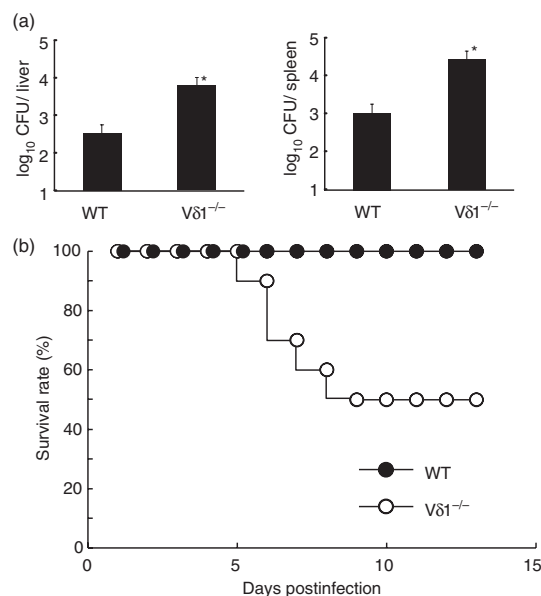


Figure 2. Exacerbated innate protective immunity against *Listeria monocytogenes* infection in the V δ 1^{-/-} mice. (a) The wild type (WT) or V δ 1^{-/-} mice were infected intraperitoneally (i.p.) with 5×10^4 colony-forming units (CFU) of *L. monocytogenes*, and bacterial counts in the liver and spleen were determined on day 5 after infection. Data that are representative of three independent experiments are shown. * $P < 0.05$ compared to WT mice. (b) The WT and V δ 1^{-/-} mice were inoculated i.p. with 2.5×10^5 CFU *L. monocytogenes*, and survival rates of the mice were monitored for 2 weeks. The survival rates of the WT and V δ 1^{-/-} mice were significantly different ($P = 0.003$). Two experiments showed nearly the same results, and representative data are demonstrated.

Formation of abscess-like lesions and increase of macrophage infiltration in the liver of the V δ 1^{-/-} mice after *L. monocytogenes* infection

To analyse the mechanism of the V γ 6/V δ 1⁺ $\gamma\delta$ T-cell-mediated protection, we compared the liver histology of the V δ 1^{-/-} mice and wild-type mice on day 5 of *L. monocytogenes* infection. The wild-type mice showed small granulomatous inflammatory lesions in sections with haematoxylin & eosin staining (arrow head in Fig. 3a upper panels), these consisted of colocalized CD11b⁺ cells and CD3⁺ cells in CF-LSM analysis (Fig. 3b, upper panels). In contrast, V δ 1^{-/-} mice showed a higher number of inflammatory lesions (Fig. 3a, arrowheads in lower panels) with large, abscess-like lesions (Fig. 3a, arrows in lower panels) in the infected liver. Furthermore, the V δ 1^{-/-} mice failed to show an organized granulomatous lesion with colocalization of CD11b⁺ and CD3⁺ cells in CF-LSM (Fig. 3b, lower panels).

Next, the infiltration of inflammatory cells in the liver of the V δ 1^{-/-} and wild-type mice was analysed 5 days after *L. monocytogenes* infection. Although the total liver mononuclear cell counts were not significantly different between

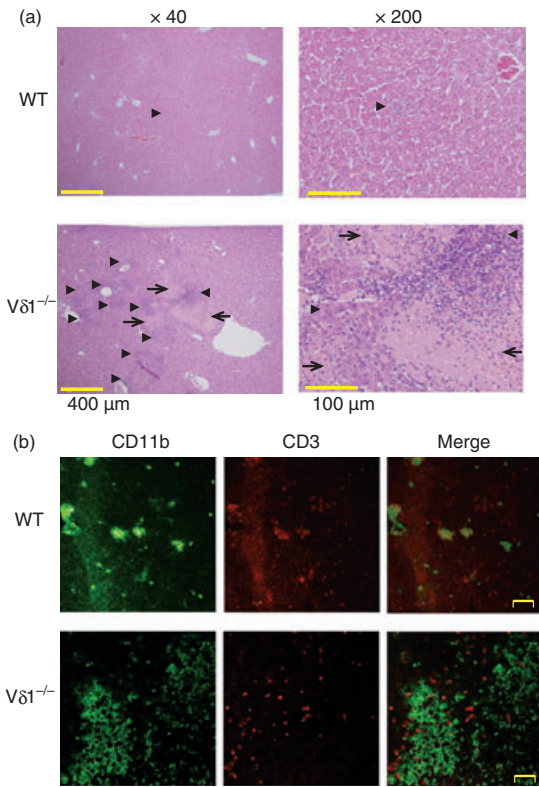


Figure 3. Histological examination of the liver of the wild-type (WT) and $V\delta 1^{-/-}$ mice on day 5 of *L. monocytogenes* infection. (a) Haematoxylin & eosin staining of the liver sections from the WT and $V\delta 1^{-/-}$ mice is demonstrated. There were only a few granulomatous lesions of inflammatory cell accumulation (arrowhead) in the WT mice (upper panels). In contrast, the liver sections from $V\delta 1^{-/-}$ mice (lower panels) showed higher numbers of inflammatory cell-accumulated lesions (arrowheads), and large abscess-like lesions (arrows). (b) Confocal laser scanning microscopy analysis of the expression of CD11b and CD3 is shown on the liver sections from the WT and $V\delta 1^{-/-}$ mice. In the WT mice, accumulated CD11b⁺ cells in the liver colocalized with CD3⁺ cells (upper panels) while colocalization of CD11b⁺ cells and CD3⁺ cells was hardly observed in the liver of the $V\delta 1^{-/-}$ mice (lower panels).

the wild-type and $V\delta 1^{-/-}$ mice, the number of macrophages was significantly higher in the $V\delta 1^{-/-}$ mice compared to the wild-type mice (Fig. 4). The number of $\gamma\delta$ T cells was slightly lower in the $V\delta 1^{-/-}$ mice, which may be because of the lack of $V\gamma 6/V\delta 1^{+}$ $\gamma\delta$ T cells. The number of natural killer T cells was also lower in the $V\delta 1^{-/-}$ mice. In contrast, the numbers of neutrophils and conventional CD4⁺ and CD8⁺ T cells in the livers of $V\delta 1^{-/-}$ mice were similar to those in the livers of wild-type mice.

IFN- γ and IL-17A production by the $V\gamma 6/V\delta 1^{+}$ $\gamma\delta$ T cells in the liver of *L. monocytogenes*-infected mice

It is well established that proinflammatory cytokines have a pivotal role in regulation of the innate immune response against *L. monocytogenes* infection. Both IFN- γ and IL-17A are reported as proinflammatory cytokines produced by $\gamma\delta$ T cells.²⁻⁵ Therefore, expression of the cytokines was analysed by FCM on the liver $\gamma\delta$ T cells of the wild-type and $V\delta 1^{-/-}$ mice after *L. monocytogenes* infection. When the $\gamma\delta$ T cells of the wild-type mice were stained with the anti-V γ mAb mixture and anti-cytokine mAb, expression of IL-17A and IFN- γ was detected not only in the anti-V γ mAb mixture-positive fraction but also in the anti-V γ mAb mixture-negative fraction represented by the $V\gamma 6/V\delta 1^{+}$ $\gamma\delta$ T cells (Fig. 5, upper panels). Interestingly, the anti-V γ mAb mixture-negative $\gamma\delta$ T cells contained more IFN- γ -producing $\gamma\delta$ T cells compared to the anti-V γ mAb mixture-positive $\gamma\delta$ T cells. Analysis of the liver $\gamma\delta$ T cells from *L. monocytogenes*-infected $V\delta 1^{-/-}$ mice showed that IFN- γ expression by the $\gamma\delta$ T cells decreased in the liver $\gamma\delta$ T cells as the result of a lack of IFN- γ -producing $V\gamma 6/V\delta 1^{+}$ $\gamma\delta$ T cells in the anti-V γ mixture-negative fraction (Fig. 5, lower panels). In contrast, expression of IL-17A was maintained in the liver $\gamma\delta$ T cells of the $V\delta 1^{-/-}$ mice because IL-17A production by the anti-V γ mAb mixture-positive $\gamma\delta$ T cells, especially $V\gamma 4^{+}$ $\gamma\delta$ T cells (data not shown),

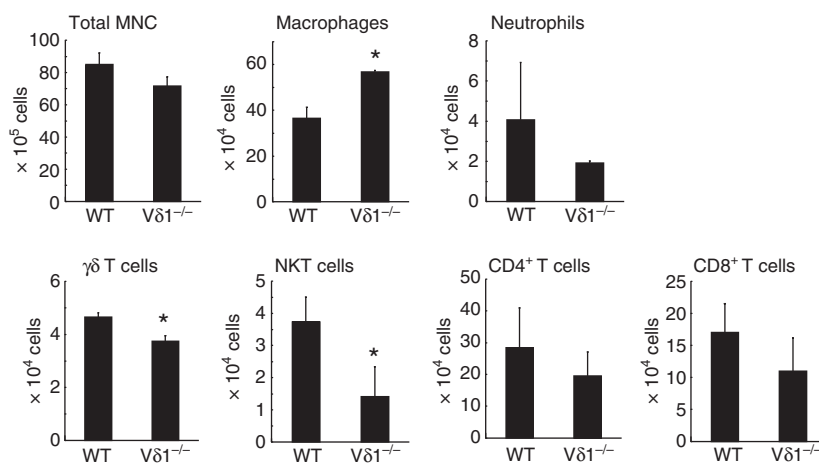


Figure 4. The number of inflammatory cells in the livers of wild-type (WT) and $V\delta 1^{-/-}$ mice on day 5 of *Listeria monocytogenes* infection. The ratios of macrophages, neutrophils and T-cell subpopulations were determined by flow cytometry and the absolute numbers of the cell populations were calculated as described in the *Materials and methods* section. * $P < 0.05$ compared to WT mice.

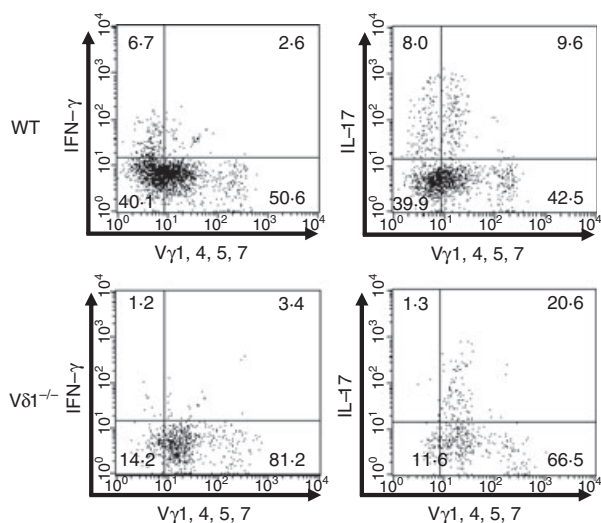


Figure 5. Expression of interleukin-17A (IL-17A) and interferon- γ (IFN- γ) by the $\gamma\delta$ T cells induced in the livers of the wild-type (WT) and V δ 1^{-/-} mice on day 5 of *Listeria monocytogenes* infection. The liver $\gamma\delta$ T cells were enriched, and stimulated *in vitro* with phorbol 12-myristate 13-acetate and calcium ionophore for 4 hr in the presence of brefeldin A. The cells were stained with anti-TCR C δ monoclonal antibody (mAb), anti-V γ mAb mixture, and anti-IL-17A or anti-IFN- γ mAb, and were analysed by flow cytometry. The analysis gate was set on the TCR C δ ⁺ T cells in all panels. Experiments were repeated three to five times, and representative data are shown.

increased in the mice. The result suggests that proinflammatory cytokines IFN- γ and IL-17A, especially IFN- γ , may participate in V γ 6/V δ 1⁺ $\gamma\delta$ T-cell-mediated regulation of innate immunity against *L. monocytogenes* infection.

Discussion

It has been reported that $\gamma\delta$ T cells have a pivotal role in innate protective immunity against *L. monocytogenes* infection in the liver.^{8–10} The $\gamma\delta$ T cells induced by *L. monocytogenes* infection have been divided based on their V γ and V δ repertoires into V γ 1⁺, V γ 4⁺, and V γ 6/V δ 1⁺ T cells.^{11–15} The role of V γ 6/V δ 1⁺ $\gamma\delta$ T cells was analysed using mice deficient for both V γ 4 and V γ 6 genes; however, function of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells could not be separated from that of V γ 4⁺ $\gamma\delta$ T cells in the mice.¹³ In the present report, the importance of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells in the protective immunity against *L. monocytogenes* was demonstrated using V δ 1^{-/-} mice, which selectively lack V γ 6/V δ 1⁺ $\gamma\delta$ T cells in the *L. monocytogenes*-infected liver.

Various stimuli induce V γ 6/V δ 1⁺ $\gamma\delta$ T cells in inflammatory sites. It has been reported that systemic infection of *L. monocytogenes* induces V γ 6/V δ 1⁺ $\gamma\delta$ T cells expressing invariant junctional sequences developed in the fetal thymus.^{14,15} The invariant V γ 6/V δ 1⁺ $\gamma\delta$ T cells are also induced in the peritoneal cavity of mice after i.p. inocula-

tion of *Escherichia coli*.^{19,24} Furthermore, in a murine autoimmune orchitis model,^{25,26} infiltration of the invariant V γ 6/V δ 1⁺ $\gamma\delta$ T cells was detected in the lesions produced by the autoimmune inflammatory response in the testis. All the results suggest that V γ 6/V δ 1⁺ $\gamma\delta$ T cells are induced in the inflammatory site independent of the recognition of exogenous bacterial antigens. Consistent with these observations, V γ 6/V δ 1 soluble TCR-octamer stained murine cells such as keratinocytes and fibroblasts in the absence of exogenous antigen, indicating that the V γ 6/V δ 1 TCR is specific for self-surface molecules or molecular complexes.^{2,27} Although ligand specificity of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells is still unclear, the observations suggest that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells participate in immune surveillance in inflammatory sites through recognition of self antigen.

The V γ 6/V δ 1⁺ $\gamma\delta$ T cells may participate in innate immunity against infections through production of proinflammatory cytokines. Our data demonstrated that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells produced proinflammatory cytokines IL-17A and IFN- γ . Interferon- γ is an important cytokine against *L. monocytogenes* infection at an innate immunity level.^{28,29} It is possible that formation of early granulomatous lesions in the *L. monocytogenes*-infected liver depends on the V γ 6/V δ 1⁺ $\gamma\delta$ T-cell-derived IFN- γ because the importance of the IFN- γ in granuloma formation is well established.^{29,30} Interleukin-17A produced by the $\gamma\delta$ T cells also has a pivotal role in innate protection against *L. monocytogenes* (Hamada *et al.* unpublished observation). Production of IL-17A by the V γ 6/V δ 1⁺ $\gamma\delta$ T cells was also reported in *E. coli* infection.³¹ Therefore, both IL-17A and IFN- γ produced by the V γ 6/V δ 1⁺ $\gamma\delta$ T cells may be important in innate protective immunity against various pathogens. However, we estimate that IFN- γ production by the V γ 6/V δ 1⁺ $\gamma\delta$ T cells would be more important in V γ 6/V δ 1⁺ $\gamma\delta$ T-cell-mediated protection because IL-17A production of the $\gamma\delta$ T cells was compensated by V δ 1-negative $\gamma\delta$ T cells but infection was exacerbated in the V δ 1^{-/-} mice. Further analysis is required to clarify the relative importance of IFN- γ compared to IL-17A in the V γ 6/V δ 1⁺ $\gamma\delta$ T-cell-mediated protection.

Although our data support a proinflammatory role of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells, it is possible that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells are also regulatory against inflammation. Histological analysis of the liver of *L. monocytogenes*-infected V δ 1^{-/-} mice showed abscess-like lesions which were indistinguishable from those in the mice that were deficient for all $\gamma\delta$ T cells,^{9,10} suggesting the possibility that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells participate in innate immunity not only through the induction of inflammatory cells, but also through the suppression of an excessive inflammatory response that damages infected tissues. In agreement with this, depletion of $\gamma\delta$ T cells accelerated the inflammatory response in the testis of mice with autoimmune orchitis with V γ 6/V δ 1⁺ $\gamma\delta$ T-cell induction in the auto-

immune lesions.³² The result suggests a regulatory mechanism for V γ 6/V δ 1⁺ $\gamma\delta$ T cells against inflammation. V γ 6⁺ $\gamma\delta$ T cells were also shown to suppress *Bacillus subtilis*-induced pulmonary fibrosis through suppression of CD4⁺ and CD8⁺ T-cell responses.³³ Therefore it is possible that the V γ 6/V δ 1⁺ $\gamma\delta$ T cells differentiate not only into proinflammatory (IFN- γ -producing or IL-17A-producing) T cells but also into anti-inflammatory T cells, and optimize the inflammatory response to eliminate pathogens and protect organs from tissue injury. Further research is required to clarify the anti-inflammatory function of the V γ 6/V δ 1⁺ $\gamma\delta$ T cells.

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