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Differential requirement of RasGRP1 for $\gamma\delta$ T cell development and activation

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Abstract

$\gamma\delta$ T cells ($\gamma\delta$ T) belong to a distinct T cell lineage that performs immune functions different from $\alpha\beta$ T cells ($\alpha\beta$ T). Previous studies have established that Erk1/2 MAPKs are critical for positive selection of $\alpha\beta$ T cells. Additional evidence also suggests that elevated Erk1/2 activity promotes $\gamma\delta$ T cell generation. RasGRP1, a guanine nucleotide releasing factor for Ras, plays an important role in positive selection of $\alpha\beta$ T cells by activating the Ras-Erk1/2 pathway. In this report, we demonstrate that RasGRP1 is critical for TCR-induced Erk1/2 activation in $\gamma\delta$ T cells but exerts different roles for $\gamma\delta$ T cell generation and activation. Deficiency of RasGRP1 does not obviously affect $\gamma\delta$ T cell numbers in the thymus but leads to increased $\gamma\delta$ T cells, particularly CD4⁻CD8⁺ $\gamma\delta$ T cells, in the peripheral lymphoid organs. The virtually unhindered $\gamma\delta$ T cell development in the *RasGRP1*^{-/-} thymus proved to be cell intrinsic, while the increase in CD8⁺ $\gamma\delta$ T cells is caused by non-cell-intrinsic mechanisms. Our data provides genetic evidence that decreased Erk1/2 activation in the absence of RasGRP1 is compatible for $\gamma\delta$ T cell generation. Although RasGRP1 is dispensable for $\gamma\delta$ T cell generation, RasGRP1-deficient $\gamma\delta$ T cells are defective in proliferation following TCR stimulation. Additionally, RasGRP1-deficient $\gamma\delta$ T cells are impaired to produce IL-17 but not IFN γ . Together, these observations have revealed that RasGRP1 plays differential roles for $\gamma\delta$ and $\alpha\beta$ T cell development but is critical for $\gamma\delta$ T cell proliferation and production of IL-17.

Introduction

Two lineages of T cells marked by the expression of two distinct antigen receptors, $\alpha\beta$ and $\gamma\delta$ T cell receptors (TCRs), are generated during intrathymic development. T cell

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development in the thymus can be divided into CD4⁻CD8⁻ (double negative, DN), CD4⁺CD8⁺ (double positive, DP), and CD4⁺CD8⁻ or CD4⁻CD8⁺ (single positive, SP) stages. DN thymocytes contain the most immature T cells and can be further divided from DN1 to DN4 based on CD25 and CD44 or cKit expression (1). Functional TCRs must be generated through somatic V(D)J recombination in the TCR loci for generation of either $\alpha\beta$ T cells ($\alpha\beta$ T) or $\gamma\delta$ T cells ($\gamma\delta$ T) (2). V(D)J recombination in the TCR loci is tightly regulated in a developmental-stage-specific manner. At the DN 2 and 3 stages, TCR γ , δ , and β loci rearrange. Formation of functional $\gamma\delta$ TCR directs progenitor cells to the $\gamma\delta$ lineage (3). TCR β associates with the pre-TCR α chain to form the pre-TCR, which drives DN thymocyte maturation to the DP stage and full commitment to the $\alpha\beta$ T cell lineage (4). DN2 thymocytes are mostly committed to the T cell lineage. $\gamma\delta$ T lineage commitment mainly occurs at the DN2 stage but can also happen at the DN3 stage (5). At the DP stage, the TCR α gene rearranges and in-frame rearranged α gene produces a functional chain to associate with the TCR β chain to drive DP thymocytes to mature to the SP stage (6). In normal thymus and peripheral lymphoid organs, $\gamma\delta$ T is the minor lineage while $\alpha\beta$ T is the dominant lineage. Most $\gamma\delta$ T cells reside in the DN population, and $\gamma\delta$ T cells expressing CD4 or CD8 co-receptor are rare.

It has been well documented that expression of a functional $\gamma\delta$ TCR or $\alpha\beta$ TCR in developing thymocytes is essential for the generation of their respective T cell lineages. Defects in formation of a functional $\gamma\delta$ TCR or $\alpha\beta$ TCR can cause a complete absence of $\gamma\delta$ or $\alpha\beta$ T cell lineage, respectively (3, 7, 8). Our knowledge of TCR signal transduction has primarily come from studies on the $\alpha\beta$ TCR, as $\gamma\delta$ T cells are rare. It is well known that TCR stimulation leads to the activation of PLC γ 1 via orchestrated actions of proximal protein kinases such as Lck, Zap70, and Itk, and adaptor molecules such as SLP-76 and LAT (9-12). Activated PLC γ 1 produces two critical second messengers, diacylglycerol (DAG) and inositol-1,4,5-trisphosphate (IP₃), that are crucial for relaying proximal signaling to the activation of distal signaling cascades (13). IP₃ binds to its receptor in ER, leading to depletion of calcium from ER and subsequent calcium influx through the CRAC channel, which leads to the activation of the calcineurin-NFAT pathway (14). DAG associates with multiple effector molecules, including the RasGRP1, PKC θ , and PKDs, to induce the activation of downstream signaling cascades such as the RasGRP1-Ras-Erk1/2 and PKC θ -CARM1/Bcl10-IKK-NF κ B pathways (13, 15, 16).

Evidence suggests that $\alpha\beta$ TCR and $\gamma\delta$ TCR signaling share at least some common features. Deficiency of some proximal signaling molecules such as Lck, SLP76, and LAT impacts both $\alpha\beta$ T cell and $\gamma\delta$ T cell generation (17). Additionally, expression of an $\alpha\beta$ TCR at early DN stages in transgenic mice can drive thymocytes to adopt the $\gamma\delta$ fate even though they express the $\alpha\beta$ TCR, suggesting that the timing of $\gamma\delta$ TCR versus $\alpha\beta$ TCR signaling rather than the quality of the signaling plays an important role in $\gamma\delta$ versus $\alpha\beta$ lineage fate decision (18). While similarities between $\alpha\beta$ and $\gamma\delta$ TCRs exist, differences between them have also been reported. Murine $\alpha\beta$ TCR but not $\gamma\delta$ TCR contains the CD3 δ chain (19, 20). The threshold of $\gamma\delta$ TCR signaling appears lower than $\alpha\beta$ TCR. Under the same stimulating condition, $\gamma\delta$ TCR induces stronger Erk1/2 activation than $\alpha\beta$ TCR. Furthermore, strong TCR signaling and Erk1/2 activation has been reported to promote $\gamma\delta$ differentiation (19, 21-24). However, substantial genetic evidence demonstrating the requirement of Erk1/2 for $\gamma\delta$ T cell lineage development is lacking.

RasGRP1, a guanine nucleotide exchange factor for Ras, is critical for the activation of the Ras-Erk1/2 pathway in $\alpha\beta$ T cells (25, 26). RasGRP1 promotes positive selection of conventional $\alpha\beta$ T cells, particularly those expressing TCR with low affinity to self-peptide-MHC complex (27). We have recently found that RasGRP1 also promotes the development of the invariant NKT cells (28). However, positive selection of thymocytes with relative

high affinity to self-peptide-MHC complex, such as regulatory T cells, is less dependent on RasGRP1 (29). While the importance of RasGRP1 in $\alpha\beta$ T cells is becoming clear, whether RasGRP1 functions as the upstream activator for Erk1/2 during TCR $\gamma\delta$ signaling and its importance for $\gamma\delta$ T development and activation are not well understood. In this report, we demonstrate that RasGRP1 is important for TCR-induced Erk1/2 activation in $\gamma\delta$ T cells but is dispensable for $\gamma\delta$ T generation. Deficiency of RasGRP1 does not obviously affect $\gamma\delta$ T cell numbers in the thymus but leads to increased $\gamma\delta$ T cells, particularly CD4⁻CD8⁺ $\gamma\delta$ T cells, in the peripheral lymphoid organs. Although RasGRP1 is not required for $\gamma\delta$ T cell ontogeny, it is critical for TCR-induced $\gamma\delta$ T cell proliferation and production of IL-17.

Materials and methods

Mice

The C57BL/6J mice were purchased from the Jackson Laboratory. The *RasGRP1*^{-/-} mice were previously described (25) and were backcrossed onto B6 background for 9 generations. All mice used were between 6 and 8 weeks of age according to a protocol approved by the Duke University Institute Animal Care and Use Committee. Thymocytes, splenocytes, and lymph node cells were prepared following standard procedures.

Antibodies and flow cytometry

Cells were stained with antibodies in PBS containing 2% FCS. PE-Cy7-conjugated anti-CD4 (RM4-5), APC-Cy7-conjugated anti-CD8 α (53-6.7), APC-conjugated anti-CD8 β .2 (53-5.8), FITC- or APC-conjugated anti-TCR β (H57-597), FITC- or PE-conjugated anti-TCR $\gamma\delta$ (GL3), PE-conjugated anti-IL-17A (TC11-18H10), APC-conjugated anti-IFN γ (XMG1.2), PE/Cy7-conjugated anti-ICOS (C398.4A), PE-conjugated anti-CD5 (53-7.3), APC-conjugated anti-CD25 (PC61), FITC-conjugated anti-CD122 (TM- β 1), Biotin-conjugated anti-NKG2D (C7), APC-conjugated anti-CD44 (IM7), Biotin-conjugated anti-OX40 (OX-86), Biotin-conjugated anti-CD27 (LG.3A10), APC-conjugated anti-CD90.1 (HIS51), and PE- or APC-conjugated anti-CD90.2 (30-H12) were purchased from Biolegend or BD Biosciences. Biotin-conjugated antibodies were stained with PE-, PE-Cy5-, or APC-conjugated streptavidin. For intracellular staining, cells were permeabilized using the Foxp3 staining kit (eBioscience) after cell surface staining, followed by APC-conjugated anti-TCR β or unconjugated anti-Ki-67 (B56, BD Biosciences), anti-IL-17A, or anti-IFN γ . An Alexa Fluor[®] 488-conjugated goat anti-mouse IgG (H+L) (Invitrogen) was used to detect anti-Ki67 antibodies. Anti-TCRV γ 1.1 (clone 2.11) and -TCRV δ 6.3 (clone 9D3) antibodies were previously reported (30). Cell death was identified using a 7AAD or Live/Dead[®] fixable dead cell stain kit (Invitrogen). Pacific Blue- or APC-conjugated Annexin V (BD Biosciences) was used to detect apoptotic cells. Data of stained samples were collected using a Canto II flow-cytometer (BD Biosciences) and analyzed using FlowJo software (TreeStar, Inc.).

Bone marrow reconstitution

WT C57BL/6J mice were lethally irradiated (1100 rad) four hours before adoptive transfer. Bone marrow cells from age- and sex-matched Thy1.1⁺ B6 and Thy1.2⁺ *RasGRP1*^{-/-} were mixed at a 1:1 ratio. Ten to twenty million mixed cells were then intravenously injected into each recipient mouse. The resulting chimeric mice were analyzed 7 to 8 weeks later.

Western blot

Total $\gamma\delta$ T cells were isolated from WT and *RasGRP1*^{-/-} thymocytes, splenocytes, and LN cells using MACS purification and then FACS sorting using a MoFlo Cell Sorter (Beckman Coulter), with post-sort purity >98%. Thymic and splenic $\gamma\delta$ T cells were also similarly

purified from these mice. Four hundred thousand sorted $\gamma\delta$ T cells were rested in 0.5ml PBS at 37°C for 30 minutes. Cells were then either left untreated or stimulated with an anti-CD3e antibody (500A2, 5 μ g/ml, BD Biosciences) for 5 minutes. The 500A2 antibody is capable of inducing TCR signaling without cross-linking by a secondary antibody. Cells were lysed in 1% NP-40 lysis buffer (1% NP-40, 150mM NaCl, 50mM Tris, pH 7.4) with protease and phosphatase inhibitors. Proteins in lysates were subjected to Western blot analysis with anti-phospho-Erk1/2, total Erk1/2 antibodies (Cell Signaling Technology), and an anti-RasGRP1 antibody (Santa Cruz Biotechnology). For loading control, the blots were stripped and reprobed with anti- β -actin (Sigma).

$\gamma\delta$ T cell proliferation and cytokine production

Sorted WT and *RasGRP1*^{-/-} $\gamma\delta$ T were labeled with 10 μ M CFSE at room temperature for 9 minutes as previously described (31). Cells were seeded at 5 \times 10⁵ cells/well in a 48-well plate precoated with PBS or with 10 μ g/ml soluble anti-CD3e (2C11). After incubation at 37°C for 72 hours, cells were stained for death with Live/Dead® fixable staining before being analyzed by flow cytometry.

For cytokine production, 2 \times 10⁶ freshly isolated thymocytes, splenocytes, and LN cells were seeded in each well in a 48 well-plate. The cells were left unstimulated or stimulated with PMA (25 ng/ml) and ionomycin (500 ng/ml) in the presence of a Golgi Plug for 5h. Cells were then surface stained for TCR $\gamma\delta$, TCR β , CD4, CD8, CD27, and CD44, followed by intracellular staining for IFN γ and IL-17A.

Real-time PCR

Total RNAs were extracted from sorted cells and cDNAs were obtained using the Superscript III First-Strand Synthesis System (Invitrogen). Real-time PCR was prepared using the RealMasterMix (Eppendorf) and performed on the Mastercycler®Ep realplex² system (Eppendorf). Primers for RasGRPs and actin were published previously (25) or are listed as follows: Egr1: forward, 5'-gtcctttctgacatcgctctga-3' and reverse 5'-cgagtcgttggtgggata-3'; Egr2: forward, 5'-ttgaccagatgaacggagtg-3' and reverse 5'-cagagatgggagcgaagcta-3'; Id3: forward, 5'-cacttacctgaactcaacgcc-3' and reverse 5'-cccattctcggaaaagccag-3'. Relative mRNA expression was normalized with β -actin and was presented as an arbitrary unit (a.u.) of fold change using the 2^{- $\Delta\Delta$ CT} method.

Statistics

For statistical analysis, two-tail Student *t*-tests were performed: *, p<0.05. **, p<0.01, ***, p<0.001.

Results

RasGRP1 is dispensable for intrathymic $\gamma\delta$ T cell development

We first examined mRNA levels of RasGRP1 and other RasGRP family members in $\gamma\delta$ T cells since it has been unclear whether these molecules are expressed in these cells. RasGRP1, 2, and 4 mRNAs could be detected in both $\gamma\delta$ T cells and $\alpha\beta$ T cells sorted from the thymus and spleens (Figures 1A and 1B). $\alpha\beta$ T cells appeared to express higher levels of RasGRP1 and RasGRP2 than $\gamma\delta$ T cells. The differences were particularly obvious in the peripheral lymphoid organs. In contrast, expression of RasGRP4 was similar between $\alpha\beta$ T and $\gamma\delta$ T cells. RasGRP3 was undetectable in either $\gamma\delta$ T cells or $\alpha\beta$ T cells.

To determine the role of RasGRP1 in $\gamma\delta$ T cells, we analyzed *RasGRP1*^{-/-} mice. As previously reported (21, 24), CD4⁺CD8⁻ and CD4⁻CD8⁺ single-positive (SP) thymocytes were markedly decreased in *RasGRP1*^{-/-} (KO) mice as compared to WT mice (Figures 1C

and 1D). In addition, we also found that the CD4⁺CD8⁺ double positive (DP) thymocyte number as well as total thymic cellularity of *RasGRP1*^{-/-} mice was decreased about 50% (Figure 1D). Further analysis of DN thymocytes revealed an increase in CD25⁺CD44⁻ DN3 cells and a decrease in CD25⁻CD44⁻ DN4 cells in *RasGRP1*^{-/-} mice (Figure 1E). This data suggested RasGRP1 deficiency resulted in a partial blockage of DN3 to DN4 transition, a process called β -selection due to its requirement of the pre-TCR signal. Thus, besides promoting positive selection, RasGRP1 may also be involved in pre-TCR signaling and plays a role for β -selection.

In contrast to the striking decrease in $\alpha\beta$ T cells and DP thymocytes, the overall percentage of $\gamma\delta$ T cells in the *RasGRP1*^{-/-} thymus was increased about twofold as compared to WT controls (Figures 1F and 1G). Although rare, the relative ratio of $\gamma\delta$ T cells in the CD4 or CD8 SP population was also increased. The total thymic as well as DN and CD8⁺ $\gamma\delta$ T cell subsets in *RasGRP1*^{-/-} mice were similar to WT mice. However, the CD4⁺ $\gamma\delta$ T cell number was slightly decreased in *RasGRP1*^{-/-} mice. Thus, although RasGRP1 is critical for DP to SP maturation and is important for efficient DN3 to DN4 transition, it is virtually dispensable for intrathymic $\gamma\delta$ T cell development.

Enrichment of $\gamma\delta$ T in the peripheral lymphoid organs in *RasGRP1*^{-/-} mice

As previously reported (25, 27), CD4⁺ and CD8⁺ T cell percentages and numbers in the spleen and lymph nodes (LNs) were decreased in *RasGRP1*^{-/-} mice as compared to WT mice (Figure 2A). Different from the decrease in $\alpha\beta$ T cells, the percentages and absolute numbers of the overall $\gamma\delta$ T cells did not decrease and in fact even increased in the spleen (Figure 2B, 2D) and LNs (Figure 2C, 2E) of *RasGRP1*^{-/-} mice. The percentages of splenic total and DN $\gamma\delta$ T cells were slightly increased in the spleen but more substantially increased in the LNs in *RasGRP1*^{-/-} mice. The total and DN $\gamma\delta$ T cell numbers in the WT and *RasGRP1*^{-/-} spleens were similar but were increased two- to fourfold in *RasGRP1*^{-/-} LNs. The reason for the selective increase of $\gamma\delta$ T cells in the LNs but not in the spleen is unclear but could be caused by altered expression of chemokine receptors or homing molecules such as CD62L in the absence of RasGRP1. The total CD4SP $\gamma\delta$ T cells were slightly decreased in both *RasGRP1*^{-/-} spleens and LNs. However, CD8SP $\gamma\delta$ T cells were drastically increased both in percentages and absolute numbers in both *RasGRP1*^{-/-} spleens and LNs. While CD8 α SP $\gamma\delta$ T cells accounted for around 0.5% of CD8SP T cells in WT mice, up to 40% LN and 20% splenic CD8SP cells from *RasGRP1*^{-/-} mice were $\gamma\delta$ T cells. About 90% of CD8SP $\gamma\delta$ T cells in WT and *RasGRP1*^{-/-} thymus and LN were CD8 α ⁺ β ⁺. In the spleen, CD8 α ⁺ β ⁺ cells accounted for 65% and 85% of WT and *RasGRP1*^{-/-} CD8SP $\gamma\delta$ T cells, respectively (Figure 2F). As to be demonstrated in figure 7, the CD8⁺ $\gamma\delta$ T cells were capable of producing high levels of IFN γ . At present, it is unclear whether RasGRP1 is involved in regulating CD4/CD8 expression.

Cell-intrinsic and -extrinsic mechanisms control $\gamma\delta$ T cell development in *RasGRP1*^{-/-} mice

Since the *RasGRP1*^{-/-} mice we studied were germ-line knockout and the overall T cell lymphopenia could significantly affect $\gamma\delta$ T cell development, we investigated whether $\gamma\delta$ T cell development in *RasGRP1*^{-/-} mice was caused by cell-intrinsic mechanisms. We generated and analyzed mixed bone marrow (BM) chimeric mice reconstituted with Thy1.1⁺ WT and Thy1.2⁺ *RasGRP1*^{-/-} BM cells at a 1:1 ratio. Chimeric mice were analyzed 8 weeks after transfer. The ratio of Thy1.1⁺ WT to Thy1.2⁺ *RasGRP1*^{-/-} CD4⁻CD8⁻ DN thymocytes were about 2:1 (Figure 3A), suggesting that RasGRP1 may be involved in early T cell development. Within the DN population, Thy1.2⁺ *RasGRP1*^{-/-} CD25⁺CD44⁻ DN3 and CD25⁻CD44⁻ DN4 cells were substantially increased and decreased, respectively, as compared to Thy1.1⁺ WT controls (Figure 3B), indicating that RasGRP1 is intrinsically

important for efficient β -selection. These observations, together with those demonstrating critical roles of RasGRP1 for DP to SP maturation (25), indicate RasGRP1 plays crucial roles at multiple stages for $\alpha\beta$ T cell development in a cell-autonomous manner. Within TCR $\alpha\beta^+$ T cells, most were Thy1.1⁺ WT and Thy1.2⁺ *RasGRP1*^{-/-} cells were virtually undetectable (Figure 3C). However, the ratio of $\gamma\delta$ T cells originated from *RasGRP1*^{-/-} to WT BM was similar to the ratio of total DN cells between these two origins. As mentioned earlier, CD8⁺ $\gamma\delta$ T cells were drastically increased in *RasGRP1*^{-/-} mice, but such an increase was not observed in the chimeric mice (Figure 3D). Thus, the overall RasGRP1-independent generation of $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice was cell intrinsic. However, the expansion of CD8⁺ $\gamma\delta$ T cells in these mice was not cell intrinsic.

Differential effects of RasGRP1 deficiency on $\alpha\beta$ T and $\gamma\delta$ T cell homeostasis

The TCR-mediated Ras-Erk1/2 pathway plays a crucial role in T cell survival and proliferation. One possibility that may cause increased $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice would be that they have proliferative and/or survival advantage over $\alpha\beta$ T cells. Previous studies have demonstrated that RasGRP1 prevents conventional CD4⁺ T cells and CD8⁺ T cells as well as α NKT cells from apoptosis (28, 29). By staining freshly isolated cells with Annexin-V and 7-AAD, we also observed increased apoptosis of *RasGRP1*^{-/-} $\alpha\beta$ T cells. However, no increase of apoptosis was observed in *RasGRP1*^{-/-} $\gamma\delta$ T cells as compared with WT controls. In fact, slight decreases of apoptosis were observed in $\gamma\delta$ T cells from *RasGRP1*^{-/-} spleen and LNs (Figure 4A).

In *RasGRP1*^{-/-} mice, the total peripheral $\alpha\beta$ T cell numbers are decreased. Under such a lymphopenic condition, T cells may undergo homeostatic proliferation. Expression of nuclear Ki67, a protein that is upregulated in actively cycling cells, was obviously increased in freshly isolated *RasGRP1*^{-/-} $\alpha\beta$ T cells as compared to WT controls (Figure 4B). Consistent with increased Ki67 expression in *RasGRP1*^{-/-} $\alpha\beta$ T cells, most of these cells were CD44^{high}CD62L⁻, which was consistent with homeostatic proliferation of these cells (Figure 4C). In contrast, Ki67 expression in *RasGRP1*^{-/-} $\gamma\delta$ T cells was similar to WT control (Figure 4B). Furthermore, most *RasGRP1*^{-/-} $\gamma\delta$ T cells displayed a naïve CD44^{low}CD62L⁺ phenotype (Figure 3D). Of note, Ki67 positive CD4SP *RasGRP1*^{-/-} $\gamma\delta$ T cells were increased as compared to WT controls (Figure 4B), likely caused by homeostatic proliferation due to the decrease of this population of $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice.

Together, these observations suggest that *RasGRP1*^{-/-} $\alpha\beta$ T cells undergo enhanced homeostatic proliferation *in vivo* and that there is no obvious increase in homeostatic proliferation of most $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice except the CD4SP subset. Our data also suggest that $\gamma\delta$ T cells may sense different homeostatic cues compared with $\alpha\beta$ T cells since the overall $\alpha\beta$ T cell lymphopenia resulted in increased homeostatic proliferation of $\alpha\beta$ T but not $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice. The enrichment of $\gamma\delta$ T cells in the absence of RasGRP1 is likely an active process involved in $\gamma\delta$ T cell generation since the percentage of intracellular TCR β positive $\gamma\delta$ T cells was significantly decreased in *RasGRP1*^{-/-} mice (Figure 4E).

Requirement of RasGRP1 for TCR-induced Erk1/2 activation in $\gamma\delta$ T cells

As mentioned earlier in Figure 1A, in addition to RasGRP1, RasGRP2 and 4 are also expressed in $\gamma\delta$ T cells. We examined whether RasGRP1 deficiency may affect RasGRP2 and RasGRP4 expression. As shown in Figure 5A, RasGRP2 mRNA level decreased about 50% while RasGRP4 mRNA level increased about twentyfold in *RasGRP1*KO $\gamma\delta$ T cells compared to WT controls. Increased Erk1/2 activity has been found to promote $\gamma\delta$ T cell development (21, 23). It is further known that RasGRP1 plays an important role in TCR-induced Erk1/2 activation in total thymocytes and $\alpha\beta$ T cells (25, 27). The minimal

requirement of RasGRP1 for $\gamma\delta$ T cell development and the drastic increase in RasGRP4 expression in *RasGRP1KO* $\gamma\delta$ T cells raise the possibility that RasGRP1 may be dispensable for TCR-induced Erk1/2 activation, or elevated RasGRP4 expression may compensate the loss of RasGRP1 for TCR-induced Erk1/2 activation in $\gamma\delta$ T cells. To determine the role of RasGRP1 for TCR-induced Erk1/2 activation, we sorted $\gamma\delta$ T cells from *RasGRP1^{-/-}* and WT mice. Sorted cells were left unstimulated or stimulated with a soluble anti-CD3 antibody, and Erk1/2 phosphorylation was examined by Western blot analysis. As shown in Figure 5B, Erk1/2 phosphorylation was decreased in *RasGRP1^{-/-}* $\gamma\delta$ T cells. However, induction of Erk1/2 phosphorylation could still be detected in *RasGRP1^{-/-}* $\gamma\delta$ T cells. Thus, RasGRP1 is required for optimal TCR-induced Erk1/2 activation in $\gamma\delta$ T cells, and other Ras activation mechanisms are not able to fully compensate for the loss of RasGRP1 for TCR-induced Erk1/2 activation.

Activation of the Ras-Erk1/2 pathway leads to induction of Egr1-3, which promotes Id3 transcription. Both Id3 and Egr1/2 are involved in $\gamma\delta$ T cell development. Id3-deficient mice contain increased $\gamma\delta$ T cells that are dominantly V γ 1⁺V δ 6.3⁺ innate $\gamma\delta$ T cells, which display an activating phenotype (32-36). Egr1/2 and Id3 expression was substantially decreased in *RasGRP1^{-/-}* $\gamma\delta$ T cells (Figure 5C). The decreased expression of Id3 cells may contribute to the approximately twofold increase in V γ 1⁺V δ 6.3⁺ $\gamma\delta$ T cells in *RasGRP1^{-/-}* mice (Figures 5D, 5E).

Requirement of RasGRP1 for TCR-induced $\gamma\delta$ T cell proliferation

RasGRP1 activates the Ras-Erk1/2 pathway in peripheral T cells and is critical for peripheral $\alpha\beta$ T cell activation (25-27). The dispensable role of RasGRP1 for $\gamma\delta$ T cell development and homeostasis raises the question whether this protein participates in $\gamma\delta$ T cell activation. Due to downregulation of $\gamma\delta$ TCR following TCR activation, we sorted $\gamma\delta$ T cells and labeled them with CFSE. CFSE-labeled $\gamma\delta$ T cells were left unstimulated or stimulated with plate-bound anti-CD3 for 72 hours. While WT $\gamma\delta$ T cells proliferated vigorously, *RasGRP1^{-/-}* $\gamma\delta$ T cells were defective in proliferation (Figure 6A). No obvious difference in cell death was observed between WT and *RasGRP1KO* $\gamma\delta$ T cells during *in vitro* stimulation (Figure 6B), suggesting that defective $\gamma\delta$ T cell proliferation in the absence of RasGRP1 was not caused by increased cell death. In WT mice, anti-CD3 ϵ -induced Erk1/2 phosphorylation in splenic $\gamma\delta$ T cells was stronger than in thymic $\gamma\delta$ T cells. Both thymic and splenic *RasGRP1KO* $\gamma\delta$ T cells displayed decreased Erk1/2 phosphorylation as compared to WT control (Figure 6C). Given the role of the Ras-Erk1/2 pathway for $\alpha\beta$ T cell activation, impaired Erk1/2 activation in *RasGRP1KO* $\gamma\delta$ T cells may lead to their proliferative defect. *RasGRP1KO* $\gamma\delta$ T cells expressed slightly increased CD25 and CD122 compared to WT control. They expressed similar levels of costimulatory molecules ICOS, CD5, OX40, and NKG2D as compared to WT controls (Figure 6D), suggesting that RasGRP1 is not required for the expression of these molecules. However, whether RasGRP1 is involved in costimulatory signaling in $\gamma\delta$ T cells is unclear at present. Together, these observations indicate that RasGRP1 is crucial for TCR-induced $\gamma\delta$ T cell proliferation.

Defective IL-17 production by $\gamma\delta$ T cells in the absence of RasGRP1

$\gamma\delta$ T cells are an important innate source of IL-17 and IFN γ (37-39). To examine whether RasGRP1 deficiency affects cytokine production by $\gamma\delta$ T cells, we stimulated freshly isolated thymocytes, splenocytes, and LN cells with PMA and ionomycin for 5 hours in the presence of a Golgi plug, and we then analyzed cytokine production by intracellular staining. IL-17A production by *RasGRP1^{-/-}* $\gamma\delta$ T cells from the thymus, spleen, and LN was substantially decreased (Figure 7A). Different from IL-17, IFN γ production was increased in *RasGRP1^{-/-}* $\gamma\delta$ T cells as compared to WT $\gamma\delta$ T cells. Further comparison of cytokine expression in CD4⁺, CD8⁺, and DN $\gamma\delta$ T cells revealed that CD8⁺ $\gamma\delta$ T cells produced

higher levels of IFN γ but lower levels of IL-17A than CD4⁺ and DN $\gamma\delta$ T cells in WT mice. In the absence of RasGRP1, IFN γ expression was increased but IL-17A expression was substantially decreased in these $\gamma\delta$ T cell populations (Figure 7B).

$\gamma\delta$ T cells can be divided into functional subsets based on CD27 and CD44 expression, with the CD27⁻CD44⁺ mainly producing IL-17 and the CD27⁺CD44^{+/-} producing IFN γ (40). In *RasGRP1KO* thymus and spleen, the percentages of CD27⁻CD44⁺ $\gamma\delta$ T cells were decreased about 50% and 30% respectively as compared to WT controls, suggesting that RasGRP1 is important for the generation/maintenance of this population of $\gamma\delta$ T cells (Figure 7C). Moreover, IL-17 production by *RasGRP1KO* CD27⁻CD44⁺ $\gamma\delta$ T cells was substantially decreased compared to WT control. In contrast to CD27⁻CD44⁺ $\gamma\delta$ T cells, there were about 50%-70% increases in CD27⁺CD44⁻ $\gamma\delta$ T cells in *RasGRP1KO* thymus and spleen. Moreover, IFN γ production by splenic and LN CD27⁺CD44^{+/-} $\gamma\delta$ T cells from *RasGRP1KO* mice appeared to increase (Figure 7D). Together, these observations revealed that RasGRP1 is not only important for the generation/maintenance of the CD27⁻CD44⁺ $\gamma\delta$ T cell subset but also critical for this subset of cells to produce IL-17.

Discussion

An important signaling event following both $\alpha\beta$ TCR and $\gamma\delta$ TCR stimulation is the activation of Erk1/2. In $\alpha\beta$ T cells, RasGRP1 is a critical upstream activator for Erk1/2 through the Ras-Raf-Mek1/2 signaling cascade following TCR engagement (27). Genetic evidence has revealed that both RasGRP1 and Erk1/2 are critical for maturation of T cells adopting the $\alpha\beta$ T cell lineage, including conventional $\alpha\beta$ T cells (25, 27, 41), α NKT cells (28), and Treg (29). In this report, we have demonstrated that RasGRP1 is important for Erk1/2 in $\gamma\delta$ T cells following TCR stimulation. Thus, although additional RasGRPs and other guanine-nucleotide-releasing factors such as Sos are expressed in $\gamma\delta$ T cells, they cannot fully compensate for the loss of RasGRP1 for Erk1/2 activation. Extending previous studies demonstrating that RasGRP1 is critical for positive selection of $\alpha\beta$ T cells (25, 27), we now find that RasGRP1 is also important for efficient β -selection as evidenced by the accumulation of DN3 cells and decreased DN4 thymocytes, suggesting that RasGRP1 participates in pre-TCR signaling.

In *RasGRP1*^{-/-} mice, total as well as CD4⁻CD8⁻ DN and CD8 SP $\gamma\delta$ T cell numbers in the thymus are comparable to WT control. The virtually normal development of *RasGRP1*^{-/-} $\gamma\delta$ T cells in mixed chimeric mice further supports the minimal role of RasGRP1 for $\gamma\delta$ T cell development. CD4⁻CD8⁻ DN $\gamma\delta$ T cells, and CD8 SP $\gamma\delta$ T cells in particular, expand in *RasGRP1*^{-/-} peripheral lymphoid organs. However, such expansion of *RasGRP1*^{-/-} $\gamma\delta$ T cells is not observed in mixed BM chimeric mice, suggesting that cell-extrinsic factors lead to the expansion of $\gamma\delta$ T cells in the peripheral lymphoid organs in *RasGRP1*^{-/-} mice. These extrinsic factors could be cytokines produced by *RasGRP1*^{-/-} conventional $\alpha\beta$ T cells under the lymphopenic environment or by other cell types. Furthermore, CD4⁺Foxp3⁺ Treg numbers in *RasGRP1*^{-/-} mice were decreased (29), which could also contribute to peripheral $\gamma\delta$ T cell expansion.

It has been reported that TCR signal strength influences $\alpha\beta$ T and $\gamma\delta$ T cell lineage commitment with strong TCR signal and enhanced Erk1/2 activity favoring the $\gamma\delta$ T lineage (21-23). It is intriguing that $\gamma\delta$ T cell development is virtually intact in *RasGRP1*^{-/-} mice with an obvious decrease in Erk1/2 activation. One potential explanation is that the relative strengths of Erk1/2 signaling dictate $\gamma\delta$ T versus $\alpha\beta$ T differentiation. The commitment of the progenitor to the $\alpha\beta$ T lineage or cells that have committed to the $\alpha\beta$ T fate could be more sensitive to the decreased Erk1/2 activity than those that are committing to or have committed to the $\gamma\delta$ T lineage. It is important to note that our data does not rule out the role

of Erk1/2 for $\gamma\delta$ T cell development since a low level of Erk1/2 phosphorylation can still be induced in *RasGRP1*^{-/-} $\gamma\delta$ T cells. Furthermore, TCR stimulation induces stronger Erk1/2 activation in peripheral $\gamma\delta$ T cells than thymic $\gamma\delta$ T cells, suggesting that differential regulation of Erk1/2 activation between thymic and peripheral $\gamma\delta$ T cells. The low level of TCR-induced Erk1/2 activity in thymic $\gamma\delta$ T cells may suggest the possibility that developing $\gamma\delta$ T cells can be less sensitive to the decrease of Erk1/2 activation in the absence of RasGRP1 than peripheral $\gamma\delta$ T cells. Additionally, we have recently demonstrated that in addition to Erk1/2, RasGRP1 also functions as an upstream activator for PI3K/Akt and mTOR in thymocytes. In *RasGRP1*^{-/-} thymocytes, not only Erk1/2 but also PI3K/Akt and mTOR activation is impaired following TCR engagement (42, 43). RasGRP1 could promote $\alpha\beta$ T cell development not only via Erk1/2 but also through other downstream signaling pathways.

The Ras-Erk1/2 pathway plays an important role in transcriptional activation of *Egr1/2*, which in turn promotes *Id3* expression (44). In *RasGRP1*-deficient $\gamma\delta$ T cells, *Egr1/2* and *Id3* expression is decreased, which is consistent with the importance of RasGRP1 for Erk1/2 activation in $\gamma\delta$ T cells. Proper expression of *Id3* ensures normal $\gamma\delta$ T cell development. Loss of *Id3* leads to preferential generation and expansion of “innate” $\gamma\delta$ T cells. However, elevated *Id3* level also leads to increases in $\gamma\delta$ T cells (32, 34-36, 45). These data suggest a narrow window of *Id3* expression for proper development of innate $\gamma\delta$ T cells. Although $\gamma\delta$ T cells are enriched in *Id3*^{-/-} mice and are relatively enriched in *RasGRP1*^{-/-} mice, differences between these two strains are obvious. In *Id3*^{-/-} mice, the $V\gamma 1^+V\delta 6.3^+$ innate $\gamma\delta$ T cells are predominantly increased (32, 34, 36). In *RasGRP1*^{-/-} mice, there is less than a twofold increase in such innate $\gamma\delta$ T cells. The differences between these two strains of mice can be explained by the influence of RasGRP1 deficiency on multiple downstream effector molecules and the incomplete loss of *Id3* expression.

Most $\gamma\delta$ T cells in adult WT mice are CD4⁻CD8⁻ DN. Although CD4SP or CD8SP $\gamma\delta$ T cells can be easily detected in fetal thymus, they are rare in postnatal thymus, spleen, and LN (48, 49). In *RasGRP1*^{-/-} mice, CD8SP $\gamma\delta$ T cells are substantially increased but CD4SP $\gamma\delta$ T cells are decreased. In both WT and *RasGRP1*^{-/-} thymus, spleen, and LN, most of the CD8SP $\gamma\delta$ T cells express the CD8 α and β heterodimer. Although $\gamma\delta$ T cells are not MHC-I restricted and extracellular factors play a critical role in the expansion of CD8SP $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice, expression of the CD8 $\alpha\beta$ heterodimer could be involved in the expansion of these CD8SP $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice since the CD8 $\alpha\beta$ heterodimer is able to associate with MHC-I. Of note, IL-7 has been reported to be able to expand CD8 $\alpha^+\beta^+$ $\gamma\delta$ T cells from WT fetal thymus (50). It would be interesting to determine whether IL-7 level is elevated and is responsible for the expansion of CD8SP $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice. Enrichment of CD8SP $\gamma\delta$ T cells has also been observed in several other genetically manipulated mice such as *Id3*^{-/-} mice (34). Those *Id3*^{-/-} CD8SP $\gamma\delta$ T cells express the CD8 $\alpha\alpha$ homodimer and are thus different from the *RasGRP1*^{-/-} CD8 $\alpha^+\beta^+$ $\gamma\delta$ T cells. It is possible that combined action of altered intrinsic properties, such as abnormal expression of *Id3* and other molecules, and extrinsic factors causes the expansion of CD8SP $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice.

RasGRP1 is differentially required for $\alpha\beta$ versus $\gamma\delta$ T cell development. However, it is critical for both $\alpha\beta$ and $\gamma\delta$ T cell activation. RasGRP1 may exert its role in $\gamma\delta$ T cell activation through multiple mechanisms. For example, it can directly promote mature $\gamma\delta$ T cell activation by activating the Ras-Erk1/2 pathway, mTOR, and PI3K-Akt. Alternatively but not mutually exclusively, RasGRP1 may be required for licensing or arming $\gamma\delta$ T cells during their development to ensure competence for activation after they mature, a phenomenon that occurs during NK cell development (46, 47). Further studies utilizing conditional *RasGRP1*-deficient mice will help to distinguish such possibilities.

IL-17-producing $\gamma\delta$ T cells play important roles during microbial infection and the pathogenesis of autoimmune and inflammatory diseases (51-53). Generation of IL-17-producing $\gamma\delta$ T cells appears to be developmentally programmed during intrathymic development that is at least partially dependent on ROR γ t and the Notch-Hes pathway but is independent on Stat3 (54). RasGRP1-deficient mice contain decreased CD27⁻CD44⁺ subset but increased CD27⁺CD44⁻ subset in both thymus and peripheral lymphoid organs, suggesting that RasGRP1 is involved in differentiation of $\gamma\delta$ T cells into the different effector subsets. Although it is most likely that RasGRP1 mediates $\gamma\delta$ TCR signaling to regulate $\gamma\delta$ effector T cell differentiation, we cannot rule out that RasGRP1 may be involved in other receptor signaling to control $\gamma\delta$ T differentiation. Within the CD27⁻CD44⁺ RasGRP1KO $\gamma\delta$ T cells, the ratio of IL-17A-producing cells is also decreased, suggesting that RasGRP1 participates in IL-17A production by these cells. At present, it is unclear how RasGRP1 promotes IL-17 production by $\gamma\delta$ T cells. In $\alpha\beta$ T cells, mTOR complex 1 signaling is critical for Th17 differentiation (55, 56). Interestingly, RasGRP1 is critical for mTOR activation in T cells by activating Ras-Erk1/2 signaling (42, 43). Future studies should determine whether RasGRP1 promotes IL-17-producing $\gamma\delta$ T cells through inducing mTOR activation and whether RasGRP1 can be a potential target for $\gamma\delta$ T-cell-mediated diseases.

In summary, we identified several important functions of RasGRP1 in T cells. We demonstrated that RasGRP1 is involved in $\alpha\beta$ T cell maturation from the DN to DP stage, is important for TCR-induced Erk1/2 activation in both thymic and splenic $\gamma\delta$ T cells, is dispensable for overall $\gamma\delta$ T cell development but important for the generation/maintenance of IL-17-producing $\gamma\delta$ T cells, and, finally, is critical for TCR-induced $\gamma\delta$ T cell proliferation.

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Abbreviations

TCR	T cell receptor
mTOR	mammalian target of rapamycin
SP	single positive
DN	double negative
DP	double positive

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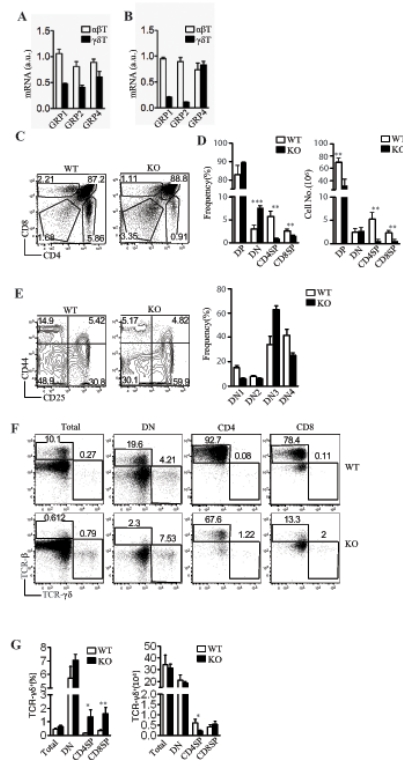


Figure 1. RasGRP1 is dispensable for $\gamma\delta$ T cell development in the thymus

(A, B) Expression of RasGRPs in $\alpha\beta$ T and $\gamma\delta$ T cells. mRNA levels of RasGRPs from sorted $\alpha\beta$ T and $\gamma\delta$ T cells from thymus (A) and spleen (B) were determined by real-time qPCR. (C) CD4 and CD8 staining of total thymocytes from WT and *RasGRP1*^{-/-} mice. (D) Decrease of DP and SP thymocytes in *RasGRP1*^{-/-} mice. Data shown are mean \pm SEM. (E) Inefficient DN3 to DN4 transition in the absence of *RasGRP1*^{-/-}. Representative FACS plots of CD25 and CD44 expression in gated CD4⁻CD8⁻ DN thymocytes are shown. (F) Representative FACS plots of TCR- $\gamma\delta$ and TCR- β staining of the indicated thymocytes. Total thymocytes as well as gated DN and SP cells are shown. (G) Percentages (left panel) and absolute number (right panel) of $\gamma\delta$ T cells within total, CD4SP, CD8SP, and DN thymocytes in WT and *RasGRP1*^{-/-} mice. Mice used in the study were 6-8 weeks of age. Data shown are representative of at least three experiments. * P <0.05, ** P <0.01, *** P <0.001 determined by the Student t -test.

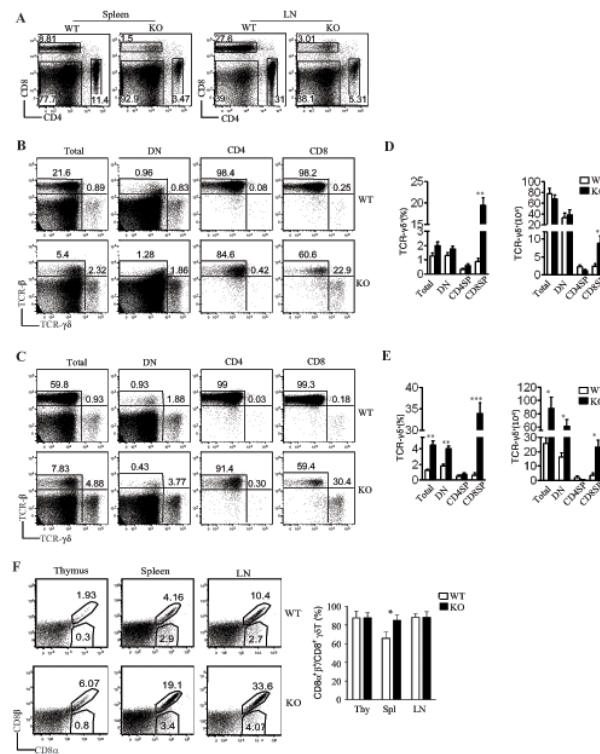


Figure 2. Enrichment of $\gamma\delta$ T cells in the peripheral lymphoid organs in *RasGRP1*^{-/-} mice
(A) Decreased CD4 and CD8 T cells in *RasGRP1*^{-/-} spleen and LNs. Representative dot plots of CD4 and CD8 staining of live-gated cells are shown. **(B, C)** Representative dot plot for assessment of $\gamma\delta$ T and $\alpha\beta$ T cells in the spleens (B) and LNs (C) in *RasGRP1*^{-/-} and WT mice. **(D, E)** Mean \pm SEM presentation of percentages and absolute numbers of $\gamma\delta$ T cells in the spleens (D) and LNs (E). **(F)** Expression of CD8 α and CD8 β heterodimer in $\gamma\delta$ T cells. Representative dot plots of CD8 α and CD8 β expression on gated TCR β ⁺TCR $\gamma\delta$ ⁺ cells from WT and *RasGRP1*^{-/-} thymus (Thy), spleen (spl), and LNs (LN). Bar graph shows mean \pm SEM (n=3) of percentages of CD8 α ⁺ β ⁺ cells within the CD8 α ⁺ $\gamma\delta$ T cells. Data shown are representative of at least three experiments. **P*<0.05, ***P*<0.01, ****P*<0.001 determined by the Student *t*-test.

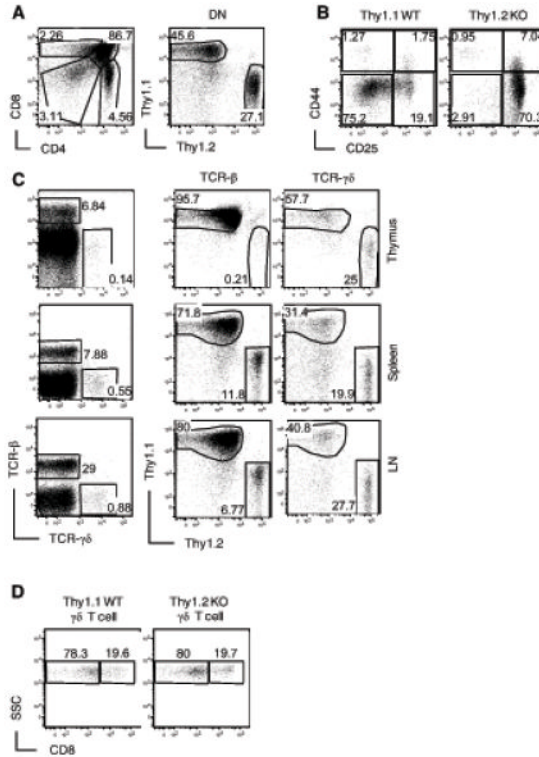


Figure 3. Contribution of cell-intrinsic and -extrinsic mechanisms for $\gamma\delta$ T cell development in *RasGRP1*-deficient mice

(A) Left panel, CD4 and CD8 α staining on total live thymocytes from lethally irradiated recipient mice 8 weeks after reconstituted with WT (Thy1.1) and *RasGRP1*^{-/-} (Thy1.2) BM. Right panels, Thy1.1 and Thy1.2 expression on gated CD4⁻CD8⁻ DN thymocytes. (B) Impaired β -selection in the absence of *RasGRP1*. Dot plots show CD44 and CD25 staining on gated Thy1.1⁺ WT and Thy1.2⁺ *RasGRP1*^{-/-} CD4⁻CD8⁻ DN thymocytes from the recipient mice. (C) Left panel, TCR β and TCR $\gamma\delta$ staining of live-gated thymocytes from recipient mice. Middle and right panels, Thy1.1 and Thy1.2 staining on gated TCR β ⁺ and TCR $\gamma\delta$ ⁺ thymocytes, respectively. (D) CD8 α staining of Thy1.1⁺ WT and Thy1.2⁺ *RasGRP1*^{-/-} $\gamma\delta$ T cells in the chimeric mice. Data are representative of three experiments.

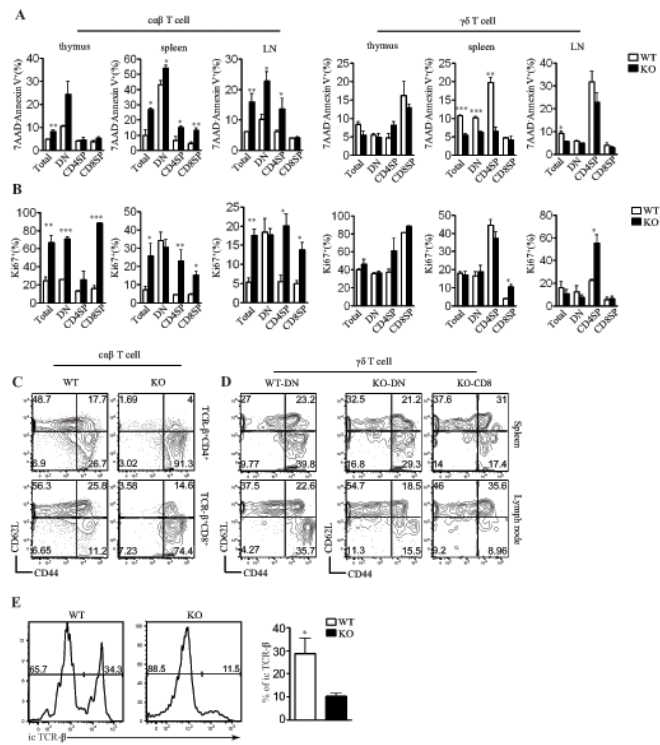


Figure 4. Assessment of T cell survival and homeostatic proliferation

(A) Increased apoptosis of $\alpha\beta$ T but not $\gamma\delta$ T cells in the absence of RasGRP1. Freshly isolated thymocytes, splenocytes, and LN cells from WT and *RasGRP1*^{-/-} mice were stained with anti-CD4, anti-CD8, anti-TCR β , anti-TCR $\gamma\delta$, 7AAD, and annexin V. Bar figures show mean \pm SEM of apoptotic cells within the indicated populations. (B) Increased Ki67⁺ $\alpha\beta$ T but not $\gamma\delta$ T cells in the absence of RasGRP1. Cells were similarly stained and analyzed as described in (A), except that Ki67 were determined by intracellular staining. (C, D) CD44 and CD62L expression on $\alpha\beta$ T (C) and $\gamma\delta$ T cells (D) from WT and *RasGRP1*^{-/-} mice. Data from WT CD8⁺ $\gamma\delta$ T cells was not presented since they were rare. (E) Decreased intracellular TCR β ⁺ cells within the $\gamma\delta$ T cells in the absence of RasGRP1. Histograms show intracellular staining of TCR β in gated $\gamma\delta$ T cells from WT and *RasGRP1*^{-/-} thymus. Data shown represent three experiments. **P*<0.05, ***P*<0.01, ****P*<0.001 determined by the Student *t*-test.

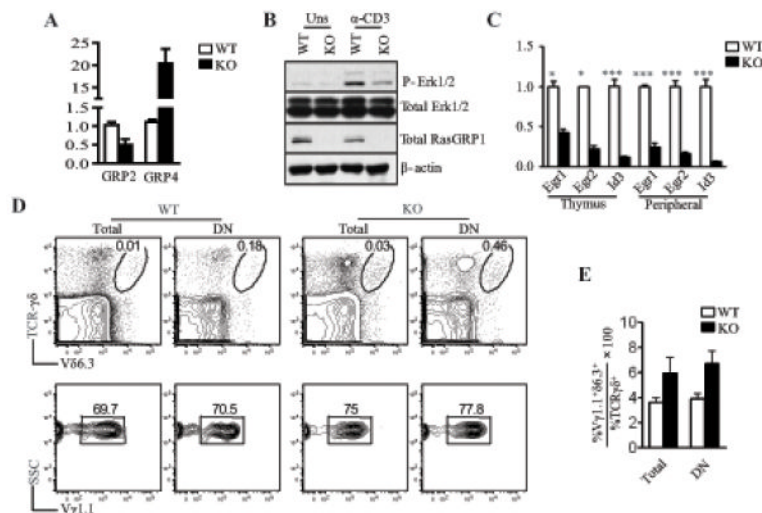


Figure 5. Requirement of RasGRP1 for TCR-induced Erk1/2 activation in $\gamma\delta$ T cells
(A) Altered RasGRP2 and RasGRP4 expression in RasGRP1KO $\gamma\delta$ T cells. mRNA levels of *RasGRP2* and *RasGRP4* in sorted WT and *RasGRP1*KO $\gamma\delta$ T cells were determined by qRT-PCR. **(B)** Decreased Erk1/2 activation in *RasGRP1*KO $\gamma\delta$ T cells. Sorted $\gamma\delta$ T cells from WT and *RasGRP1*^{-/-} mice were rested in PBS for 30 minutes and then left unstimulated or stimulated with an anti-CD3e antibody for 5 minutes. Erk1/2 phosphorylation was determined by immunoblotting analysis. **(C)** Decreased Egr1/2 and Id3 expression in *RasGRP1*^{-/-} $\gamma\delta$ T cells. RNA isolated from WT and *RasGRP1*^{-/-} $\gamma\delta$ T cells were reversely transcribed, and mRNA levels of indicated molecules were determined by real-time qPCR. Data shown are representative of two experiments. **(D)** Slight increase in V γ 1⁺V66.3⁺ $\gamma\delta$ T cells in *RasGRP1*^{-/-} mice. Upper panels show TCR $\gamma\delta$ and TCRV66.3 staining of total thymocytes from WT and *RasGRP1*^{-/-} mice. Lower panels show TCRV γ 1.1 staining of gated TCR $\gamma\delta$ ⁺TCRV66.3⁺ cells. **(E)** Percentages of TCRV γ 1.1⁺V66.3⁺ cells within the $\gamma\delta$ T cells. Data shown are representative of two (A-C) and three (D, E) experiments.

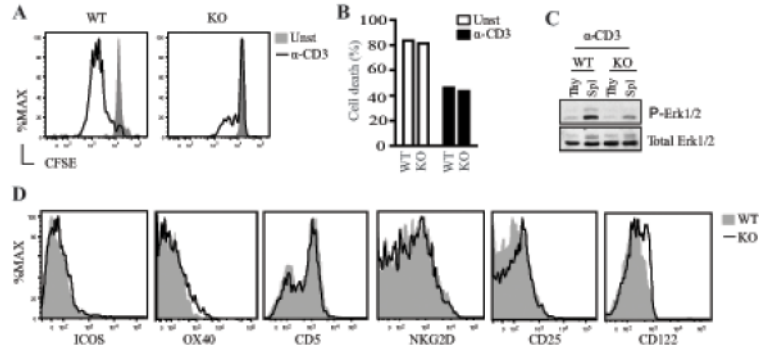


Figure 6. Requirement of RasGRP1 for TCR-induced $\gamma\delta$ T cell proliferation

(A, B) CFSE dilution assay to assess $\gamma\delta$ T cell proliferation and death. CFSE-labeled purified WT and *RasGRP1*^{-/-} $\gamma\delta$ T and $\alpha\beta$ T cells were left unstimulated or stimulated with a plate-bound anti-CD3e antibody for 72 hours. Cultured cells were stained with Live/Dead before analyzed by flow cytometry. Histograms show CFSE intensity on gated live $\gamma\delta$ T cells (A). Bar figure shows death rate of cultured $\gamma\delta$ T cells (B). (C) Impaired Erk1/2 activation in both thymic and splenic $\gamma\delta$ T cells in the absence of RasGRP1. (D) Expression of costimulatory molecules and cytokine receptors by *RasGRP1*KO $\gamma\delta$ T cells. Data shown are representative of two (A-C) and three (D) experiments.

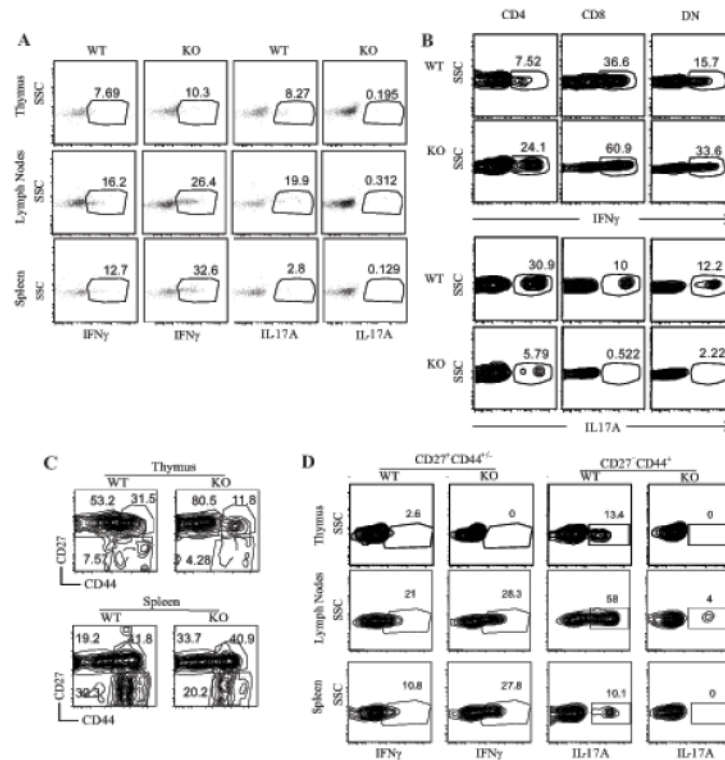


Figure 7. Defective IL-17 production by *RasGRP1*^{-/-} $\gamma\delta$ T cells

(A) WT and *RasGRP1*^{-/-} thymocytes, splenocytes, and LN cells were stimulated with PMA and ionomycin in the presence of Golgi plug for 5 hours. Stimulated cells were cell surface stained for TCR β and TCR $\gamma\delta$, followed by intracellular cytokine staining for IFN γ and IL17A. Dot plots show IL-17A and IFN γ expression in gated total TCR β ⁺TCR $\gamma\delta$ ⁺ cells. (B) IFN γ and IL-17A expression in CD4⁺, CD8 α ⁺, and DN $\gamma\delta$ T cells. LN cells were similarly treated and stained as in (A) with the addition of CD4 and CD8 α staining. Contour-plots show IFN γ and IL-17A staining in gated WT and *RasGRP1*^{-/-} CD4⁺, CD8 α ⁺, and DN $\gamma\delta$ T cells. (C) $\gamma\delta$ T cell subsets based on *ex vivo* CD27 and CD44 staining. Contour plots show CD27 and CD44 expression in gated TCR β ⁺TCR $\gamma\delta$ ⁺CD4⁻CD8⁻ thymocytes and splenocytes from WT and *RasGRP1*^{KO} mice. (D) Differential effects of *RasGRP1* deficiency on IL-17A production by CD27⁻CD44⁺ $\gamma\delta$ T cells and IFN γ expression by CD27⁺CD44⁺ $\gamma\delta$ T cells. Data shown represent three experiments.