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Tec kinase Itk in $\gamma\delta$ T cells is pivotal for controlling IgE production in vivo

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In conventional $\alpha\beta$ T cells, the Tec family tyrosine kinase Itk is required for signaling downstream of the T cell receptor (TCR). Itk also regulates $\alpha\beta$ T cell development, lineage commitment, and effector function. A well established feature of *Itk*^{-/-} mice is their inability to generate T helper type 2 (Th2) responses that produce IL-4, IL-5, and IL-13; yet these mice have spontaneously elevated levels of serum IgE and increased numbers of germinal center B cells. Here we show that the source of this phenotype is $\gamma\delta$ T cells, as normal IgE levels are observed in *Itk*^{-/-}*Tcrd*^{-/-} mice. When stimulated through the $\gamma\delta$ TCR, *Itk*^{-/-} $\gamma\delta$ T cells produce high levels of Th2 cytokines, but diminished IFN γ . In addition, activated *Itk*^{-/-} $\gamma\delta$ T cells up-regulate costimulatory molecules important for B cell help, suggesting that they may directly promote B cell activation and Ig class switching. Furthermore, we find that $\gamma\delta$ T cells numbers are increased in *Itk*^{-/-} mice, most notably the V γ 1.1⁺V δ 6.3⁺ subset that represents the dominant population of $\gamma\delta$ NKT cells. *Itk*^{-/-} $\gamma\delta$ NKT cells also have increased expression of PLZF, a transcription factor required for $\alpha\beta$ NKT cells, indicating a common molecular program between $\alpha\beta$ and $\gamma\delta$ NKT cell lineages. Together, these data indicate that Itk signaling regulates $\gamma\delta$ T cell lineage development and effector function and is required to control IgE production in vivo.

T cell development | T cell differentiation | T cell signaling

The Tec family tyrosine kinase Itk is important for signaling downstream of the T cell receptor (1). In particular, Itk-deficient T cells have defects in phospholipase C- γ (PLC- γ) phosphorylation, calcium mobilization, mitogen-activated protein kinase (MAP kinase) activation, and AP-1 and nuclear factor of activated T cells (NFAT) activation after T cell receptor (TCR) stimulation. Itk is also critical for conventional $\alpha\beta$ T cell development, selection, and function. Of particular importance, Itk signaling regulates CD4⁺ T helper cell differentiation, playing a key role in the development of Th2 responses (2). Based on this well-documented defect of *Itk*^{-/-} mice in generating Th2 effector responses and cytokine production, it was surprising to discover that these mice had spontaneously elevated levels of serum IgE (3, 4), as B cell isotype switching to IgE is highly dependent on Th2 cytokines IL-4 and IL-13 (5). As our previous studies had indicated that *Itk*^{-/-} $\alpha\beta$ TCR⁺ NKT cells (referred to as $\alpha\beta$ NKT cells) were also highly defective in producing effector cytokines such as IL-4 (6), we considered the possibility that $\gamma\delta$ TCR⁺ NKT cells were the major source of Th2 cytokines in *Itk*^{-/-} mice.

The $\gamma\delta$ T cells are a highly conserved subset of T cells that constitutes 1–5% of the lymphocytes in the blood and peripheral organs of mice but can account for up to 50% of the lymphocytes in the mucosal epithelia. As with other subsets of “innate” T cells, $\gamma\delta$ T cells express memory cell surface markers (7), and are capable of rapidly secreting effector cytokines (8). Among the many functions attributed to $\gamma\delta$ T cells, a great deal of recent interest has focused on their ability to modulate adaptive immune responses, specifically the humoral response (9).

A variety of studies have indicated that $\gamma\delta$ T cells are able to provide help for B cell responses. Initial studies performed in mice lacking $\alpha\beta$ T cells showed that B cell expansion, differentiation, and secretion of ‘T-dependent’ antibody isotypes, IgE,

and IgG₁, were all intact in these mice (10). Furthermore, TCR β ^{-/-} mice challenged repeatedly with parasitic infections could produce germinal centers and generate increased antibody production (11). Using a model of pulmonary allergic inflammation, decreased production of IgE and IgG₁ was seen in mice lacking $\gamma\delta$ T cells compared with WT mice (12). The $\gamma\delta$ T cells have also been shown to directly induce germinal center formation and Ig hypermutation (13). Interestingly, even though the $\gamma\delta$ T cells expressing CD4 account for only 5–10% of all $\gamma\delta$ cells, it is this subset that appears to be responsible for inducing germinal centers (14). Human $\gamma\delta$ T cells have also been found in germinal centers; these cells were found to up-regulate B cell costimulatory molecules such as CD40L, OX40, CD70, and inducible costimulatory molecule (ICOS) in response to TCR stimulation (15, 16). Together, these data indicate that $\gamma\delta$ T cells can promote, either directly or indirectly, the humoral immune response.

Here we show that, in the absence of Itk, $\gamma\delta$ T cell differentiation and effector function are dramatically altered. *Itk*^{-/-} mice contain increased numbers of CD4⁺ and NK1.1⁺ $\gamma\delta$ T cells that normally constitute the $\gamma\delta$ NKT population (17). These cells express PLZF, a transcription factor that uniquely defines $\alpha\beta$ NKT cells and is essential for normal $\alpha\beta$ NKT cell development (18, 19). As a consequence, *Itk*^{-/-} $\gamma\delta$ T cells produce robust amounts of Th2 cytokines when stimulated, accompanied by enhanced expression of cell surface receptors associated with B cell help, such as ICOS and CD40L and thus promote a spontaneous elevation in serum IgE levels. These data indicate a surprising role for Itk in regulating the lineage development and effector function of $\gamma\delta$ T cells, particularly in controlling the PLZF⁺ subset.

Results

$\gamma\delta$ T Cells Promote the Hyper IgE and Enriched Germinal Center Phenotype Seen in *Itk*^{-/-} Mice. In an effort to identify the cell type producing Th2 cytokines and driving IgE class-switching and secretion in unimmunized *Itk*^{-/-} mice, we considered $\gamma\delta$ T cells. To test this possibility, *Itk*^{-/-} mice were crossed to *Tcrd*^{-/-} mice (20) that lack $\gamma\delta$ T cells. As shown in Fig. 1A and reported previously (3, 4), *Itk*^{-/-} mice have elevated concentrations of serum IgE compared with WT controls. Strikingly, in *Itk*^{-/-}*Tcrd*^{-/-} double-deficient mice, serum IgE levels drop significantly compared with *Itk*^{-/-} mice. Similar results were seen upon analysis of the proportion of germinal center phenotype B cells (B220⁺ peanut agglutinin [PNA]⁺) (Fig. 1B). Individual cohorts of mice were tested at 2 months of age, 3.5 months of age, and 5 months of age, with similar results.

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The authors declare no conflict of interest.

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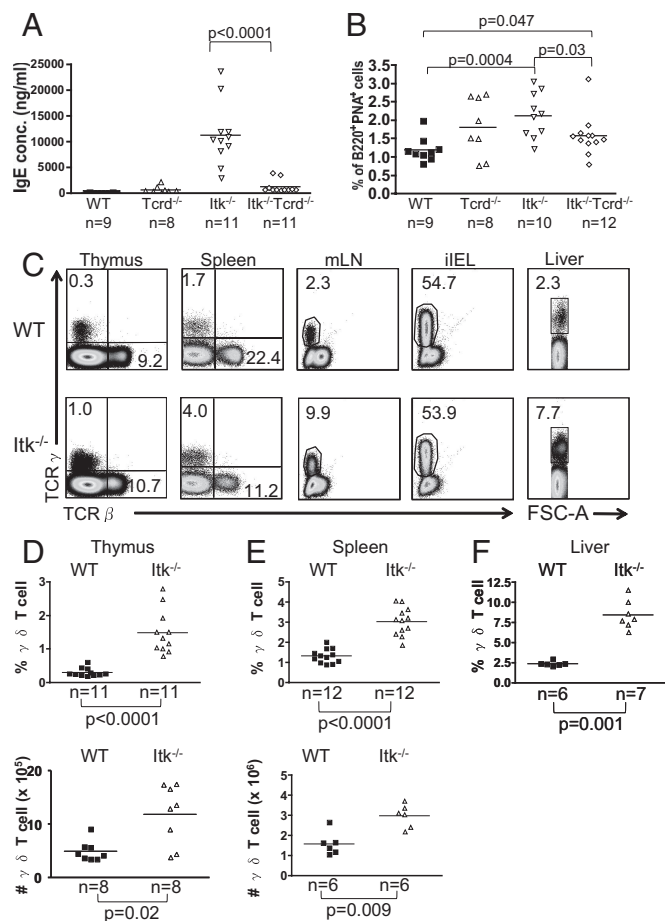


Fig. 1. $\gamma\delta$ T cells in *Itk*^{-/-} mice are responsible for the spontaneous elevation in serum IgE levels. (A) Serum obtained from WT, *Tcrd*^{-/-}, *Itk*^{-/-}, and *Itk*^{-/-}*Tcrd*^{-/-} mice were analyzed for IgE by ELISA. (B) Splenocytes from WT, *Tcrd*^{-/-}, *Itk*^{-/-}, and *Itk*^{-/-}*Tcrd*^{-/-} mice were stained with α -B220 Ab and PNA to identify germinal center B cells. Differences between WT and *Tcrd*^{-/-} ($P = 0.14$) and between *Itk*^{-/-} and *Itk*^{-/-}*Tcrd*^{-/-} ($P = 0.54$) were not statistically significant. (C) Cells were prepared from thymus, spleen, mesenteric lymph nodes (mLN), intestinal epithelium (iIELs), and liver and were stained with anti-TCR δ and anti-TCR β Abs. (D–F) Percentages and absolute numbers of $\gamma\delta$ T cells were compiled for the thymus (D) and spleen (E); percentages only were compiled for liver (F).

Itk-Deficient Mice Have Increased Numbers of $\gamma\delta$ T Cells. The finding that elevated serum IgE levels in *Itk*^{-/-} mice were dependent on $\gamma\delta$ T cells suggested that the $\gamma\delta$ T cells present in *Itk*^{-/-} mice might be altered relative to their WT counterparts. We first examined $\gamma\delta$ T cell numbers. As seen in Fig. 1C, *Itk*^{-/-} mice have increased proportions of $\gamma\delta$ T cells in the thymus, the spleen, the mesenteric lymph nodes, and the liver, but not in the intestinal intraepithelial lymphocyte (iIEL) compartment. A summary of data indicates significant increases in the proportion and absolute numbers of $\gamma\delta$ T cells in both the thymus and the spleen, as well as an increase in the percentage of $\gamma\delta$ T cells in the liver, in the absence of *Itk* (Fig. 1D–F).

Itk^{-/-} Mice Contain Increased Proportions of $\gamma\delta$ T Cell Subsets Expressing V γ 1.1+V δ 6.3⁺, CD4, and NK1.1. Analysis of TCR γ chain repertoires indicated an increased proportion in V γ 1.1⁺ $\gamma\delta$ T cells in *Itk*-deficient versus WT mice, and a concomitant decrease in the other major subsets (V γ 2⁺ and V γ 5⁺) (Fig. 2A and B). Further, the majority of V γ 1.1⁺ $\gamma\delta$ T cells in the thymus and liver of *Itk*^{-/-} mice also expressed V δ 6.3 (Fig. 2C). Overall, we find a significant increase in the absolute numbers of this $\gamma\delta$ T

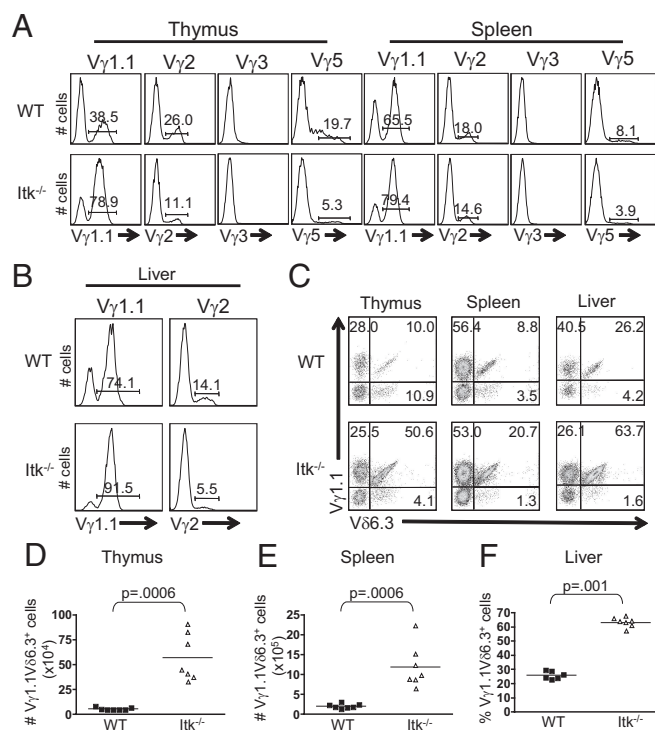


Fig. 2. *Itk*^{-/-} mice have an increased population of V γ 1.1V δ 6.3⁺ $\gamma\delta$ T cells. (A) Thymocytes and splenocytes were stained with antibodies to TCR δ and either V γ 1.1, V γ 2, V γ 3, and V γ 5. Histograms show V γ staining on gated TCR δ ⁺ cells. (B) Lymphocytes were isolated from the liver and stained with antibodies to TCR δ and V γ 1.1 or V γ 2. Histograms show V γ staining on gated TCR δ ⁺ cells. (C) Thymocytes, splenocytes, and liver cells were stained with antibodies to TCR δ , V γ 1.1, and V δ 6.3. Dot plots show V γ 1.1 vs. V δ 6.3 staining on gated TCR δ ⁺ T cells. (D) Absolute numbers of V γ 1.1V δ 6.3⁺ $\gamma\delta$ T cells in the thymus and spleen and percentages of V γ 1.1V δ 6.3⁺ $\gamma\delta$ T cells in the liver were compiled.

cell subset in thymus and spleen of *Itk*^{-/-} mice compared with controls (Fig. 2D and E), as well as an increased proportion of V γ 1.1+V δ 6.3⁺ $\gamma\delta$ T cells in *Itk*^{-/-} liver (Fig. 2F). Interestingly, V γ 1.1+V δ 6.3⁺ $\gamma\delta$ T cells are the predominant subset of $\gamma\delta$ NKT cells, and readily produce effector cytokines (17, 21, 22).

In contrast to WT $\gamma\delta$ T cells, the *Itk*^{-/-} $\gamma\delta$ T cell population shows a striking increase in the proportion of CD4⁺ cells in the thymus, spleen and liver (Fig. 3A and B). *Itk*^{-/-} mice also had elevated proportions of thymic NK1.1⁺ $\gamma\delta$ T cells (Fig. 3C and D). Interestingly, the majority of the thymic *Itk*^{-/-} CD4⁺ $\gamma\delta$ T cells and virtually all of the thymic *Itk*^{-/-} NK1.1⁺ $\gamma\delta$ T cells express high levels of the memory cell marker CD44 (Fig. 3E), as well as the activation marker CD69. As previously reported, the majority of $\gamma\delta$ T cells in the liver of WT mice are NK1.1⁺ (i.e., $\gamma\delta$ NKT cells) (17), and this is also the case in *Itk*^{-/-} liver (Fig. 3C and D).

Focusing on the V γ 1.1+V δ 6.3⁺ subset of $\gamma\delta$ T cells, we find that WT and *Itk*^{-/-} mice each have a similar proportion of NK1.1⁺ CD4⁻ cells, whereas NK1.1⁺ CD4⁺ cells are increased substantially in *Itk*^{-/-} mice [supporting information (SI) Fig. S1]. Furthermore, outside the thymus, the V γ 1.1+V δ 6.3⁺ subset is found predominantly in the liver, rather than the spleen (Fig. 3F and G), as has been reported previously for $\gamma\delta$ NKT cells (17). The $\gamma\delta$ T cells not expressing V γ 1.1V δ 6.3 correspond to the “conventional” NK1.1⁻CD4⁻CD8⁻ subset; this conventional $\gamma\delta$ T cell subset is not altered in numbers in *Itk*^{-/-} versus WT mice, indicating that ITK controls the generation of “nonconventional” CD4⁺ or NK1.1⁺, but not conventional, $\gamma\delta$ T cells.

Expanded populations of CD4⁺ $\gamma\delta$ T cells have been previ-

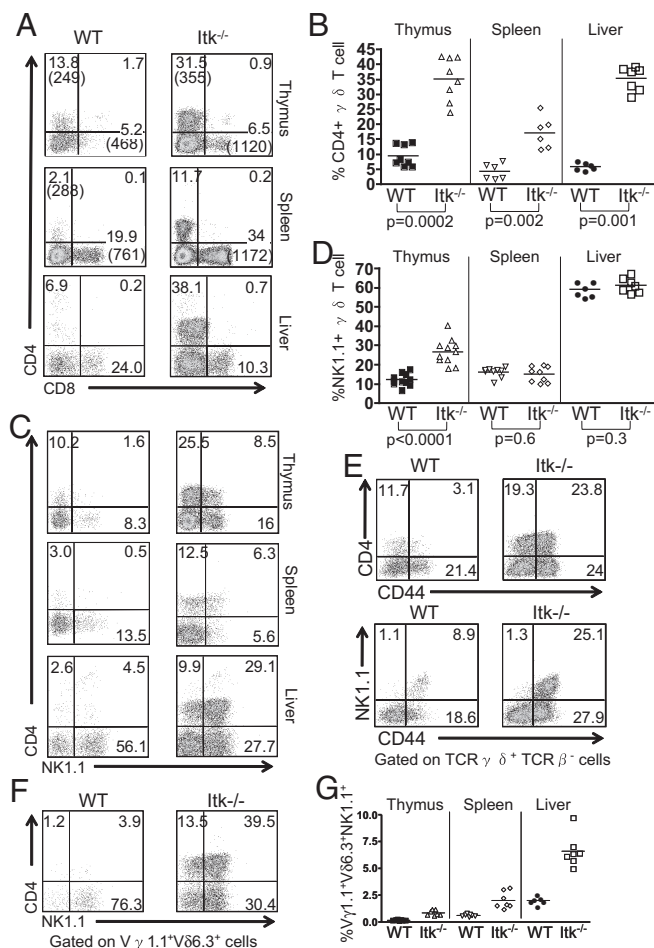


Fig. 3. Altered $\gamma\delta$ T cell subsets in the spleen and thymus of *Itk*^{-/-} mice. Cells were prepared from thymus, spleen, and liver of WT and *Itk*^{-/-} mice and analyzed by flow cytometry. (A) CD4 vs. CD8 expression on gated $\text{TCR}\delta^+\text{TCR}\beta^-$ cells. The percentages of each subpopulation are indicated, and the mean fluorescence intensities are shown in parentheses. (B) The percentages of $\text{CD4}^+\text{TCR}\delta^+$ cells in the thymus, spleen, and liver are shown. (C) CD4 vs. NK1.1 expression on gated $\text{TCR}\delta^+\text{TCR}\beta^-$ cells in the thymus, spleen, and liver. (D) The percentages of $\text{NK1.1}^+\text{TCR}\delta^+$ cells in the thymus, spleen, and liver. (E) Thymic $\text{TCR}\delta^+\text{TCR}\beta^-$ cells were analyzed for CD44 vs. CD4 (top) or NK1.1 (bottom) expression. Data are representative of two independent experiments. (F) $\text{TCR}\delta^+\text{TCR}\beta^-$ cells from the liver were analyzed for $\text{V}\gamma 1.1$, $\text{V}\delta 6.3$, NK1.1, and CD4 expression. Dot-plots show NK1.1 vs. CD4 expression on gated $\text{V}\gamma 1.1^+\text{V}\delta 6.3^+$ cells. (G) The percentages of total cells in the thymus, spleen, and liver that represent the $\text{V}\gamma 1.1^+\text{V}\delta 6.3^+\text{NK1.1}^+$ subset were calculated. Differences between the WT and *Itk*^{-/-} were statistically significant (thymus, $P = 0.0006$; spleen, $P = 0.0006$; liver, $P = 0.001$).

ously demonstrated in mice lacking expression of both $\text{TCR}\beta$ and CD5 (14). We therefore examined CD5 expression on *Itk*^{-/-} $\gamma\delta$ T cells, to determine if altered levels of CD5 might be contributing to the increased number of CD4^+ $\gamma\delta$ T cells in these mice, but detected no differences between WT and *Itk*^{-/-} mice. However, like a previously-reported subset of activated $\gamma\delta$ T cells specific for self-ligands, the MHC class Ib molecules, T10/T22 (23), the NK1.1^+ subset of thymic *Itk*^{-/-} $\gamma\delta$ T cells are all CD122-positive.

To determine whether the alterations we observed in *Itk*^{-/-} mice are intrinsic to the hematopoietic cells or to the environment, we generated bone marrow chimeras using WT or *Itk*^{-/-} bone marrow cells. These experiments demonstrated that the increased proportions of both $\gamma\delta$ T cells and germinal center B cells seen in *Itk*^{-/-} mice are intrinsic to *Itk*^{-/-} hematopoietic cells (Fig. S2). Consistent

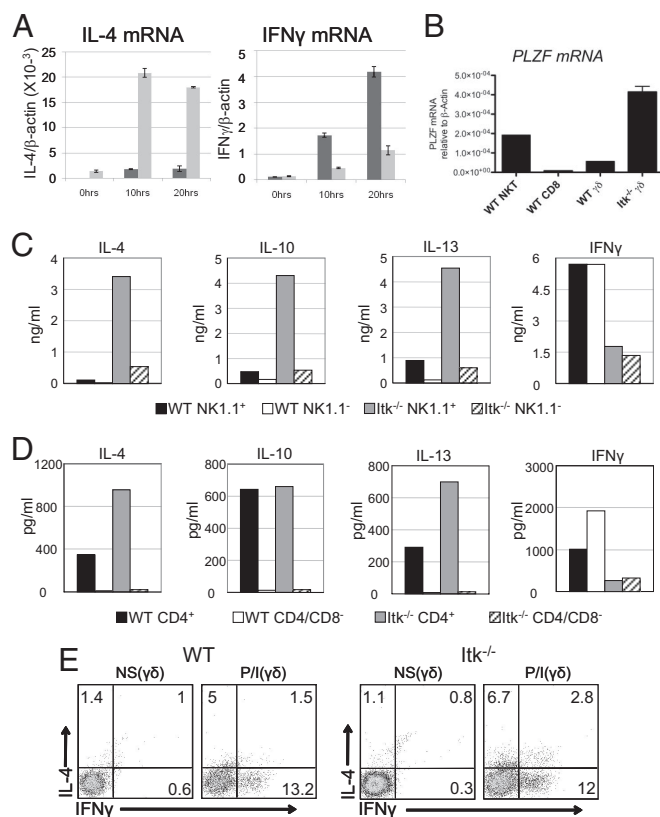


Fig. 4. *Itk*^{-/-} $\gamma\delta$ T cells produce IL-4 plus IFN- γ and express the transcription factor PLZF. Lymph nodes and spleens from WT and *Itk*^{-/-} mice were pooled, and total $\text{TCR}\delta^+$ cells (A, B) or sorted subpopulations (C, D) were analyzed. (A) 2×10^5 cells were stimulated with $10 \mu\text{g/ml}$ of anti- $\text{TCR}\delta$ for 0, 10, and 20 hours. IL-4 (left panel) and IFN- γ (right panel) mRNA expression levels normalized to β -actin were determined by real-time quantitative RT-PCR. Data shown are representative of two independent experiments. (B) Levels of PLZF mRNA normalized to β -actin were determined by real-time quantitative RT-PCR. WT peripheral $\text{CD8}^+\alpha\beta$ T cells and $\alpha\beta$ NKT cells were analyzed for comparison. Data are representative of two independent experiments. (C) 5×10^4 cells were stimulated with $10 \mu\text{g/ml}$ of anti- $\text{TCR}\delta$ for 72 hours and supernatants were analyzed for the presence of IL-4, IL-10, IL-13, and IFN- γ by cytometric bead array (CBA). Data are representative of three independent experiments. (D) 3×10^4 cells were stimulated as in (B). Supernatants were analyzed for the presence of IL-4, IL-10, IL-13, and IFN- γ by CBA. Data are representative of two independent experiments. (E) Nonstimulated (NS) and stimulated (P/I) WT (left) and *Itk*^{-/-} (right) $\gamma\delta$ T cells were analyzed for intracellular IL-4 and IFN- γ production. Cells were stimulated with 10 ng/ml PMA and $2 \mu\text{g/ml}$ Ionomycin (P/I) for 4 hours. Data are representative of four independent experiments.

with these data, we also find that the predominant Tec kinase expressed in WT $\gamma\delta$ T cells is *Itk* (Fig. S3).

Enhanced Expression of IL-4 and PLZF in *Itk*-Deficient $\gamma\delta$ T Cells. To determine the basis for the altered function of *Itk*^{-/-} $\gamma\delta$ T cells, we examined mRNA levels for T-bet, Eomesodermin, and GATA-3 in WT versus *Itk*^{-/-} $\gamma\delta$ T cells. We found that *Itk*^{-/-} $\gamma\delta$ T cells consistently expressed higher levels of GATA-3 mRNA and protein compared with WT $\gamma\delta$ T cells, while the NK1.1^+ subset of *Itk*^{-/-} $\gamma\delta$ T cells have reduced levels of mRNA for both T-bet and Eomesodermin (Fig. S3). These findings suggested that *Itk*^{-/-} $\gamma\delta$ T cells might produce a distinct pattern of cytokines compared with WT $\gamma\delta$ T cells. We first examined mRNA levels for the signature cytokines, IL-4 and IFN- γ , after $\gamma\delta$ T cell activation *in vitro*. As shown in Fig. 4A, in response to $\gamma\delta$ TCR stimulation *in vitro*, *Itk*^{-/-} $\gamma\delta$ T cells had constitutively elevated levels of IL-4 mRNA before stimulation and exhibited dramat-

ically enhanced induction of IL-4 mRNA at 10 hours and 20 hours poststimulation compared with WT $\gamma\delta$ T cells. Basal levels of IFN γ mRNA were similar between the WT and *Itk*^{-/-} $\gamma\delta$ T cells; after stimulation, both cell types produced IFN γ , although WT $\gamma\delta$ T cells showed higher levels of IFN γ mRNA compared with $\gamma\delta$ T cells lacking *Itk*.

Based on the high proportion of V γ 1.1⁺V δ 6.3⁺ $\gamma\delta$ T cells present in *Itk*^{-/-} mice and the previous association of this $\gamma\delta$ T cell subset with dual production of IL-4 and IFN γ (21, 22), we also examined WT and *Itk*^{-/-} $\gamma\delta$ T cells for expression of the transcription factor PLZF. PLZF has recently been found to be critical for the development and effector function of TCR $\alpha\beta$ ⁺ NKT cells, where it promotes the simultaneous production of IL-4 and IFN γ (18, 19). Interestingly, we found that splenic *Itk*^{-/-} $\gamma\delta$ T cells express substantially higher amounts of PLZF mRNA than do WT $\gamma\delta$ T cells (Fig. 4B). Furthermore, among *Itk*^{-/-} $\gamma\delta$ T cells, the NK1.1⁺ fraction expresses particularly high levels of PLZF mRNA. These findings support the conclusion that $\gamma\delta$ T cell development is altered in the absence of *Itk*.

To assess levels of cytokine protein secretion, individual subsets of *Itk*^{-/-} and WT $\gamma\delta$ T cells were purified and stimulated. As previous studies have found that NK1.1⁺ $\gamma\delta$ T cells and CD4⁺ $\gamma\delta$ T cells produce the highest levels of cytokines, particularly “Th2” cytokines (24–28), we compared NK1.1⁺ to NK1.1⁻, and CD4⁺ to CD4⁻ $\gamma\delta$ T cell subsets. After 72 hours of stimulation, WT NK1.1⁺ $\gamma\delta$ T cells secreted more IL-4, IL-10, and IL-13 than their NK1.1⁻ counterparts; in contrast, both NK1.1⁺ and NK1.1⁻ subsets of WT $\gamma\delta$ T cells secreted large amounts of IFN γ (Fig. 4C). Consistent with the analysis of cytokine mRNA levels, we observed elevated secretion of the Th2 cytokines IL-4, IL-10, and IL-13 by the *Itk*^{-/-} NK1.1⁺ $\gamma\delta$ T cells when compared with WT NK1.1⁺ cells; furthermore, *Itk*^{-/-} NK1.1⁺ $\gamma\delta$ T cells secreted much higher amounts of these cytokines per cell relative to the *Itk*^{-/-} NK1.1⁻ subset. In addition, we found that both subsets of *Itk*^{-/-} $\gamma\delta$ T cells secreted IFN γ , but at a lower level than the WT cells.

Comparison of the CD4⁺ versus CD4⁻ subsets of $\gamma\delta$ T cells confirmed previous data (27, 28) that $\gamma\delta$ T cells expressing CD4 are the major cytokine-producing population, particularly for the “Th2” cytokines. As shown in Fig. 4D, *Itk*^{-/-} CD4⁺ $\gamma\delta$ T cells produce elevated levels of IL-4 and IL-13 compared with WT CD4⁺ $\gamma\delta$ T cells, but produce similar levels of IL-10. None of these cytokines were detected in supernatants of stimulated CD4⁻ $\gamma\delta$ T cells from either *Itk*^{-/-} or WT mice. As noted above, WT $\gamma\delta$ T cells (both CD4⁺ and CD4⁻ subsets) secrete higher levels of IFN γ than *Itk*^{-/-} $\gamma\delta$ T cells.

On a population basis, *Itk*^{-/-} $\gamma\delta$ T cells secreted both IL-4 and IFN γ after stimulation. To determine whether individual cells were dual producers of both effector cytokines, we stimulated WT and *Itk*^{-/-} $\gamma\delta$ T cells *in vitro* with phorbol 12-myristate 13-acetate (PMA) and ionomycin and then examined IL-4 and IFN γ production by intracellular cytokine staining (Fig. 4E). As expected, a larger proportion of WT $\gamma\delta$ T cells produce IFN γ in response to stimulation compared with those that produce IL-4, and few cells produce both cytokines. In contrast, *Itk*^{-/-} $\gamma\delta$ T cells include a significantly larger population that produces IL-4 than is seen in the WT $\gamma\delta$ T cell subset (*Itk*^{-/-}, 6.9 ± 1.1; WT, 3.5 ± 0.8; *n* = 7; *P* = 0.04); additionally, a greater proportion of *Itk*^{-/-} $\gamma\delta$ T cells produces both IL-4 plus IFN γ compared with WT $\gamma\delta$ T cells (*Itk*^{-/-}, 3.9 ± 0.8; WT, 1.7 ± 0.3; *n* = 7; *P* = 0.02). Because this pharmacological stimulation bypasses the need for ITK in TCR-mediated signaling, these data indicate that a larger proportion of *Itk*^{-/-} $\gamma\delta$ T cells are programmed to produce IL-4, as well as IL-4 plus IFN γ , before their activation. This latter finding, together with the data demonstrating increased numbers of CD4⁺ and NK1.1⁺ $\gamma\delta$ T cells in the *Itk*^{-/-} mice, strongly suggests that $\gamma\delta$ T cell development is altered in the absence of

Itk leading to a striking increase in a PLZF-positive, IL-4-producing $\gamma\delta$ T cell population.

***Itk*^{-/-} $\gamma\delta$ T Cells Up-Regulate Surface Receptors That Promote B Cell Help.**

We next examined $\gamma\delta$ T cells for the expression of costimulatory molecules that provide B cell help, such as CD40L, CD70, OX40, and ICOS (15). For these experiments, WT and *Itk*^{-/-} thymocytes and splenocytes were evaluated directly *ex vivo* and, in addition, were cultured for 24 hours in the presence of stimulatory anti-TCR δ antibodies. Although analysis of splenic $\gamma\delta$ T cells from WT and *Itk*^{-/-} mice did not reveal any major changes in co-stimulatory markers, we did see a small increase in the proportion of constitutively ICOS-positive $\gamma\delta$ T cells in the spleens of *Itk*^{-/-} mice. Inasmuch as *Itk*^{-/-} mice have a two-fold increase in the absolute number of total $\gamma\delta$ T cells in the spleen compared with WT, this difference amounts to a \approx 10-fold increase in ICOS-positive splenic $\gamma\delta$ T cells, and thus could account for the enhanced B cell activation observed in *Itk*^{-/-} mice.

More strikingly, levels of ICOS were increased on a large proportion of the thymic *Itk*^{-/-} $\gamma\delta$ T cells compared with controls, but remained unaltered following stimulation (Fig. 5A). Evaluation of the ICOS^{hi} fraction of *Itk*^{-/-} thymic $\gamma\delta$ T cells indicated that nearly all of these cells were CD4⁺, and a substantial proportion were also NK1.1⁺ (Fig. 5B). Little to no difference was seen in the basal levels of CD40L and OX40 when comparing thymic *Itk*^{-/-} $\gamma\delta$ T cells to WT $\gamma\delta$ T cells directly *ex vivo* (Fig. 5A). However, after 24 hours of *in vitro* stimulation on anti-TCR δ -coated plates, *Itk*^{-/-} cells up-regulated both CD40L and OX40, whereas WT $\gamma\delta$ T cells did not. This up-regulation of CD40L and OX40 was detected on all subsets of *Itk*^{-/-} $\gamma\delta$ T cells (Fig. 5B). Finally, we could not detect expression of CD70 on either WT or *Itk*^{-/-} thymic $\gamma\delta$ T cells.

Discussion

Overall, our data indicate that $\gamma\delta$ T cell development is significantly altered in the absence of *Itk*, yielding increased numbers of $\gamma\delta$ T cells and a shift in the major effector functions mediated by these cells. Most strikingly, *Itk*^{-/-} mice have elevated numbers of $\gamma\delta$ T cells expressing CD4 and NK1.1. Furthermore, unlike the $\gamma\delta$ T cells in WT mice, the *Itk*^{-/-} $\gamma\delta$ T cells secrete large quantities of the Th2 cytokines, IL-4, IL-10, and IL13, in addition to the IFN γ typically secreted by activated WT $\gamma\delta$ T cells, correlating with high levels of the transcription factor, PLZF. These findings strongly suggest that *Itk* signaling plays a key role in regulating $\gamma\delta$ T cell lineage development.

Surprisingly, these altered $\gamma\delta$ T cells are responsible for promoting significant levels of spontaneous IgE secretion in *Itk*^{-/-} mice. Based on the findings presented here, it is likely that the high levels of IL-4 and IL-13 produced by the activated *Itk*^{-/-} $\gamma\delta$ T cells are a major factor in this response. Our data indicate that activated *Itk*^{-/-} $\gamma\delta$ T cells express elevated amounts of B cell co-stimulatory molecules, such as ICOS, CD40L, and OX40, further suggesting that the $\gamma\delta$ T cells may be directly providing help to the B cells, leading to B cell activation and Ig class switching.

In humans, a variety of studies have implicated $\gamma\delta$ T cells in allergic airway inflammation (29, 30) and, specifically, in promoting B cell activation and IgE class switching (31, 32). One interesting clinical report found that IL-4-producing $\gamma\delta$ T cells were likely responsible for a case of hyper IgE syndrome (33). Studies performed *in vitro* with human $\gamma\delta$ T cells showed that these cells, in combination with IL-4, can induce B cell activation, Ig isotype switching, and secretion of IgE (34). Further, these findings correlate well with observations that human $\gamma\delta$ T cells can express ICOS, CD40L, OX40, and CD70 (15). Our data indicate that the Tec family tyrosine kinase, *Itk*, plays a key role in regulating this potentially highly detrimental function of $\gamma\delta$ T cells.

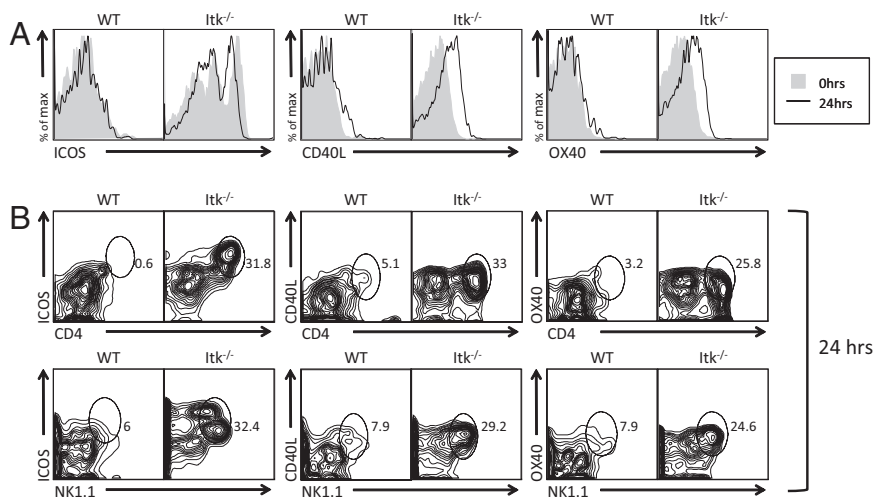


Fig. 5. $Itk^{-/-}$ $\gamma\delta$ T cells up-regulate costimulatory molecules involved in B cell help. (A) Thymocytes from WT and $Itk^{-/-}$ mice were pooled and left nonstimulated (0 hours) or were stimulated with 10 $\mu\text{g/ml}$ α -TCR δ for 24 hours. Nonstimulated and stimulated TCR δ^{+} TCR β^{-} T cells were then analyzed for the expression of ICOS, CD40L, and OX40. (B) WT and $Itk^{-/-}$ thymocytes stimulated for 24 hours with 10 $\mu\text{g/ml}$ α -TCR δ antibodies were stained for CD4, ICOS, CD40L, OX40, and NK1.1 expression.

Recently, $\gamma\delta$ T cells have also been implicated in the elevated IgE concentrations seen in mice carrying mutations in additional T cell signaling proteins. For instance, in the absence of the E3-ubiquitin ligase, Itch, $\gamma\delta$ T cells secrete IL-4 and promote IgE production in nonimmunized mice (29). More strikingly, mice expressing a mutant allele of the adapter protein linker of activated T cells (LAT), which lacks the three c-terminal tyrosines, succumb to a fatal lymphoproliferative disorder that is mediated by $\gamma\delta$ T cells (35). In these LAT mutant mice, the $\gamma\delta$ T cells accumulate to large numbers, and show a phenotype remarkably similar to those lacking Itk. For instance, the LAT mutant $\gamma\delta$ T cells secrete IL-4, rather than IFN- γ , and many of them express the CD4 co-receptor; in addition, the mice also have elevated levels of serum IgE. As this mutant LAT protein does not support $\alpha\beta$ T cell development, these altered $\gamma\delta$ T cells are the only source of T cell help for B cell activation and IgE class switching. As Itk and LAT interact in a TCR-dependent signaling complex in $\alpha\beta$ T cells, the similarities in the $\gamma\delta$ T cell phenotype in these two lines of mice strongly suggest that these proteins are also in the same signaling pathway downstream of the $\gamma\delta$ TCR, and furthermore, that this pathway regulates the development and effector function of $\gamma\delta$ T cells. As Itk has previously been shown to suppress the development of innate $\alpha\beta$ lineage T cells and to promote the development of conventional $\alpha\beta$ T cells (36), a similar function for Itk may be required in $\gamma\delta$ T cells; thus, in the absence of Itk, enhanced development of innate (e.g., PLZF $^{+}$, NK1.1 $^{+}$) $\gamma\delta$ T cells occurs, leading to increased numbers of effector cytokine-producing $\gamma\delta$ T cells in $Itk^{-/-}$ mice.

Interestingly, a recent report by Jensen et al. demonstrates that $\gamma\delta$ T cells, like $\alpha\beta$ T cells, are found as both naïve and effector subsets (23). Effector-type $\gamma\delta$ T cells express higher levels of CD44, NK1.1, and CD122 relative to the naïve subset and in addition, show an altered cytokine secretion profile. Furthermore, the presence of the effector $\gamma\delta$ T cell population correlated with the expression of the TCR ligand for these $\gamma\delta$ T cells, indicating that ligand recognition was responsible for their activated phenotype. As a large population of $Itk^{-/-}$ $\gamma\delta$ T cells exhibit a similar effector cell phenotype and produce effector cytokines such as IFN- γ and IL-4; these findings suggest that ligand recognition by $Itk^{-/-}$ $\gamma\delta$ T cells may contribute to their activation *in vivo* and their role in promoting IgE production in nonimmunized mice.

These findings have substantial relevance to the potential effects of small molecule inhibitors of Itk. Given the importance of Itk in Th2 development and cytokine production by CD4 $^{+}$ $\alpha\beta$ T cells, this kinase is currently being targeted for the development of drugs to treat asthma and other allergic diseases (5, 37). It would be unfortunate if Itk inhibition also led to the aberrant activation of $\gamma\delta$ T cells and thus to enhanced production of IgE. Elevated levels of serum IgE would, in turn, lead to up-regulation of the Fc ϵ RI on mast cells (38), promoting hyperresponsiveness of these cells to IgE-mediated receptor aggregation. In light of the findings presented here, further studies on the role of Itk in human $\gamma\delta$ T cells are clearly warranted.

Materials and Methods

Mice. $Itk^{-/-}$ mice (39) are on the C57BL/10 strain. Tcrd $^{-/-}$ mice (20) on the B57BL/6 background (Jackson Laboratories) were crossed to $Itk^{-/-}$ mice to obtain $Itk^{-/-}$ Tcrd $^{-/-}$ mice. Wild-type mice were $Itk^{+/+}$ Tcrd $^{+/+}$ littermates or C57BL/10 mice (Jackson Laboratories). All mice used were 2–3 months of age and were maintained at the University of Massachusetts Medical School under specific pathogen-free conditions after institutional animal care and use committee approval.

Cell Preparations, Antibodies, and Flow Cytometry. Liver lymphocytes were isolated by collagenase digestion of minced liver followed by Ficoll gradient centrifugation; iIELs were isolated by incubation of cleaned intestine followed by Ficoll gradient centrifugation. The following antibodies were purchased from BD Pharmingen: TCR δ (GL3)-FITC, V γ 2-FITC, V γ 3-FITC, V δ 6.2/6.3-PE, TCR β -allophycocyanin, TCR β -PE, TCR β -biotin (bio), CD4-allophycocyanin, CD4-PE, CD8 α -PerCP-Cy5.5, NK1.1-PE-Cy7, CD5-CyChrome, IL-4-PE, IFN- γ -allophycocyanin, B220-allophycocyanin, streptavidin (strep)-allophycocyanin, and OX40-biotin. TCR δ -allophycocyanin, ICOS-PE, and CD40L-allophycocyanin were purchased from eBioscience. PNA-FITC was purchased from Vector Laboratories. Strep-Cascade Yellow was purchased from Invitrogen Molecular Probes. V γ 1.1-bio was a kind gift from Lynn Puddington (University of Connecticut Health Center, Storrs, CT). Cells (500,000–2,000,000 events) were collected on a LSRII (BD Biosciences) flow cytometer. Data were analyzed using FlowJo software (Tree Star).

Quantitative Real-Time PCR. RNA and cDNA were prepared from sorted cells as previously described (40). Real-time PCR was also performed as previously described (6) on an i-Cycler (Bio-Rad). A cDNA clone encoding PLZF was a kind gift from Albert Bendelac (University of Chicago, Chicago).

In Vitro Stimulation Assays. Wild-type and $Itk^{-/-}$ TCR γ^{+} NK1.1 $^{+}$, TCR γ^{+} NK1.1 $^{-}$, TCR γ^{+} CD4 $^{+}$, and TCR γ^{+} CD4 $^{-}$ subsets from were activated *in vitro* with 10 $\mu\text{g/ml}$

of anti-TCR γ biotin (BD Pharmingen) for 72 hours, and supernatants were collected and IL-4, IL-10, IL-13, and IFN γ were measured by cytometric bead array (CBA) (BD Pharmingen). Cells used for quantitative PCR were stimulated for 10 and 20 hours and examined for IL-4 and IFN γ mRNA. For intracellular cytokine staining, cells from WT and *Itk*^{-/-} mice were stimulated as previously described (6). Cells were then surface stained for anti-TCR γ and anti-NK1.1, fixed, and permeabilized using a Cytofix/Cytoperm kit (BD Pharmingen) and stained for IL-4 and IFN γ . For ICOS, CD40L and OX40 expression on $\gamma\delta$ T cells, cells were stimulated for 24 hours with 10 μ g/ml anti-TCR γ . Cells were then surface stained with anti-ICOS, anti-CD40L, and anti-OX40 antibodies.

Serum Analysis. Blood was collected from WT, *Tcrd*^{-/-}, *Itk*^{-/-}, and *Itk*^{-/-}*Tcrd*^{-/-} mice. Serum was obtained by spinning the blood at 5000 rpm for 5 minutes and

removing the supernatant. Supernatants were analyzed by enzyme-linked immunosorbent assay (ELISA) for IgE.

Statistical Analysis. InStat software (GraphPad) was used to perform two-tailed nonparametric Mann-Whitney tests.

For additional details, see *SI Materials and Methods*.

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