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Beyond $\alpha\beta/\gamma\delta$ lineage commitment: TCR signal strength regulates $\gamma\delta$ T cell maturation and effector fate

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Abstract

Signaling by the $\gamma\delta$ T cell receptor (TCR) is required not only for $\alpha\beta/\gamma\delta$ lineage commitment but also to activate and elicit effector functions in mature $\gamma\delta$ T cells. Notably, at both of these stages, the signal delivered by the $\gamma\delta$ TCR is more robust than the one delivered by either the preTCR or the $\alpha\beta$ TCR. Recent studies now provide evidence that signaling by the $\gamma\delta$ TCR is also required at other stages during $\gamma\delta$ T cell development. Remarkably, the strength of the $\gamma\delta$ TCR signal also plays a role at these other stages, as evidenced by the findings that genetic manipulation of $\gamma\delta$ TCR signal strength affects $\gamma\delta$ T cell maturation and effector fate. In this review, we discuss how a strong TCR signal is a recurring theme in $\gamma\delta$ T cell development and activation.

Keywords

T cell development; Signal transduction; $\gamma\delta$ T cell receptor; Effector function

1. Enhanced signaling proficiency of the γδ T cell receptor

There are two T cell lineages, $\alpha\beta$ and $\gamma\delta$, that are defined by the antigen-binding heterodimers contained within their respective antigen receptors. While it is still not known why two T cell lineages have been conserved in all jawed vertebrates, years of study demonstrate that $\gamma\delta$ T cells recognize antigens differently than $\alpha\beta$ T cells, acquire effector functions faster than $\alpha\beta$ T cells, and play specialized roles in immunity (reviewed in Refs. [1–6]). However, despite this knowledge, the signaling properties of the $\gamma\delta$ T cell receptor (TCR) remain poorly understood.

To learn more about $\gamma\delta$ TCR signal transduction, we directly compared the signaling ability of the $\gamma\delta$ TCR with that of the more extensively studied $\alpha\beta$ TCR [7]. Because there are no known antigens for murine $\alpha\beta$ - and $\gamma\delta$ TCRs with similar binding kinetics [8], we chose to crosslink the respective TCRs with the same anti-CD3 ϵ monoclonal antibody (mAb) to initiate the TCR signaling cascade. When signal transduction by the two TCR isoforms was compared, the magnitude of the $\gamma\delta$ TCR signal was found to be greater than that of the $\alpha\beta$ TCR in assays that measure the mobilization of calcium and activation of the mitogen-

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activated protein kinase (MAPK) ERK (extracellular signal-regulated kinase) [7]. Importantly, the enhanced signaling proficiency of the $\gamma\delta$ TCR affected the kinetics of T cell activation, as the proliferative response of stimulated $\gamma\delta$ T cells was greater than that of stimulated $\alpha\beta$ T cells, regardless of the concentration of anti-CD3 mAb used [7]. This difference in proliferative response was due to the dependence of $\alpha\beta$ T cells but not $\gamma\delta$ T cells on CD28 costimulated by anti-CD3 ϵ and anti-CD28 mAbs was comparable to that of $\gamma\delta$ T cells stimulated by anti-CD3 ϵ mAb alone [7]. As would be expected, the discovery of the enhanced signaling proficiency of the $\gamma\delta$ TCR had a significant impact not only on our perception of $\gamma\delta$ T cells but also on our present understanding of T cell development and function.

2. $\gamma\delta TCR$ signal strength and the $\alpha\beta/\gamma\delta$ lineage fate decision

Both $\alpha\beta$ and $\gamma\delta$ T cells arise from a common thymic precursor but diverge into separate lineages early in ontogeny [9]. Once they diverge, $\alpha\beta$ and $\gamma\delta$ lineage cells follow different developmental pathways. Differentiation along the $\alpha\beta$ lineage begins at the immature CD4⁻ CD8⁻ (double negative; DN) stage. Following expression of, and signaling by, the pre-T cell receptor (preTCR), immature $\alpha\beta$ lineage cells undergo a strong proliferative burst, transition to the CD4⁺ CD8⁺ (double positive; DP) stage and initiate rearrangement at the *TCR*\alpha locus [10].DP thymocytes that are capable of expressing amature $\alpha\beta$ TCR then undergo a selection process based on the affinity of their TCRs for self-peptides/self-MHC and eventually emerge as functionally competent CD4⁺ or CD8⁺ single positive (SP) thymocytes [11,12]. Differentiation along the $\gamma\delta$ lineage also begins at the immature DN stage; however, it does not proceed through developmental stages defined by preTCR expression, CD4/CD8 coreceptor expression or extensive proliferation. Instead, $\gamma\delta$ lineage cells express only the mature $\gamma\delta$ TCR complex, remain DN, and undergo a small proliferative burst relative to $\alpha\beta$ lineage (*i.e.*, preTCR⁺) DN thymocytes [13–15].

Several models have been proposed to explain the mechanism(s) by which an immature DN thymocyte chooses to become either an $\alpha\beta$ or $\gamma\delta$ T cell (reviewed in Refs. [16–21]). Of these, the two which are currently in favor are the stochastic model and the signal strength model (Fig. 1). The stochastic model proposes that the fate of the cell is determined randomly prior to expression of a TCR and that the isoform of the TCR must match the predetermined fate of the cell in order for that cell to survive and mature (Fig. 1A). Evidence supporting this model is the heterogeneous expression of intracellular and surface proteins, which include transcription factors and cytokine receptors, in the progenitor pool that gives rise to $\alpha\beta$ and $\gamma\delta$ T cells [22,23]. This heterogeneity may reflect the activation of $\alpha\beta$ or $\gamma\delta$ lineage-specific molecular programs prior to the expression of a functional TCR isoform.

SOX13, which encodes a high-mobility-group transcription factor, is one of the genes that is differentially expressed in the progenitor pool (*i.e.*, approximately 50% of immature DN thymocytes express this gene) [23]. Notably, *SOX13* is also expressed in $\gamma\delta$ thymocytes but not in DP thymocytes, suggesting that *SOX13* expression in immature DN thymocytes marks cells committing to the $\gamma\delta$ lineage (Table 1) [23]. To determine whether Sox13 plays a role in the $\alpha\beta/\gamma\delta$ lineage decision, Melichar et al. [23] generated transgenic mice in which Sox13 is expressed in all immature DN thymocytes. In both fetal and adult Sox13 Tg mice, there was a striking decrease in the number of $\alpha\beta$ lineage cells (*i.e.*, DP thymocytes) but no concomitant increase in the number of $\gamma\delta$ thymocytes. Further analysis demonstrated that, in Sox13 Tg mice, the proliferative status of immature DN thymocytes from Sox13 Tg mice were found to express TCR γ transcripts, which are normally not detected in wild-type DP thymocytes [23]. Because of this latter finding, the authors concluded that Sox13 is able to

impose a $\gamma\delta$ lineage-specific molecular program when its expression is enforced in $\alpha\beta$ lineage cells [23]. However, since proliferation is required to silence transcription at the TCR γ locus in $\alpha\beta$ lineage cells as they transition from the DN to DP stage [24], an alternative explanation for this result, which is consistent with the thymic phenotype of Sox13 Tg mice, is that Sox13 regulates cellular proliferation, in that cells expressing no or low levels of Sox13 have a higher proliferative capacity than those expressing high levels of Sox13. Therefore, considering that $\alpha\beta$ lineage cells are more dependent than $\gamma\delta$ lineage cells on cellular proliferation for their development, it follows that overexpression of Sox13 would have a greater impact on their generation than on the generation of $\gamma\delta$ lineage cells. Interestingly, in the absence of Sox13, the converse phenotype is observed, namely a significant decrease in the number of $\gamma\delta$ thymocytes and no change in the number of DP thymocytes compared to wild-type mice [23]. However, it is important to note that because SOX13 is expressed in multiple tissues (reviewed in Ref. [25]) and because $Sox13^{-/-}$ mice die prematurely for an as yet unknown reason [23], it is not clear whether the defects in $\gamma\delta$ T cell development that are observed in $Sox13^{-/-}$ fetuses are in fact due to the loss of Sox13in $\gamma\delta$ T cell precursors. Thus, more studies are required to elucidate the functional significance of *SOX13* expression in both immature DN thymocytes and $\gamma\delta$ lineage cells.

The signal strength model differs from the stochastic model in that TCR signaling plays a primary (i.e., deterministic), rather than a secondary (i.e., confirmatory), role in the lineage decision process (Fig. 1B). Specifically, it posits that the strength of the signal delivered by the antigen receptor instructs lineage choice, with immature DN thymocytes receiving a strong signal choosing the $\gamma\delta$ T cell fate and those receiving a weak signal choosing the $\alpha\beta$ T cell fate [21]. Although the signal that directs lineage choice can potentially be delivered by any TCR isoform, under normal conditions, it is the $\gamma\delta$ TCR that transduces the strong signal and the preTCR that transduces the weak signal [26,27]. Genetic manipulation of the strength of the signal through a single TCR isoform, namely the $\gamma\delta$ TCR, provides strong evidence in support of this model. First, these studies demonstrated that the critical factor in dictating the $\alpha\beta/\gamma\delta$ lineage decision is the strength of the TCR signal strength, not the type of TCR expressed [26,27]. Moreover, the genetic manipulations affected lineage choice in a consistent manner, with attenuation of $\gamma\delta TCR$ signal strength favoring the $\alpha\beta$ fate and augmentation of $\gamma\delta$ TCR signal strength favoring the $\gamma\delta$ fate [26,27]. Recently, the extracellular signal-related kinase (ERK)-early growth response gene (Egr)-inhibitor of DNA binding 3 (Id3) or ERK-Egr-Id3 pathway was identified as a signaling pathway that is activated in $\gamma\delta$ lineage cells as a consequence of a strong TCR signal [27,28]. Accordingly, alterations in the expression levels of either EGR1 or ID3 in immature DN thymocytes had a significant effect on $\alpha\beta/\gamma\delta$ lineage commitment (Table 1) [27,28]. Despite the fact that these data support the signal strength model, it is also conceivable that TCR signal strength does not determine lineage fate but instead confirms the fate decision of pre-committed immature DN thymocytes. However, a recent study demonstrated that the progeny of a single thymocyte, expressing either the $\gamma\delta TCR$ or the preTCR and destined to adopt the $\alpha\beta$ lineage fate, can be redirected to the γδ lineage by a strong TCR signal, indicating that precommitment does not occur prior to TCR signaling [29]. These results suggest that, regardless of which molecules are expressed in immature thymocytes, or their potential to influence the decision process, the strength of the TCR has the final say.

3. γδTCR signal strength and γδ T cell maturation

The maturation stages of $\gamma\delta$ thymocytes are not as well-defined as those of $\alpha\beta$ thymocytes and, to date, very few surface antigens have been identified to define their maturation stages. $\gamma\delta$ TCR surface expression is first detected on DN thymocytes expressing the CD25 and CD24 surface antigens [15,26,30]. Subsequent signaling by the $\gamma\delta$ TCR is necessary to downregulate CD25 expression, as virtually all TCR $\gamma\delta^+$ thymocytes in mice bearing

mutations in critical components of the TCR-coupled signaling pathway retain CD25 expression [26,30]. Interestingly, not all $\gamma\delta$ thymocytes downregulate CD24 expression in the thymus. Although a small percentage of CD24^{lo} TCR $\gamma\delta^+$ cells can be detected in the thymus [15,31,32], the vast majority of $\gamma\delta$ thymocytes that are exported to the periphery are CD24^{hi}, suggesting that these recent thymic emigrants undergo some of their maturation in the periphery [31,32].

Notably, decreasing, not increasing, $\gamma\delta$ TCR signal strength has an effect on $\gamma\delta$ T cell maturation [27,33]. Lck is a positive regulator of both $\alpha\beta$ - and $\gamma\delta$ TCR signaling [33–36]; therefore, $\gamma\delta$ TCR signal strength can be attenuated by reducing or eliminating the expression of Lck. When $\gamma\delta$ T cell maturation was examined in $\gamma\delta$ TCR Tg Lck^{+/-} or Lck^{-/-} mice, significant decreases in the numbers of both CD24^{lo} $\gamma\delta$ thymocytes [27] and peripheral CD24^{hi} $\gamma\delta$ T cells [33] were observed. Interestingly, augmenting $\gamma\delta$ TCR signal strength, by reducing the expression levels of the negative regulator Fyn [33,37], had no effect on $\gamma\delta$ T cell maturation as evidenced by the wild-type numbers of CD24^{hi} $\gamma\delta$ T cells in the periphery of $\gamma\delta$ TCR Tg Fyn^{+/-} mice [33]. Together, these data indicate that a relatively strong $\gamma\delta$ TCR signal is required following $\gamma\delta$ T cell commitment for the maturation, survival and/or export of thymic $\gamma\delta$ T cells.

γδTCR signal strength and acquisition of effector fates

Recent studies have demonstrated that $\gamma\delta$ T cells have the potential to adopt multiple effector fates and, importantly, that these effector fates are predetermined in the thymus [38–40]. For at least some of these effector fates, there is evidence that interactions between the $\gamma\delta$ TCR and endogenous self-ligand are required for their fate selection [38,40]. Accordingly, because of this dependence on ligand-induced signaling, these effector fate decisions are sensitive to alterations in $\gamma\delta$ TCR signal strength. However, as ligand may play a role in the $\alpha\beta/\gamma\delta$ lineage decision [27], it is not clear whether these effector fate decisions occur concurrently with or subsequent to the $\gamma\delta$ lineage fate decision (Fig. 2).

The V γ 1⁺ V δ 6.3/6.4⁺ $\gamma\delta$ T cell subset shares properties with NKT cells, in that they express promyelocytic leukemia zinc finger protein or PLZF, a transcription factor required for the development of functional NKT cells, and they produce IL-4 and/or IFNy following TCR engagement [40–44]. The restricted expression of PLZF to a $\gamma\delta$ T cell subset with limited V γ and V δ usage suggests that TCR specificity plays a critical role in the development of these NKT-like cells [39]. Consistent with this idea is the finding that the expression of a $V\gamma 1/$ V δ 6.4 $\gamma\delta$ TCR transgene supports the generation of PLZF⁺ $\gamma\delta$ T cells [39]. Remarkably, PLZF expression can also be induced in immature $\gamma\delta$ thymocytes with a diverse repertoire (*i.e.*, extensive V γ and V δ usage) following TCR cross-linking [40]. Together, these results demonstrate that a strong TCR signal, delivered either by the interaction of the V γ 1⁺ V δ 6.3/6.4⁺ $\gamma\delta$ TCR with self-ligands or by treatment with anti-TCR $\gamma\delta$ or specific anti-V γ mAbs, is required for the development of this $\gamma\delta$ T cell subset [40]. Paradoxically, when TCR signal strength is attenuated, such as in mice deficient for the Tec kinase Itk or in mice expressing a signaling mutant form of LAT or SLP-76, the development of $V\gamma 1^+ V\delta 6.3/6.4^+$ PLZF⁺ $\gamma\delta$ T cells is favored [30,45–47]. One possible explanation for these data is that the signal generated by the interaction between the V γ 1⁺ V δ 6.3/6.4⁺ $\gamma\delta$ TCR and self-ligands is strong enough to surmount an attenuated TCR signal and to promote the development of this $\gamma\delta$ T cell subset. It would be interesting to examine the development of PLZF⁺ $\gamma\delta$ T cells in mice in which $\gamma\delta$ TCR signal strength has been augmented to determine whether the generation of V γ 1⁺ V δ 6.3/6.4⁺ PLZF⁺ $\gamma\delta$ T cells is affected and whether PLZF expression can be induced in mature $\gamma\delta$ thymocytes bearing V γ and V δ gene segments other than V $\gamma1$ and V\delta6.3/6.4.

A second effector response of $\gamma\delta$ T cells is the production of IFN γ . IFN γ -producing effectors are defined by the expression of the tumor necrosis factor receptor member CD27, with a subset of these CD27⁺ $\gamma\delta$ T cells expressing CD122 [38,39]. CD122⁺ CD27⁺ $\gamma\delta$ T cells are generated in the thymus through interactions with self-ligands [38]. When stimulated by TCR cross-linking, these CD122⁺ CD27⁺ $\gamma\delta$ T cells are capable of rapidly producing IFN γ [33,38]. As encounter with self-antigen is required for their development, alterations in $\gamma\delta$ TCR signal strength have been shown to affect the generation of CD122⁺ CD27⁺ $\gamma\delta$ T cells. Significantly, both weakening and strengthening the $\gamma\delta$ TCR signal, by reducing the cellular levels of Lck and Fyn, respectively, resulted in fewer peripheral CD122⁺ CD27⁺ $\gamma\delta$ T cells compared to wild-type mice [33]. These results suggest that selection of CD122⁺ CD27⁺ $\gamma\delta$ T cells, at least for those bearing the V $\gamma\delta$ /V δ 1 $\gamma\delta$ TCR transgene, occurs over a relatively narrow signaling range.

CD122⁻ CD27⁺ $\gamma\delta$ T cells, which also have the potential to be IFN γ -producers, differ from CD122⁺ CD27⁺ $\gamma\delta$ T cells in many ways. First, CD122⁻ CD27⁺ $\gamma\delta$ T cells require several days following TCR engagement to differentiate into IFN γ -producing effectors [39]. Second, although they also arise in the thymus, there is no evidence that CD122⁻ CD27⁺ $\gamma\delta$ T cells are selected by interactions with self-ligands [39]. However, there is evidence that their development is dependent on interactions with a quorum of DP thymocytes capable of producing cytokines such as lymphotoxin [39,48]. As would be predicted, alterations in $\gamma\delta$ TCR signal strength have no apparent effect on the generation of CD122⁻ CD27⁺ $\gamma\delta$ T cells in wild-type, Lck^{+/-} and Fyn^{+/-} mice [32]. These results indicate that the selection of CD122⁻ CD27⁺ $\gamma\delta$ T cells, compared to that of CD122⁺ CD27⁺ $\gamma\delta$ T cells, is less dependent on TCR signaling.

The third known effector fate of $\gamma\delta$ T cells is to produce IL-17. $\gamma\delta$ T cells destined to be IL-17 producers express the IL-23 receptor [49–51] but not CD122 and CD27 [38,39]. Current data indicate that encounter with self-antigens in the thymus is not required for the generation of this $\gamma\delta$ T cell effector subset [38]. Accordingly, when $\gamma\delta$ TCR signal strength is either attenuated by reducing Lck levels or augmented by reducing Fyn levels, the expression of *IL23R* and *IL12RB1* (subunits of the IL-23 receptor [52]) is unaffected [33]. A recent report demonstrates that the acquisition of IL-17-producing ability requires TGF- β signaling, as mice deficient in TGF β 1 and Smad3, an intermediate in the TGF- β signaling pathway, have a selective impairment in the generation of IL-17-producing $\gamma\delta$ T cells [53]. Thus, these results suggest that signals other than those delivered through the $\gamma\delta$ TCR play a role in shaping the $\gamma\delta$ effector repertoire.

5. Concluding remarks

Data is accumulating to support a model in which TCR signal strength has a critical and deterministic role at multiple points in $\gamma\delta$ T cell development, affecting lineage choice, maturation and acquisition of effector functions. An intriguing concept emerging from these findings is that TCR signal strength may be utilized to achieve entirely different outcomes in $\alpha\beta$ and $\gamma\delta$ lineage cells. In the $\alpha\beta$ lineage, it is well established that the strength/duration of signal regulates the outcome of thymocyte selection, with relatively weak signals promoting positive selection. In striking contrast, strong or sustained TCR signals appear to be required for efficient $\gamma\delta$ lineage commitment and maturation. It is tempting to speculate that these differential signaling requirements may reflect a hierarchical and functional relationship between $\alpha\beta$ and $\gamma\delta$ T cells. $\gamma\delta$ T cells recognize unprocessed antigen, are not necessarily dependent on costimulation by APCs, and appear to require strong interaction with self-ligands for their maturation. These properties may be optimized for the

development of rapidly responding T cell populations with a limited TCR repertoire, which represent a first line of defense against common insults. In contrast, $\alpha\beta$ T cells are self-MHC restricted, require weak interactions with self-ligand for development (and in fact are deleted by strong interactions) and are strictly dependent upon costimulation for their full activation. This latter system favors the formation of a highly diverse TCR repertoire that requires appropriate antigen processing and presentation as well as costimulation for full activation in order to prevent autoimmunity. Further investigation into the role of TCR signaling potential in the $\gamma\delta$ T cell developmental program should provide further insights into the shared and unique aspects of $\alpha\beta$ and $\gamma\delta$ T cell maturation.

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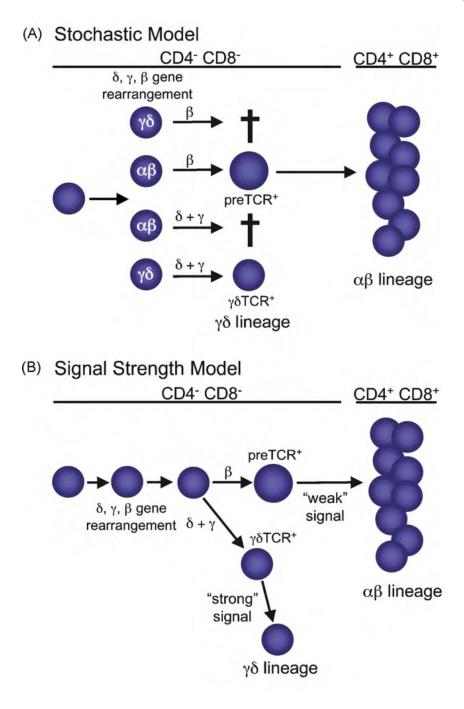


Fig. 1.

Current models of $\alpha\beta/\gamma\delta$ lineage commitment. (A) Stochastic model: in this model, the lineage fate decision occurs randomly prior to the expression of either the preTCR or the $\gamma\delta$ TCR. Nonetheless, the expressed TCR isoform must match the predetermined fate of the immature thymocyte in order for that cell to mature. If the isoform does not match the predetermined fate, then the immature thymocyte undergoes apoptosis. (B) Signal strength model: in this model, the strength of the TCR signal dictates the fate of the cell, with cells receiving a strong signal choosing the $\gamma\delta$ T cell fate and cells receiving a weak signal choosing the $\alpha\beta$ T cell fate. In a wild-type thymus, it is usually the $\gamma\delta$ TCR that delivers a strong TCR signal and the preTCR that delivers a weak TCR signal.

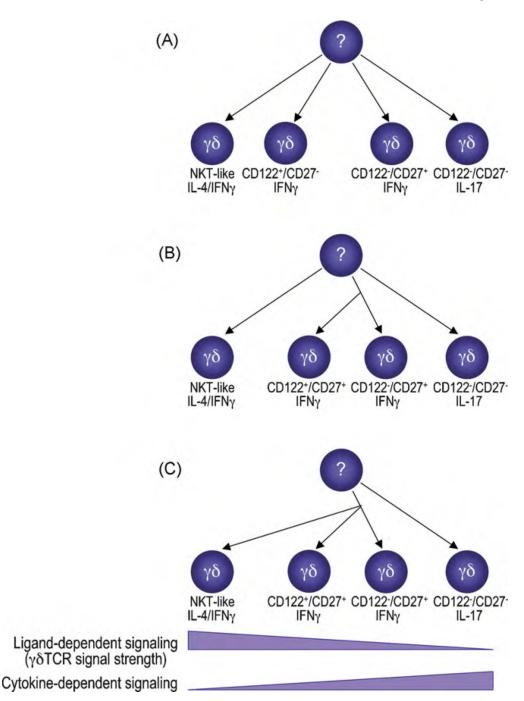


Fig. 2.

Proposed scenarios by which thymic $\gamma\delta$ T cells acquire effector functions. $\gamma\delta$ T cells acquire the potential to differentiate into effectors, which are able to produce IL-17, IFN γ or both IFN γ and IL-4, in the thymus. As it is not known whether the acquisition of effector fate occurs concurrently with or subsequent to commitment to the $\gamma\delta$ lineage, we have designated the cell that has the potential to give rise to these different effectors by a "?". Moreover, based on current data, we have ordered the different effector fates relative to one another according to their dependence on ligand- and cytokine-induced signaling. (A) In the first scenario, all effector fates represent different lineages, with the resulting effector fate depending on the specificity of the $\gamma\delta$ TCR in addition to the availability of self-ligands and

various cytokines. (B) In the second scenario, $CD27^+ \gamma \delta$ thymocytes, which all have the potential to become IFN γ -producers, represent a single effector lineage that can give rise to $CD122^+ CD27^+ \gamma \delta$ thymocytes following encounter with self-antigen. (C) In the last scenario, we have grouped NKT-like, $CD122^+ CD27^+$ and $CD122^- CD27^+ \gamma \delta$ subsets into one lineage based on their potential to produce IFN γ . As NKT-like and $CD122^+ CD27^+ \gamma \delta$ subsets require interactions with self-antigen, we propose that it is the strength of the signal delivered by this interaction that dictates effector fate, with cells receiving the stronger signal becoming NKT-like $\gamma \delta$ T cells.

Table 1

Recently identified genes that play a role in the commitment and/or development of $\gamma\delta$ lineage cells.

Gene	Reference
SOX13	Melichar et al. [23]
EGR1	Haks et al. [27]
ID3	Lauritsen et al. [28]

SOX13, sex-determining region (Sry)-related high-mobility-group (HMG) box; EGR1, early growth response 1; ID3, inhibitor of DNA binding 3.