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$\gamma\delta$ T Cells Promote the Maturation of Dendritic Cells During West Nile Virus Infection

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Abstract

$\gamma\delta$ T cells are important for the early control of West Nile virus (WNV) dissemination. Here, we investigated the role of $\gamma\delta$ T cells in regulation of CD4⁺ T cell response following WNV challenge. Splenic dendritic cells (DCs) of WNV-infected $\gamma\delta$ T cell-deficient (TCR $\delta^{-/-}$) mice displayed lower levels of CD40, CD80, CD86 and major histocompatibility complex (MHC) class II expression and interleukin-12 (IL-12) production than those of wild-type mice. Naïve DCs co-cultured with WNV-infected $\gamma\delta$ T cells had enhanced levels of co-stimulatory molecules, MHC class II expression and IL-12 production. Further, co-culture of CD4⁺ T cells from OT II transgenic mice with DCs of WNV-infected TCR $\delta^{-/-}$ mice induced less interferon- γ (IFN- γ) and IL-2 production than with those of wild-type controls. Viral antigens were detected in WNV-infected $\gamma\delta$ T cells. WNV infection or toll-like receptor (TLR) agonist treatment of $\gamma\delta$ T cells induced the production of IFN- γ , tumor necrosis factor- α (TNF- α) and IL-6, which were known to promote DC maturation. Nevertheless, levels of TLRs 2, 3, 4 and 7 expression of WNV-infected $\gamma\delta$ T cells were not different from those of non-infected cells. Overall, these data suggest that WNV-induced $\gamma\delta$ T cell activation promotes DC maturation and initiates CD4⁺ T cell priming.

Keywords

West Nile virus; Dendritic cell; $\gamma\delta$ T cell

Introduction

West Nile virus (WNV), a mosquito-borne neurotropic flavivirus, has caused annual outbreaks of viral encephalitis in North America for nearly a decade (Campbell, *et al.*, 2002, Pletnev, *et al.*, 2006). Although most WNV infections in humans are asymptomatic, severe neurological disease (including encephalitis) and death have been observed with a higher frequency in the elderly and immunocompromised (Campbell, *et al.*, 2002, Pletnev, *et al.*, 2006). Human vaccines are not available yet.

The murine model has been used as an effective tool to investigate host immunity to WNV infection in humans. Type I interferon (IFN) and $\gamma\delta$ T cells provide immediate control of virus dissemination (Anderson & Rahal, 2002, Wang, *et al.*, 2003, Samuel & Diamond, 2005). B cells and specific antibodies are critical in the control of disseminated WNV infection, but are not sufficient to eliminate it from the host (Roehrig, *et al.*, 2001, Diamond, *et al.*, 2003). $\alpha\beta$ T cells (Diamond, *et al.*, 2003) provide long-lasting protective immunity and contribute to host survival following WNV infection. CD4⁺ T cells provide help for antibody responses and sustain WNV-specific CD8⁺ T cell responses in the central nervous system (CNS) (Sitati & Diamond, 2006), whereas CD8⁺ effector T cells are important in clearing WNV infection from tissues and preventing viral persistence (Wang, *et al.*, 2003, Shrestha & Diamond, 2004). The development of memory T cells following WNV infection remains poorly understood.

Dendritic cells (DCs) represent the most important antigen presenting cells exhibiting the unique capacity to initiate primary T cell responses. Upon microbial infection, DC maturation is an innate response that leads to adaptive immunity to foreign antigens (De Smedt, *et al.*, 1996, Bennett, *et al.*, 1998). Maturation of DCs results in the expression of high levels of major histocompatibility complex (MHC) and co-stimulatory molecules such as CD40, CD80 and CD86 and is often associated with the secretion of interleukin-12 (IL-12) (Inaba, *et al.*, 2000, Fujii, *et al.*, 2004). Proinflammatory cytokines such as tumor necrosis factor-alpha (TNF- α) and IFN- γ , promote this process (Dieli, *et al.*, 2004, Conti, *et al.*, 2005). Furthermore, increasing evidence suggests that the cross talk between $\gamma\delta$ T cells and DCs contributes to DC maturation (Leslie, *et al.*, 2002, Collins, *et al.*, 2005, Munz, *et al.*, 2005). Nevertheless, the *in vivo* mechanisms underlying this process are not clearly identified.

We have recently shown that $\gamma\delta$ T cells expanded quickly in response to WNV infection and produced significant amount of IFN- γ (Wang, *et al.*, 2003). $\gamma\delta$ T cell-deficient (TCR $\delta^{-/-}$) mice had a reduced CD8⁺ T cell memory response and were more susceptible to secondary WNV infection, suggesting a role of $\gamma\delta$ T cells in linkage of innate immunity to adaptive immune responses (Wang, *et al.*, 2006). In this study, we investigate the role of $\gamma\delta$ T cells in regulating DC maturation and initiating CD4⁺ T cell priming following WNV challenge.

Materials and Methods

Mice

6-10-week-old C57BL/6 (B6) mice and OT II transgenic mice were purchased from the Jackson Laboratory Bar Harbor, ME). TCR $\delta^{-/-}$ mice were a kind gift from Dr. E. (Fikrig (Yale University, New Haven). Groups were age and sex-matched for each experiment and were housed under identical conditions. All animal experiments were approved by the Institutional Animal Care and Use Committee at Colorado State University.

Infection in mice

WNV NY99-6480 was passaged three times in C6/36 *Aedes albopictus* cells to make a virus stock for both cell culture and *in vivo* studies. Mice were inoculated intraperitoneally with 100 PFU of WNV NY99-6480 isolate.

Purification of DCs, CD4⁺, and $\gamma\delta$ T cell subsets

DCs, CD4⁺ and $\gamma\delta$ T cells were purified from pooled spleens of 3–5 mice by a positive selection method, using anti-CD11c, anti-mPDCA-1, anti-CD4 magnetic beads or FITC-conjugated anti-mouse TCR $\gamma\delta$ BD Biosciences, San Diego, CA) followed by anti-FITC magnetic beads according to the manufacturer's instructions (Miltenyi Biotec, Auburn, CA).

The purity of these cells is 82–95%. For FACS purification of $\gamma\delta$ T cells, splenocytes were enriched for T cells using anti-CD90 magnetic beads (Miltenyi Biotec), stained with FITC labeled anti-TCR $\gamma\delta$ and sorted based on expression of TCR $\gamma\delta$ (MoFlo, DakoCytomation). The purity of $\gamma\delta$ T cells was 94.3%.

Flow cytometry

Antibodies for CD40, CD80, CD86, I-A^b and CD11c were purchased from BD Biosciences. Following staining, cells were fixed in PBS with 1% paraformaldehyde and examined using a Cyan flow cytometer (Dako Cytomation). Data were analyzed using Summit 4 software (Dako Cytomation). For intracellular cytokine staining, splenocytes were stimulated at 2×10^6 cells/well with 10 μ g/ml LPS (Sigma-Aldrich, St. Louis, MO) or 0.5 μ g/ml CL097 (Invivogen, San Diego, CA) with Golgi-Plug for 4 h at 37°C. Cells were harvested, stained with FITC-anti-CD11c, fixed in Fixation/Permeabilization solution before adding PE- anti-IL-12 (eBioscience, San Diego, CA) or rat IgG2a (BD Biosciences).

In vitro DC maturation and T cell priming assays

Naïve DCs were co-cultured with $\gamma\delta$ T cells from non-infected or WNV-infected mice at 1:1 ratio in 24-well plates at 37 °C for 24 h. Cells were harvested and stained with antibodies for cell surface markers. CD11c⁺ cells were gated for analysis. DCs were also co-cultured with *in vitro* WNV-infected $\gamma\delta$ T cells. CD4⁺ T Cells and DCs were purified from splenocytes of OT II transgenic mice or WNV-infected mice at day 3 post-infection. CD4⁺ T Cells (1×10^5) were mixed with DCs at 1:1 ratio and treated with or without OVA residue 323–339 (10 μ g/ml, Genscript Corporation, Piscataway, NJ).

WNV infection or stimulation with TLR agonist in $\gamma\delta$ T cells

$\gamma\delta$ T cells (1×10^5 cells/well) were cultured for 2 days at 37 °C in RPMI-1640 medium (Invitrogen, Carlsbad, CA) in 96-well plates coated with 5 μ g/ml anti-CD3 (eBioscience). Cells were infected with WNV at a MOI of 0.5 for 1 h, washed and incubated in the above medium containing 5 ng/ml recombinant human IL-2 (eBioscience). H36.12j cells (macrophage cell line, American Type Culture Collection, Manassas, VA) were infected with WNV (MOI = 1) and harvested at day 4 post-infection. In some experiments, $\gamma\delta$ T cells were stimulated with poly I: C (Amersham Pharmacia, New Jersey, 50 μ g/ml) or CL097 (Invivogen, 0.5 μ g/ml).

Quantitative PCR (Q-PCR) and PCR for determining viral load and TLR levels

RNA was extracted using RNAeasy extraction kit (Qiagen, Valencia, CA) and was used to synthesize complementary (c)DNA using the ProSTAR First-strand RT-PCR kit (Stratagene, Cedar Creek, TX). The sequences of the primersets for WNV envelope gene (*WNE*), *Tlr2*, *Tlr3*, *Tlr4* and *Tlr7* cDNA and PCR conditions were described previously (Lanciotti, *et al.*, 2000, Schulz, *et al.*, 2005, Chen, *et al.*, 2006). Q-PCR analysis was performed with RT2 Real-Time SYBR Green / Fluorescein PCR master mix (Superarray, Frederick, MD) on an iCycler (Bio- Rad, Hercules, CA). The ratio of the amount of amplified gene compared with the amount of β -actin cDNA represented the relative levels in each sample. Regular PCR was performed as follows: 94°C for 2 min followed by 35 cycles of 94°C for 1 min, 60°C for 1 min and 72°C for 1.5 min, and final extension at 72°C for 5 min. The primer pairs used were: TLR2, 5'-CAG ACG TAG TGA GCG AGC TG-3' and 5'-GGC ATC GGA TGA AAA GTG TT-3'; TLR3, 5'-CCC CCT TTG AAC TCC TCT TC-3' and 5'-TTT CGG CTT CTT TTG ATG CT -3'; TLR4, 5'-GCT TTC ACC TCT GCC TTC AC-3' and 5'-CGA GGC TTT TCC ATC CAA TA-3'; TLR7, 5'-CAT CAG AGG CTC CTG GAT GA-3' and 5'-AGG CAT GTC CTA GGT GGT GA-3'. The sequences of the primersets for β -actin were described earlier (Farrar & Street, 1995).

Fluorescence microscopy with $\gamma\delta$ T-cells

Cells were fixed in acetone at -20°C for 30 min, rehydrated in PBS and stained with FITC-conjugated anti-CD3, Phycoerythrin (PE)-conjugated anti-TCR β (eBioscience) and the flavivirus E protein-specific monoclonal antibody 4G2 followed by Alexa Fluor 350-conjugated anti-mouse IgG2a (Invitrogen) at 37°C . Images were acquired on an Olympus IX71 Inverted Microscope at $20\times$ magnification.

Plaque assay

Vero cells were seeded in DMEM (Invitrogen) with 10% FBS and incubated with serial dilutions of culture supernatant for 2 h. DMEM containing 5% FBS and 1% low-melting-point agarose was added and incubated for 4 days. A second overlay of 1% agarose-medium containing 0.01% neutral red was added to visualize plaques.

Cytometric bead array and ELISA

Culture supernatant was measured for cytokine production using a mouse Th1/Th2 kit or an inflammation kit by a FACSArray analyzer (BD Biosciences). Cytokine levels were also measured by ELISAs (BD Biosciences & PBL Interferon Source).

Statistical analysis

Data analysis was performed using Prism software (Graph-Pad) statistical analysis. Values for phenotype analysis, viral burden, and cytokine production experiments were presented as means \pm SEM. *P* values of these experiments were calculated with a non-paired Student's *t* test or Mann-Whitney test. Statistical significance was accepted at $P < 0.05$.

Results

DC activation and maturation was reduced in TCR $\delta^{-/-}$ mice during WNV infection

Our previous results have shown that TCR $\delta^{-/-}$ mice were much more susceptible to a LD₅₀ dose of WNV infection than wild-type controls (Wang, *et al.*, 2003). Further, TCR $\delta^{-/-}$ mice that survived a LD₅₀ dose of WNV challenge were more susceptible to the secondary infection than wild-type mice (Wang, *et al.*, 2006). To investigate the role of $\gamma\delta$ T cells in regulating CD4⁺ T cell response, we assessed splenic DCs phenotype and functionality in wild-type and TCR $\delta^{-/-}$ mice following infection with a LD₅₀ dose of WNV. At day 3 post-infection, the expression of CD40, CD80, CD86, and MHC class II on CD11c⁺ splenocytes of wild-type mice was increased by percentage and mean fluorescence intensity (MFI) ($P < 0.05$ or 0.01 , Fig. 1A, Table 1 and Suppl. Fig. 1). In WNV-infected TCR $\delta^{-/-}$ mice, percentage of CD80⁺CD11c⁺, CD86⁺CD11c⁺ splenocytes or MFI on these cells were also increased ($P < 0.05$ or 0.01); while CD40 and MHC class II expression were only elevated by percentage ($P < 0.01$, Fig. 1A, Table 1 & Suppl. Fig. 1). Interestingly, in comparison to wild-type mice, expression of all these surface molecules was significantly lower in CD11c⁺ splenocytes of WNV-infected TCR $\delta^{-/-}$ mice by percentage (12–28%, Fig. 1A) or by MFI (16–32% except for CD80, Table 1 & Suppl. Fig. 1, $P < 0.05$ or 0.01). Similar results were observed at day 5 post-infection, though the magnitude of increase of these surface molecules expression was reduced in both groups of mice by 5–40% as compared to day 3 (data not shown). There were no differences in the expression of the above surface molecules in CD11c⁺ cells between naïve wild-type and TCR $\delta^{-/-}$ mice (Fig. 1A & Table 1 and Suppl. Fig. 1).

Proinflammatory cytokines, including TNF- α , IFN- γ and IL-12, are important for DC maturation (Sallusto & Lanzavecchia, 1994, Inaba, *et al.*, 2000, Le Bon, *et al.*, 2001). We next measured cytokine production in CD11c⁺ DCs from wild-type and TCR $\delta^{-/-}$ mice using

ex vivo intracellular cytokine staining. There were no differences in the percentage of IL-12-producing DCs between naive wild-type and TCR $\delta^{-/-}$ mice upon stimulation with the TLR4 agonist LPS or the TLR7 agonist CL097 (Fig. 1B left panel). However, the percentage of IL-12-producing DCs in WNV-infected TCR $\delta^{-/-}$ mice stimulated with LPS or CL097 were 54% or 26% lower than those of WNV-infected wild-type mice ($P < 0.01$, Fig. 1B right panel). There were no significant differences in TNF- α and IFN- γ production in CD11c $^{+}$ DCs between these two groups (data not shown). Overall, these data suggest that $\gamma\delta$ T cells are involved in the process of DC maturation during WNV infection.

DCs exposed to WNV-infected $\gamma\delta$ T cells acquire the functional and phenotypic characteristics of mature cells

To verify that $\gamma\delta$ T cells are involved in DC activation during WNV infection, we performed *ex vivo* DC maturation assays. In this assay, CD11c $^{+}$ DCs were purified from naïve B6 mice and co-cultured with $\gamma\delta$ T cells from non-infected or day 2 post-infected mice. After 24 h co-culture, cells were harvested and gated on CD11c $^{+}$ population for phenotypic analysis. Interestingly, CD40, CD80, CD86 and MHC class II expression on CD11c $^{+}$ cells were enhanced after co-culture with naive $\gamma\delta$ T cells by MFI and/or percentage (Figs. 2A & 2B, $P < 0.05$). Further, DCs co-cultured with $\gamma\delta$ T cells from WNV-infected mice had a higher level of expression of these cell surface molecules than those co-cultured with naïve $\gamma\delta$ T cells or DC alone (Figs. 2A & 2B, $P < 0.05$). We also co-cultured immature DCs with *in vitro* WNV-treated $\gamma\delta$ T cells. At 24 h, IL-12 levels in co-culture of DCs with WNV-treated and anti-CD3 stimulated $\gamma\delta$ T cells were about 150% higher than those of DCs alone (Fig. 2C, $P < 0.05$). DCs co-cultured with non-infected and anti-CD3-stimulated $\gamma\delta$ T cells also induced higher levels of IL-12 (about 60%) than DCs alone (Fig. 2C, $P < 0.05$). $\gamma\delta$ T cells, naive or activated by anti-CD3-stimulation and/or WNV infection, did not produce IL-12 (data not shown).

DCs from WNV-infected TCR $\delta^{-/-}$ mice could not prime CD4 $^{+}$ T cells as efficiently as those of wild-type mice

To further understand the role of $\gamma\delta$ T cells in regulating CD4 $^{+}$ T cell response during WNV infection, we tested the capability of DCs from WNV-infected mice to activate naïve CD4 $^{+}$ T cells *in vitro*. Purified naïve CD4 $^{+}$ T cells from OT II transgenic mice were co-cultured with DCs from WNV-infected wild-type or TCR $\delta^{-/-}$ mice in the presence of OVA 323–339. At 24 h post co-culture, OT II CD4 $^{+}$ T cells co-cultured with DCs from wild-type mice produced about 46% more IFN- γ ($P < 0.01$) or 15% IL-2 ($P < 0.05$) respectively than those cocultured with DCs of TCR $\delta^{-/-}$ mice (Table 2). At 72 h, IFN- γ but not IL-2 production remained higher in co-culture with wild-type DCs than TCR $\delta^{-/-}$ DCs (data not shown). These data suggests that the antigen-presenting capacity of DCs might be reduced or impaired in the absence of $\gamma\delta$ T cells.

WNV antigens were detected in the infected $\gamma\delta$ T cells

We have recently demonstrated that splenic T cells are permissive to WNV infection and support a short-term virus replication (Wang, *et al.*, 2008). Here, we asked whether $\gamma\delta$ T cells could be infected by WNV. Splenic $\gamma\delta$ T cells were purified and stimulated *in vitro* with anti-CD3 for 2 days before WNV infection. The purity of $\gamma\delta$ T cells was close to 94% as analyzed by flow cytometry (Fig. 3A). Immunofluorescence staining demonstrated CD3 $^{+}$ /TCR $\gamma\delta^{+}$ /WNV $^{+}$ populations in these cells at day 2 post-infection (Fig. 3B). Plaque assay showed WNV replicated productively in purified $\gamma\delta$ T cells at days 1 and 2 post-infection, but decreased at day 3 ($P < 0.05$, Fig. 3C). Further, Q-PCR analysis of day 4 post-infected $\gamma\delta$ T cells revealed a low but significant level of virus infection as compared to WNV-infected H36.12j cells (Fig. 3D).

WNV infection or TLR agonist stimulation of $\gamma\delta$ T cells induces proinflammatory cytokine production. However, the expression of TLRs on $\gamma\delta$ T cells was not changed after infection

The production of proinflammatory cytokines, including IFN- γ , TNF- α and IL-6 from $\gamma\delta$ T cells was increased as early as day 2 post-infection (data not shown) and became more dramatically enhanced at day 4 post-infection (Fig. 4A–C, $P < 0.05$ or 0.01). The TLR family plays a fundamental role in host innate immunity by mounting a rapid and potent inflammatory response to pathogen infection via recognition of conserved structural patterns in diverse microbial molecules. The expression of TLR2, TLR3, TLR4 and TLR7/8 on $\gamma\delta$ T cells has been reported (Shimura, *et al.*, 2005, Beetz, *et al.*, 2007, Peng, *et al.*, 2007, Beetz, *et al.*, 2008). Among them, TLR3 and TLR7-induced Type I IFNs and proinflammatory cytokine production are known to play important roles in host immunity following WNV infection (Wang, *et al.*, 2004, Daffis, *et al.*, 2008, Town, *et al.*, 2009). Here, we found $\gamma\delta$ T cells stimulated with TLR agonists such as CL097 (TLR7) or poly I: C (TLR3) also produced significant amount of IFN- γ , TNF- α , and/or IL-6 (Figs. 4D–F, $P < 0.01$). Nevertheless, there were no significant differences in the levels of TLR2, TLR3, TLR4 and TLR7 expression between WNV-infected $\gamma\delta$ T cells and non-infected controls at 6 h (Figs. 5B–5E) or 24 h (Figs. 5A–5E) post-infection.

Discussion

Although several important immune factors have been recognized to be critical for immediate control of WNV dissemination, the development of long-lasting protective immunity against WNV is not well understood. In the present study, we investigated the role of $\gamma\delta$ T cells in regulating DC maturation and CD4⁺ T cell priming following WNV challenge. We found that DC activation and maturation was reduced in TCR $\delta^{-/-}$ mice during WNV infection. Immature DCs co-cultured with $\gamma\delta$ T cells of WNV-infected mice or *in vitro* infection had enhanced levels of co-stimulatory molecule expression and IL-12 production. Co-culture of CD4⁺ T cells of OT II mice with DCs of WNV-infected wild-type mice induced more IFN- γ and IL-2 production than with DCs of TCR $\delta^{-/-}$ mice. Moreover, WNV infection of $\gamma\delta$ T cells induces proinflammatory cytokine production without changes on TLR expression levels. Collectively, our data suggest that $\gamma\delta$ T cells are involved in DC maturation and CD4⁺ T cell priming following WNV challenge.

Increasing evidence suggests that both the crosstalk between $\gamma\delta$ T cells and DCs and proinflammatory cytokines contribute to DC maturation (Ismaili, *et al.*, 2002, Leslie, *et al.*, 2002, Collins, *et al.*, 2005, Munz, *et al.*, 2005, Conti, *et al.*, 2005). Here, we have observed that naïve DCs co-cultured with non-infected $\gamma\delta$ T cells have enhanced levels of co-stimulatory molecules and MHC class II expression. This suggests that crosstalk between $\gamma\delta$ T cell and DC is necessary for DC maturation. WNV-infected $\gamma\delta$ T cells produce proinflammatory cytokines, including IFN- γ , TNF- α and IL-6. Upregulation of co-stimulatory molecules and MHC class II expression was significantly higher on DCs that were co-cultured with WNV-infected $\gamma\delta$ T cells than with naïve $\gamma\delta$ T cells. These data further demonstrates that the secreting factors from WNV-infected $\gamma\delta$ T cells are also important for promoting DC maturation.

Current understanding of the biological role of $\gamma\delta$ T cell receptors during pathogen infection remains elusive. Unlike $\alpha\beta$ T cells, there are few antigens that are recognized by $\gamma\delta$ T cell receptor (Born & O'Brien, 2009). Although $\gamma\delta$ T cells support a short-term WNV replication and are activated after infection, it is not clear whether any viral antigen is recognized by T cell receptor. TLR3 and TLR7-induced Type I IFNs and proinflammatory cytokine production play important roles in host immunity, following WNV infection (Wang, *et al.*, 2004, Daffis, *et al.*, 2008, Town, *et al.*, 2009). It is likely that $\gamma\delta$ T cells are induced by WNV infection via the innate immune receptors, such as TLRs (Peng, *et al.*, 2007, Beetz, *et*

al., 2008). This is further supported by the fact that both WNV infection and TLR agonist stimulation induce $\gamma\delta$ T cells to produce IFN- γ , TNF- α and IL-6 cytokines. Expression of TLRs on $\gamma\delta$ T cells has been shown to be upregulated after microbial infection or burn injury followed by activation (Mokuno, *et al.*, 2000, Shibata, *et al.*, 2007, Schwacha & Daniel, 2008). Nevertheless, we noted no differences in the levels of expression of TLRs after WNV infection. While the direct role of TLRs in $\gamma\delta$ T cell activation upon WNV infection is still under investigation, it is also possible that WNV infection of $\gamma\delta$ T cells induces the non-TLR innate immune receptors, such as RIG-I and MDA5, which have been reported to be involved in WNV recognition (Fredericksen & Gale, 2006, Fredericksen, *et al.*, 2008). Alternatively, $\gamma\delta$ T cells and DCs are also known to exert regulatory influences on each other. For example, induction of human $\gamma\delta$ T cells by poly I:C, a ligand for TLR3, depends on DCs mediated by Type-I IFNs (Kunzmann, *et al.*, 2004). Thus, TLR signaling might be involved in $\gamma\delta$ T cell activation indirectly through induction of WNV permissive DCs in a three-way process.

The role of T cells in protecting the host against WNV infection has been the subject of recent investigations. CD4⁺ T cells are known to provide help for antibody responses and to sustain WNV-specific CD8⁺ T cell responses in the CNS, enabling viral clearance (Sitati & Diamond, 2006). CD8⁺ T cells have important functions in clearing infection from peripheral tissues and CNS, and in preventing viral persistence (Shrestha & Diamond, 2004, Brien, *et al.*, 2007). Therefore, enhancement of memory T cell response is an important strategy for future flavivirus vaccine development. Although human and mouse $\gamma\delta$ T cells differ in the subsets and ligand recognition, they share substantial similarity in effector function and the protective role in pathogen infection (Girardi, 2006). The exploration of parallel activities mediated by murine $\gamma\delta$ T cells will continue to provide insights into immunosurveillance and immune regulation in human diseases. $\gamma\delta$ T cells are also known to display numerical and functional alteration in the elderly (Cardillo, *et al.*, 1993, Argentati, *et al.*, 2002), which are the potentially susceptible population for WNV encephalitis. Information gained from this study will possibly enhance our understanding of host immunity against WNV, and provide critical insights for new strategies for future flavivirus vaccine development.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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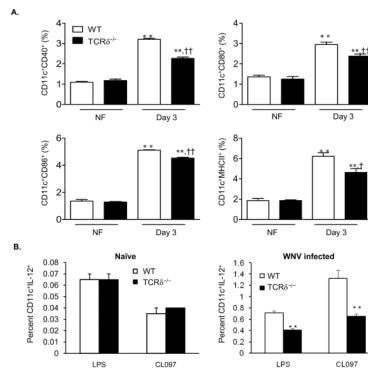
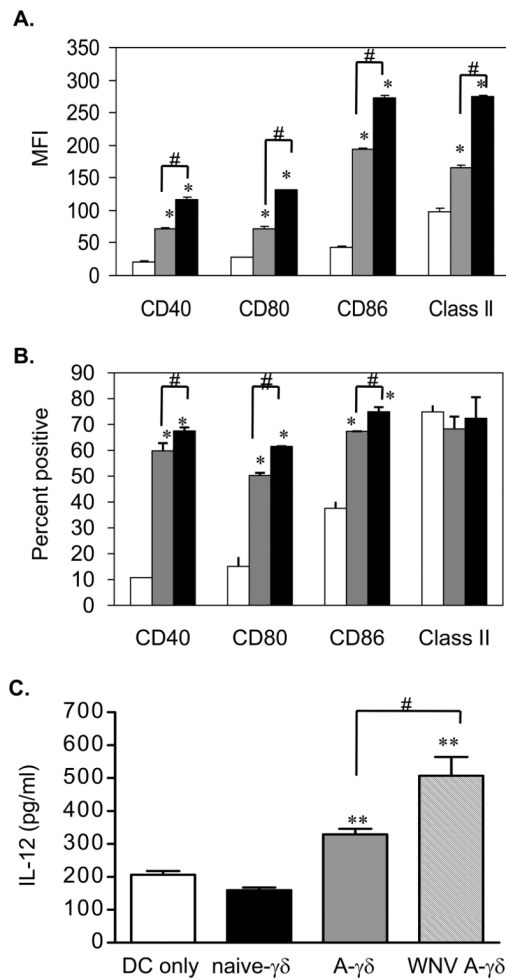


Figure 1. DC maturation in WNV-infected mice. *A*, The percentages of CD40⁺ CD11c⁺, CD80⁺CD11c⁺, CD86⁺CD11c⁺ or MHCII⁺CD11c⁺ in the spleens. Data are presented as means \pm SEM, $n = 4$ or 5 . ** $P < 0.01$ for WNV-infected (IF) vs. non-infected (NF). † $P < 0.05$ or †† $P < 0.01$ for wild-type (WT) vs. TCRδ^{-/-} mice. *B*, The percentages of IL-12⁺ CD11c⁺ naïve (left panel) or day 2 WNV-infected (right panel) splenocytes after *ex vivo* stimulation with LPS or CL097. Data are presented as means \pm SEM, $n = 4$ or 5 . ** $P < 0.01$ for wild-type vs. TCRδ^{-/-} mice. P values were calculated with a non-paired Student's *t* test.

**Figure 2.**

In vitro DC maturation assay. Cell surface molecule expression of CD11c⁺ cells alone (white bar) or co-culture with $\gamma\delta$ T cells of naïve (grey bar) or WNV-infected (black bar) mice as determined by MFI (A) and the percentage (B). C, IL-12 production from co-culture of naïve DCs with *in vitro* WNV infected $\gamma\delta$ T cells measured by ELISA. A- $\gamma\delta$: anti-CD3 activated $\gamma\delta$ T cells. WNV A- $\gamma\delta$: anti-CD3 treated $\gamma\delta$ T cells infected with WNV. Data are presented as means \pm SEM, $n = 6$. * $P < 0.05$ or ** $P < 0.01$ for co-cultured vs. DC alone. # $P < 0.05$ for IF vs. NF. P values were calculated with a non-paired Student's *t* test. Results are representative of two independent experiments.

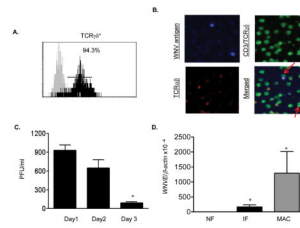


Figure 3.

In vitro WNV infection of $\gamma\delta$ T cells. **A**, Splenic $\gamma\delta$ T cells were stained with antibody to $\gamma\delta$ and analyzed by flow cytometer. Dark area represents cells stained with anti- $\gamma\delta$; gray area, unstained cells. **B**, Immunofluorescence photomicrographs of WNV-infected $\gamma\delta$ T cells at day 2 post-infection. Infected cells were stained for WNV antigen (blue), CD3/TCR $\gamma\delta$ (green) and TCR $\alpha\beta$ (red). Arrows point to CD3⁺/ $\gamma\delta$ ⁺WNV⁺ cells. **C**, Plaque assay of virus titer. * $P < 0.05$ for day 3 vs. day 1. **D**, WNV infection in splenic $\gamma\delta$ T cells as measured by Q-PCR. Negative and positive controls represent uninfected $\gamma\delta$ T cells and WNV-infected H36.12j cells (MAC), respectively. * $P < 0.05$ for non-infected vs. WNV-infected cells. P values were calculated with a non-paired Student's *t* test.

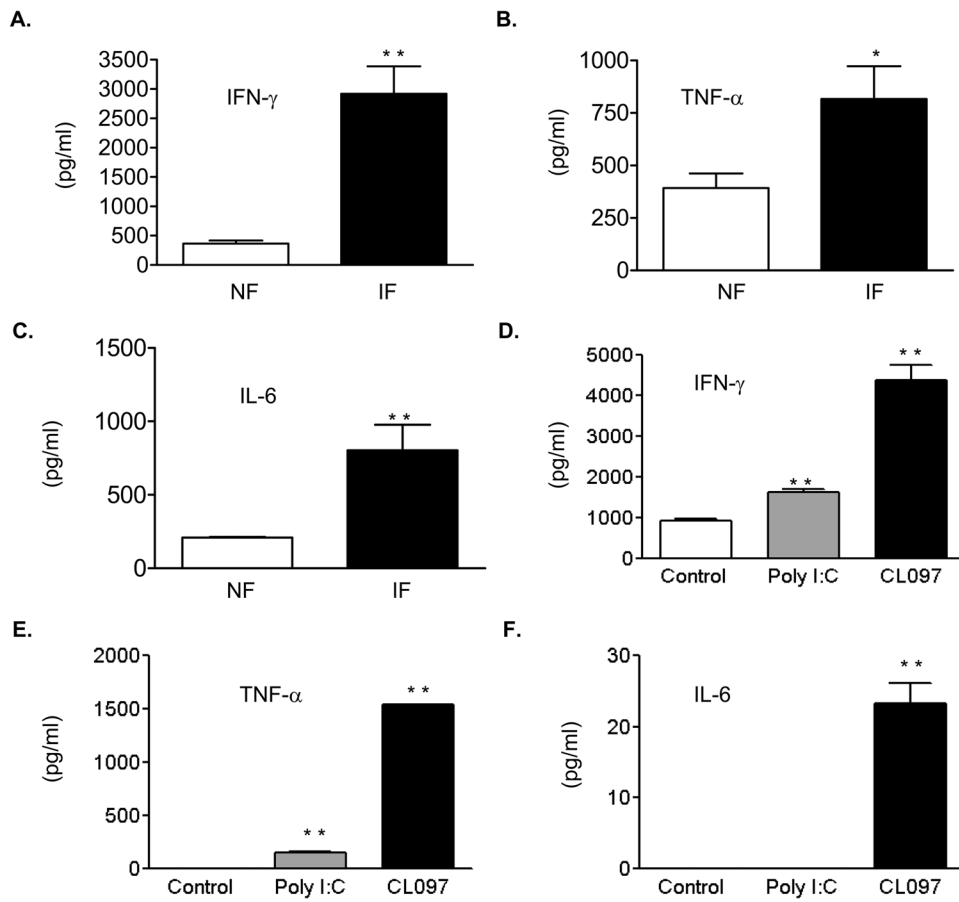


Figure 4. WNV infection or TLR agonist stimulation of $\gamma\delta$ T cells induces proinflammatory cytokines. *A–C*, Supernatant was collected at day 4 post-infection. *D–F*, Supernatant was collected 24 h post-treatment. Data are presented as means \pm SEM, $n = 6$. * $P < 0.05$ or ** $P < 0.01$ for non-infected or non-treated vs. WNV-infected or treated. P values were calculated with a non-paired Student's *t* test.

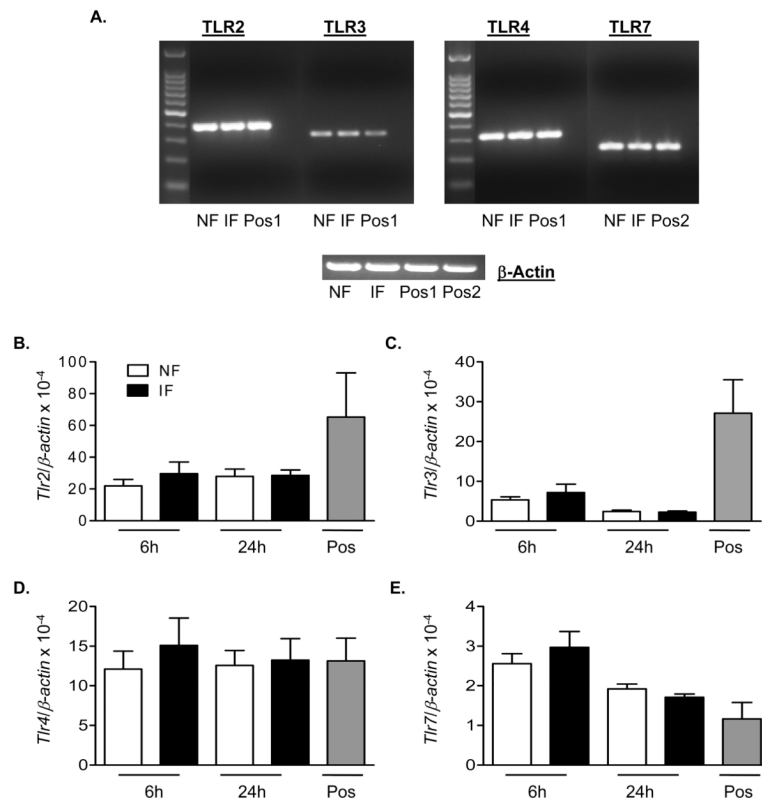


Figure 5.

Levels of TLR expression on WNV-infected $\gamma\delta$ T cells. Splenic $\gamma\delta$ T cells were infected with WNV and harvested at indicated time points. A, PCR amplification of TLRs (top panel) or β -actin (bottom panel) of WNV-infected or non-infected $\gamma\delta$ T cells harvested at 24 h post-infection. B–E, Q-PCR analysis of TLRs. B, TLR2. C, TLR3. D, TLR4. E, TLR7. cDNA of CD11c⁺ cells or plasmacytoid DCs were used as positive control 1 & 2. Data are presented as means \pm SEM, $n = 7$ or 8. P values were calculated with a Mann-Whitney test.

Table 1

Mean fluorescence intensity of CD40, CD80, CD86 and MHC class II expression on DCs from wild-type and *TCR $\delta^{-/-}$* mice following WNV infection:

Cell surface marker	Wild-type NF	Wild-type IF	<i>TCR$\delta^{-/-}$</i> NF	<i>TCR$\delta^{-/-}$</i> IF
CD40	40.2 \pm 3.1	65.2 \pm 3.2*	42.1 \pm 1.6	44.6 \pm 4.0 ^{††}
CD80	43.0 \pm 2.2	56.0 \pm 3.6*	38.0 \pm 2.5	52.9 \pm 1.8*
CD86	41.3 \pm 5.9	85.5 \pm 1.1**	39.7 \pm 5.4	70.7 \pm 0.6**, ^{††}
MHC class II	332.0 \pm 53	517.0 \pm 12.2**	338.7 \pm 55	434.0 \pm 12.3 ^{††}

Values are means \pm SEM, n = 4.

* $P < 0.05$ or

** $P < 0.01$ for WNV-infected (IF) vs. non-infected (NF).

^{††} $P < 0.01$ for wild-type vs. *TCR $\delta^{-/-}$* mice. Data shown were representative of two independent experiments.

P values were calculated with a non-paired Student's t test.

Table 2*In vitro* T cell priming assay:

Group	IFN-γ(pg/ml)	IL-2 (pg/ml)
Wild-type DCs + T cells	48.0 \pm 4.8	33.7 \pm 5.7
TCR $\delta^{-/-}$ DCs + T cells	43.3 \pm 3.2	33.3 \pm 5.8
CD4 ⁺ T cells + OVA	33.3 \pm 8.4	33.3 \pm 2.6
Wild-type DCs + OVA + CD4 ⁺ T cells	31,242 \pm 1623	18,953 \pm 759
TCR $\delta^{-/-}$ DCs + OVA + CD4 ⁺ T cells	21,640 \pm 835 ^{**}	16,543 \pm 392 [*]

Purified naïve CD4⁺ T cells from OT II transgenic mice were co-cultured with DCs from WNV-infected wild-type or TCR $\delta^{-/-}$ mice in the presence or absence of OVA 323–339. At 24 h post coculture, supernatant was harvested and measured for cytokine production using mouse Th1/Th2 cytokine kit. Data are presented as means \pm SEM, $n = 3$.

* $P < 0.05$ or

** $P < 0.01$ for wild-type vs. TCR $\delta^{-/-}$.

P values were calculated with a non-paired Student's t test. Data shown were representative of two independent experiments.