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TCR $\gamma\delta$ signalling in $\gamma\delta$ cell development; strength isn't everything

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Abstract

$\gamma\delta$ cells are conserved across ~450 million years of evolution, from which they share the distinction, alongside $\alpha\beta$ T cells and B cells, of forming antigen receptors by somatic gene recombination. However, much about these cells remains unclear. Indeed, although $\gamma\delta$ cells display “innate-like” characteristics exemplified by rapid tissue-localised responses to stress-associated stimuli, their huge potential for T cell receptor (TCR) $\gamma\delta$ diversity also suggests “adaptive-like” potential. Clarity requires a better understanding of TCR $\gamma\delta$ itself, not only through identification of TCR-ligands, but also by correlating thymic TCR $\gamma\delta$ signalling with commitment to $\gamma\delta$ effector fates. Here, we propose that thymic TCR $\gamma\delta$ -ligand engagement *versus* ligand-independent signalling differentially imprints innate-like *versus* adaptive-like characteristics on developing $\gamma\delta$ cells, which fundamentally dictate their peripheral effector properties.

Unresolved role for TCR $\gamma\delta$ in $\gamma\delta$ cell development

$\gamma\delta$ T cells are predominantly tissue-resident lymphocytes that display diverse responses against pathogens and tumours [1]. Indeed, novel immunotherapies that target $\gamma\delta$ cells are now being explored to combat chronic viral infections, atopic and autoimmune pathologies, and various cancers [2,3]. Stimulation through TCR $\gamma\delta$ is critical for $\gamma\delta$ cell function [4]. However, by contrast to TCR $\alpha\beta$, signalling through which is absolutely required for thymic $\alpha\beta$ T cell development, the role of TCR $\gamma\delta$ in $\gamma\delta$ cell development remains controversial. TCR $\gamma\delta$ signalling is clearly necessary for commitment to the $\gamma\delta$ lineage, but the initiation, regulation and molecular nature of this commitment signal are still uncertain. Moreover, ligand-mediated positive and negative selection through TCR $\gamma\delta$ remain poorly understood, as too is the correlation between thymic TCR $\gamma\delta$ signalling and subsequent $\gamma\delta$ effector fates. Here, we build on recent studies that assess the initiation and consequences of TCR $\gamma\delta$ signalling in immature thymocytes [5-7], to propose that thymic TCR $\gamma\delta$ -ligand engagement *versus* ligand-independent TCR $\gamma\delta$ signalling may differentially impose innate-like *versus* adaptive-like features on developing $\gamma\delta$ cells.

Heterogeneity of peripheral $\gamma\delta$ subsets

Discussion of the role of TCR $\gamma\delta$ in $\gamma\delta$ cell development first requires appreciation of the heterogeneous nature of peripheral and thymic $\gamma\delta$ subsets. Functionally distinct $\gamma\delta$ subsets have been extensively characterised by surface phenotype. For example, dendritic epidermal T cells (DETC), that reside in murine epidermis and predominantly express a TCR $\gamma\delta$ that uses

TCR γ variable-region-5 (V γ 5) and TCR δ variable-region-1 (V δ 1) (nomenclature from [8]), are CD44⁺CD62L⁻, express CD103 (α_E integrin), and are CD122⁺ consistent with their dependence on IL-15 (**Figure 1 and Table 1**) [9,10]. DETC also readily secrete IFN- γ when activated. This CD44⁺CD62L⁻CD122⁺ IFN- γ -secreting phenotype is also shared by a minor population of lymphoid $\gamma\delta$ cells (~0.5%) whose TCR binds to MHC class IB molecules T10^b and T22^b, but only in mouse strains expressing T10^b and T22^b [7]. Moreover, it also characterises a CD90^{dull}CD27⁺ “NKT-like” $\gamma\delta$ subset that uses a restricted V γ 1⁺V δ 6.3⁺ (or V δ 6.4⁺) TCR and is known to secrete both IFN γ and IL-4 [11].

By contrast to DETC, V γ 4-biased $\gamma\delta$ cells (i.e. a $\gamma\delta$ subset with over-representation of V γ 4-containing TCR γ chains) of the murine dermis secrete IL-17A, are CD44⁺CD122⁻, and express CCR6 and the scavenger receptor SCART2 [9,12,13] (**Figure 1 and Table 1**). These cells are likely CD27⁻, as they closely resemble IL-17A-secreting V γ 6-biased $\gamma\delta$ cells from the peritoneal cavity and female reproductive tract that are CD27⁻CD44⁺CD122⁻ and CD25⁺ [14,15]. A CD27⁻CD44⁺CD62L⁻CD122⁻ phenotype accompanied by CCR6 and SCART2 expression is also shared by a minor population of IL-17A-producing $\gamma\delta$ cells from the secondary lymphoid organs [13,16,17]. Nonetheless, the majority of lymphoid $\gamma\delta$ cells from naive mice secrete large amounts of IFN- γ when activated, but no IL-17A [16]. These cells are CD27⁺, with a contrasting (for example, to DETC) CD44⁻CD62L⁺CD122⁻ phenotype (**Figure 1 and Table 1**). Finally, V γ 7-biased $\gamma\delta$ intraepithelial lymphocytes (IELs) of the gut are CD27⁺ and express IFN- γ on activation [18]. These cells are often described as “partially activated”, and may be under constant stimulation from gut-associated antigens [19,20].

Thymic $\gamma\delta$ subsets

TCR $\gamma\delta^+$ thymocytes are first evident from embryonic day-14 (E14), being initially dominated by a population of V γ 5⁺V δ 1⁺ DETC progenitors [21]. Shortly after a population of V γ 6⁺V δ 1⁺ progenitors emerge that are destined for the female reproductive tract, peritoneal cavity and tongue. Thymic terminal transferase (TdT) is not expressed during these prenatal stages, resulting in simple V-D-J joins that characterize the canonical TCRs of fetal-derived $\gamma\delta$ cells [21]. By contrast, postnatal thymic precursors of gut and lymphoid $\gamma\delta$ cells possess diverse V-D-J joins in their V γ 1, 2, 4 and 7-containing TCRs. This sequential progression of $\gamma\delta$ cell output is partly due to ordered V γ -region transcription and rearrangement. Nonetheless, other intrinsic differences between fetal/adult thymic progenitors [22,23], and requirement for age-specific thymic stromal factors [24], are also thought to influence subsequent $\gamma\delta$ cell fate.

In the adult murine thymus, CD27, CD25, CD24 and CD44 identify five distinct $\gamma\delta$ cell populations [16,25] (**Figure 1 and Table 1**). The most immature TCR $\gamma\delta^+$ cells are CD27⁺CD25⁺CD24⁺CD44⁻, express low TCR levels but are highly proliferative [16,25]. These progenitors down-regulate CD25 to become CD27⁺CD25⁻CD24⁺CD44⁻ cells, that can possibly already colonise the periphery [26]. They also likely represent precursors for three “mature” $\gamma\delta$ thymocyte populations that lack surface expression of CD24. This includes a CD27⁻ subset that is CD44⁺CD62L⁻, largely CCR6⁺ [17], and is already committed to IL-17A secretion [16]. By contrast, mature CD27⁺CD24⁻ $\gamma\delta$ thymocytes, that have potential to secrete IFN- γ , can be further sub-divided into CD44⁺CD62L⁻CD122⁺ and CD44⁻CD62L⁺CD122⁻ subsets. The former lack CCR9 and are largely NK1.1⁺ [17,20], being enriched for precursors of NKT-like $\gamma\delta$ cells [11]. Conversely, CD27⁺CD24⁻CD44⁻CD62L⁺CD122⁻ $\gamma\delta$ thymocytes are likely progenitors to

those of similar phenotype in peripheral lymphoid organs (see previous section). Thus, the thymus generates distinct $\gamma\delta$ populations with clear phenotypic links to peripheral $\gamma\delta$ subsets.

Thymic commitment to a $\gamma\delta$ cell fate

$\gamma\delta$ and $\alpha\beta$ T cells share a common $CD4^-CD8^-$ double negative (DN) thymic progenitor in which $TCR\gamma$, $TCR\delta$, and $TCR\beta$ rearrangements initiate [21]. DN cells that express a preTCR ($TCR\beta$ paired with invariant preTCR α chain), traverse a “ β -selection” checkpoint to a $CD4^+CD8^+$ double positive (DP) stage that marks commitment to the $\alpha\beta$ lineage [27]. By contrast, $TCR\gamma\delta$ expression appears to commit DN cells to a $\gamma\delta$ fate. These observations initially suggested a qualitatively instructional role for preTCR and $TCR\gamma\delta$ in $\alpha\beta$ versus $\gamma\delta$ fate determination [28]. However, this model failed to explain development of $TCR\gamma\delta$ -dependent DP cells in preTCR-deficient mice [29,30], or that precocious expression of transgenic- $TCR\alpha\beta$ induced appearance of “ $\gamma\delta$ -like” cells [31].

A competing “pre-commitment” model for $\alpha\beta$ versus $\gamma\delta$ lineage choice alternatively proposed that fate determination occurred prior to TCR expression. This initially correlated to heterogeneity in CD127 expression in $CD44^+CD25^+$ DN (DN2) cells; $CD127^{hi}$ cells being biased toward the $\gamma\delta$ lineage [32]. More recently, expression status of Sox-13 has been similarly implicated [33], while commitment potential to the $\gamma\delta$ lineage clearly varies with both ontogeny [22], and with the developmental stage at which $TCR\gamma\delta$ is initially expressed [34]. Nonetheless, the extent to which subsequent $TCR\gamma\delta$ signalling can override these pre-committed states remains unclear [35].

“Strong” TCR signalling promotes a $\gamma\delta$ cell fate

Available data now best fit a model in which quantitative differences in TCR signal strength, irrespective of TCR identity, dictate $\alpha\beta$ versus $\gamma\delta$ fate determination; “strong” signalling promotes a $\gamma\delta$ fate, and “weaker” signalling generates $\alpha\beta$ -committed DP cells [36,37]. Operationally, this equates to an instructional model, as TCR $\gamma\delta$ largely provides strong signals while preTCR signalling is weaker. Although stronger signalling from TCR $\gamma\delta$ appeared to correlate with increased ERK1/2 phosphorylation, induction of Egr family transcription factors, and up-regulation of Id3 [36], the molecular pathways that define $\gamma\delta$ commitment are only now being defined (as discussed later). Recent investigations using Delta-like-1-expressing OP9 (OP9-DL1) stromal cell co-culture of TCR $\gamma\delta^{(+)}$ DN3 thymocytes supported a “signal-strength” model [35]. Thus, strong TCR $\gamma\delta$ signalling combined with age and/or stage-specific pre-commitment factors promote a $\gamma\delta$ fate (**Figure 2**).

Generating strong TCR signals; engaging thymic TCR $\gamma\delta$ ligands

The paucity of known murine TCR $\gamma\delta$ ligands has made investigation of ligand engagement during thymic $\gamma\delta$ cell development problematic. Nonetheless, at least three $\gamma\delta$ subsets are implicated in thymic ligand binding; thymus leukemia (TL)-specific $\gamma\delta$ cells, NKT-like $\gamma\delta$ cells, and DETC.

Interaction of TL-specific $\gamma\delta$ cells (in either KN6 or G8 TCR $\gamma\delta$ -transgenic mice) with cognate T10^b or T22^b ligand (from the MHC TL region) during thymic development was variously reported to cause tolerance, deletion, or trafficking of cells to the gut epithelium [38-40].

However, recent experiments with a T22-tetrameric FACS-staining reagent have instead suggested that thymic ligand-engaging TL-specific $\gamma\delta$ cells develop to secrete IFN- γ , whereas thymic ligand-naive TL-specific $\gamma\delta$ cells secrete IL-17A [7]. This study further reported that T22-tetramer-negative $\gamma\delta$ cells, which constitute ~99% of those observed in wild-type mice, share phenotypic features of ligand-naive (i.e. IL-17A-secreting) TL-specific $\gamma\delta$ cells, somewhat contradicting the perceived view of $\gamma\delta$ cells as predominantly IFN- γ -secreting. Importantly, the development of ligand-naive $\gamma\delta$ cells was suggested to result from TCR-oligomerization-mediated ligand-independent TCR $\gamma\delta$ signalling (see below) [7]. A subsequent report additionally suggested that TL-specific TCR $\gamma\delta^{(+)}$ IELs also lack evidence of thymic ligand engagement [20].

TCR-ligand-mediated selection is also assumed for thymic development of NKT-like $\gamma\delta$ cells, as their characteristic $V\gamma 1^+V\delta 6.3^+$ (or $V\delta 6.4^+$) TCR displays restricted $V\delta$ -CDR3 length and amino acid composition [11,41]. Like TCR $\alpha\beta^+$ NKT cells, NKT-like $\gamma\delta$ cells are dependent on SAP signalling [42] and the transcription factors ThPOK [43] and PLZF [44], the latter being necessary for IL-4 and IFN- γ secretion [44]. However, although PLZF is induced by TCR $\gamma\delta$ cross-linking [44], disruption of the LAT–Itk–Id3 signalling pathway, that functions downstream of TCR $\gamma\delta$, paradoxically promotes expansion of $V\gamma 1^+V\delta 6.3^+$ cells [42,45-47]. Consistent with this, attenuation of TCR signalling appeared to expand the $V\gamma 1^+V\delta 6.3^+$ subset, and elevated PLZF levels in those expanded cells [42]. Thus, very strong ligand-dependent TCR $\gamma\delta$ signalling may not favour development of NKT-like $\gamma\delta$ cells.

Finally, selection through TCR $\gamma\delta$ is also implicated in DETC development, which correlates with thymic stromal expression of immunoglobulin superfamily gene *Skint1* [48,49]. Although

not necessarily a direct ligand for the TCR, $V\gamma 5^+V\delta 1^+$ fetal thymic progenitors that engage $Skint1^+$ stromal cells upregulate *Egr3* that, together with NFAT and $\text{NF}\kappa\text{B}$ signalling, promote the DETC phenotype that involves up-regulation of *Tbx21* and IFN- γ -secreting potential [5]. By contrast, $V\gamma 5^+V\delta 1^+$ progenitors that develop in the absence of *Skint1* fail to induce *Egr3* and *Tbx21*, but express both *Sox13* and *Rorc* that jointly promote what can be called a “non-selected” phenotype that includes IL-17A-producing potential. Importantly, the reciprocal regulation of *Egr3* versus *Sox13/Rorc* could be demonstrated in adult $\gamma\delta$ thymocytes by cross-linking with agonist anti-TCR δ antibody. Thus, this study begins to provide critical insight into the molecular mechanisms that relate $\gamma\delta$ cell functional specification to thymic ligand engagement.

Ligand-independent TCR $\gamma\delta$ signalling

Despite the acknowledged presence of certain thymic TCR $\gamma\delta$ -ligands, and that TCR $\gamma\delta$ signalling is considered ligand-driven in peripheral immune responses, ligand engagement may not mediate all instances of TCR $\gamma\delta$ signal initiation in DN thymocytes. Ligand-independent signal initiation has long been demonstrated for preTCR [50], being variously ascribed to pT α -mediated lipid-raft association [51], preTCR oligomerization mediated by the extracellular Ig-loop of pT α [52,53], or to an intrinsically low signalling threshold in DN thymocytes [54]. Thymic ligand-independent signalling was similarly proposed for TCR $\gamma\delta$, possibly mediated by oligomerization of the variable region of TCR δ [7]. In addition, a recent study suggested that “TCR $\gamma\delta$ ” complexes that lack variable domains, or that lack both variable and constant Ig-like domains, can still initiate signals that drive RAG-2-deficient thymocytes toward a “ $\gamma\delta$ -like” fate

[6]. This implies that appropriate surface pairings of TCR γ and TCR δ chains that possibly bring CD3 ϵ -containing signalling modules into close proximity of available Lck is sufficient for TCR $\gamma\delta$ signal initiation in DN thymocytes. Thus, strong TCR $\gamma\delta$ signalling may not only be a consequence of ligand engagement; additionally, efficiently paired TCR γ /TCR δ chains that are expressed at the cell surface above a certain critical threshold will also commit DN progenitors to a $\gamma\delta$ cell fate [6].

Mapping thymic TCR $\gamma\delta$ signalling to peripheral $\gamma\delta$ effector fate

Studies on DETC, NKT-like, and antigen-experienced TL-specific $\gamma\delta$ cells clearly associate thymic ligand binding with a CD44⁺CD62L⁻CD122⁺ phenotype and IFN- γ -secreting potential [5,7,44]. Nonetheless, the majority of CD27⁺ lymphoid $\gamma\delta$ cells also produce abundant IFN- γ when activated [16], despite displaying a contrasting CD44⁻CD62L⁺CD122⁻ phenotype that implies an absence of thymic TCR-ligand engagement. Thus, we suggest that thymic ligand-independent TCR $\gamma\delta$ signalling may be sufficient to promote $\gamma\delta$ cell commitment to subsequent IFN- γ production (and not to IL-17A production as suggested in previous reports [7]). Moreover, we also propose that a significant component of thymic TCR-ligand engagement may actually be interaction with ligand-presenting cells that provide critical additional signals for subsequent $\gamma\delta$ cell effector function; this may include provision of SAP-dependent signalling for NKT-like $\gamma\delta$ cells, or access to Skint1 for developing DETC.

By contrast to IFN- γ secretion, IL-17A production by $\gamma\delta$ cells has been proposed as a default pathway in which TCR $\gamma\delta$ ⁺ thymocytes do not encounter agonist TCR-ligand [7]. However, IL-17A-secreting CD27⁻ $\gamma\delta$ thymocytes display a uniformly “activated” CD44⁺CD62L⁻ thymic

phenotype similar to ligand-experienced $\gamma\delta$ cells (although without CD122 or NK1.1 expression) [17,25]. Indeed, early studies suggested that $V\gamma6^+V\delta1^+$ thymic progenitors (that mature to secrete IL-17A) undergo ligand-driven TCR selection for canonical CDR3 sequences to a similar degree as $V\gamma5^+V\delta1^+$ DETC progenitors [55]. This notwithstanding, it is unlikely that any such IL-17A-inducing thymic TCR-ligand would behave as a full TCR $\gamma\delta$ agonist [5,7].

Whatever the nature of the inductive event for IL-17A-secreting potential, the thymic progenitors of IL-17A-secreting $\gamma\delta$ cells appear to enter a complex program of development [5] that results from some significant degree of fetal thymus-associated pre-commitment [14]. It also appears to require signalling pathways that involve B lymphoid kinase [56], TGF β 1 [57], and Hes-1 [15]. Thus, it presently remains unclear whether an IL-17A-secreting $\gamma\delta$ fate truly represents a ligand-independent $\gamma\delta$ cell developmental pathway, or whether fetal/neonatal $\gamma\delta$ progenitors of IL-17A-secreting $\gamma\delta$ cells must also interact with thymic ligands that results in distinct but overlapping phenotypic changes to those observed for DETC and NKT-like $\gamma\delta$ cells.

Implications for $\gamma\delta$ cell function

The common description of $\gamma\delta$ cells as “innate-like” perhaps more accurately reflects tissue-associated $\gamma\delta$ subsets with highly focused TCR specificities that represent prototypic stress-surveillance lymphocytes with rapid responses to autologous stress-antigens [1]. As discussed above, precursors of these populations are likely selected on thymic ligands that endow them with specific effector functions, or reinforce homing to certain body locations [11,23]. These cells respond *en masse* to local tissue insults and make critical contributions to both immune protection and tissue integrity [1]. IFN- γ -secreting DETC represent a well-studied example of

these stress-surveillance lymphocytes. However, it is tempting to also speculate whether IL-17A-producing $\gamma\delta$ subsets, such as $V\gamma 6^+V\delta 1^+$ cells from the female reproductive tract and peritoneal cavity, or $V\gamma 4$ -biased dermal $\gamma\delta$ cells, might also contribute to stress-surveillance. These subsets are predominantly tissue-located, and share the “activated” $CD44^+CD62L^-$ phenotype of DETC and NKT-like $\gamma\delta$ cells that could suggest some degree of thymic $TCR\gamma\delta$ ligand engagement [16,25]. Certainly their rapid responsiveness to cytokines such as IL-1 and IL-23 would be consistent with an “innate-like” existence [58,59]. Investigations that determine whether IL-17A-producing $\gamma\delta$ progenitors require thymic TCR-ligand interaction for development should clarify this issue. Moreover, they should test the hypothesis that “innate-like” $\gamma\delta$ subsets require $TCR\gamma\delta$ ligand engagement during their thymic development.

Adaptive T cell responses are generally defined as clonal expansions of relatively few antigen-specific lymphocytes. Interestingly, this feature may well be shared by $CD27^+CD44^-CD62L^+CD122^-$ $\gamma\delta$ cells that comprise the majority of $\gamma\delta$ cells in secondary lymphoid organs [16]. This subset rapidly and extensively expands to secrete abundant IFN- γ on activation through $TCR\gamma\delta$ (and CD27), and was shown to include Murid herpesvirus-4 responsive cells [59]. Thus, $\gamma\delta$ cells which lack evidence of thymic TCR-ligand engagement (i.e. with a $CD27^+CD44^-CD62L^+CD122^-$ phenotype), that likely develop as a consequence of ligand-independent $TCR\gamma\delta$ signalling, appear to demonstrate peripheral “adaptive-like” responses, that includes abundant secretion of IFN- γ , on recognising non-thymic (possibly pathogen-associated) antigens. Nonetheless, there is little evidence of $\gamma\delta$ memory cell generation or “fixing” of TCR specificities in the $TCR\gamma\delta$ repertoire as a consequence of these expansions

[7,26]; instead expanded $\gamma\delta$ cell clones appear relatively short-lived, being replaced by “fresh” naive $\gamma\delta$ cells that presumably maintain a diverse TCR $\gamma\delta$ repertoire.

Similarities with non-T lymphocyte development

Despite persistent temptation to align $\gamma\delta$ cells with their $\alpha\beta$ T cell cousins, comparison with non-T lymphocytes may instead reveal much about $\gamma\delta$ cell biology. For example, developing B cells whose B cell receptors (BCRs) engage self-antigen often develop as B-1 B cells, a subset with "innate-like" features that includes rapid functional responses and a restricted BCR repertoire [60]. Conversely, B cells expressing BCRs with no apparent self-reactivity primarily differentiate into "conventional" follicular B cells with classic “adaptive-like” qualities. Interestingly, BCR signalling of “intermediate” strength in response to limiting self-antigen drives marginal zone B cell generation in a BAFF-dependent manner [61]. Here, BCR signalling induces expression of non-canonical NF κ B pathway substrate p100 which suppresses survival and differentiation unless converted to active p52 by BAFF signalling [62]. This demonstrates that different B cell fates are generated by different “qualities” of BCR signalling that may or may not require ligand engagement and/or input from additional signalling pathways [60]. Thus, an alternative developmental perspective from a distantly-related lymphocyte relative may provide fresh insight on the generation of different $\gamma\delta$ effector fates that possibly result from similar differences in TCR $\gamma\delta$ signalling and co-stimulation.

Concluding remarks

Recent studies have reinforced the importance of $\gamma\delta$ cell responses in infections, cancer and autoimmunity [1]. Indeed, the administration of autologous activated human $V\gamma 9^+V\delta 2^+$ $\gamma\delta$ cells now represents a promising approach for immunotherapy in diverse disease scenarios [2,3]. Clearly, an improved knowledge of $\gamma\delta$ cell biology is essential, and great strides have been taken to characterise $\gamma\delta$ subset phenotypes and functions throughout the body. A thorough understanding of thymic $\gamma\delta$ cell development is equally important, at the forefront of which is elucidation of TCR $\gamma\delta$ -mediated selection events. Here, we propose that thymic engagement of TCR $\gamma\delta$ ligands generates “innate-like” $\gamma\delta$ subsets with rapid cytokine responses to stress-associated stimuli. By contrast, $\gamma\delta$ cells that do not recognise thymic ligands develop as “adaptive-like” $\gamma\delta$ cells that expand clonally to secrete IFN- γ in response to non-thymic and potentially foreign TCR $\gamma\delta$ -ligands. Thus, we contend that thymic TCR selection plays a critical role in ascribing appropriate functional responses to self/non-self TCR $\gamma\delta$ specificities.

Box 1. Outstanding Questions

- What are the cognate ligands for murine TCR $\gamma\delta$? Do these include self-ligands? Have canonical $\gamma\delta$ TCRs been preserved through evolution for self-ligand recognition?
- To what extent are thymic progenitors pre-committed to “innate-like” $\gamma\delta$ effector fates? Do “innate-like” $\gamma\delta$ and $\alpha\beta$ T cells share a common thymic progenitor?
- To what extent is thymic TCR $\gamma\delta$ -agonist engagement required for $\gamma\delta$ cell development? What contribution does thymic ligand-independent TCR $\gamma\delta$ signalling make to the peripheral $\gamma\delta$ pool? How does ligand-independent TCR $\gamma\delta$ signalling initiate?
- How is the IL-17A-secreting effector program initiated during thymic $\gamma\delta$ cell development? And how do additional inputs, such as TGF β or Notch signalling, affect this process?
- What are the signalling pathways, molecular mechanisms, and transcription factors that regulate thymic commitment to diverse $\gamma\delta$ cell effector fates?

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Figure Legends

Figure 1. Thymic and peripheral $\gamma\delta$ subsets. Phenotypic characterisation of thymic and peripheral $\gamma\delta$ subsets based on surface expression of CD25 (purple segments; “25”), CD24 (green segments; “24”), CD27 (yellow segments; “27”), CD44 (blue segments; “44”) and CD122 (red segments; “122”). The predominant cytokines (IFN- γ , IL17A and IL-4) produced by a subset on activation *ex vivo* are illustrated. The term “V γ 4-biased” indicates a $\gamma\delta$ subset that uses the V γ 4 region of TCR γ more frequently than expected. TCR $\gamma\delta$ expression is first seen in the thymus on CD27⁺CD24⁺CD25⁺ progenitors (**a**) that do not express CD44 or CD122. This population rapidly down-regulates CD25 and increases TCR $\gamma\delta$ surface expression to generate immature CD27⁺CD24⁺CD25⁻CD44⁻CD122⁻ cells (**b**) that are likely precursors of at least three mature CD24⁻ thymic subsets. TCR-agonist engagement preserves CD27 expression and up-regulates CD44, CD122 (**c**). This population likely includes precursors of peripheral NKT-like $\gamma\delta$ cells (**d**) that predominantly express a V γ 1⁺V δ 6.3⁺ (or V δ 6.4⁺) TCR; and possibly TL-specific $\gamma\delta$ cells that develop in a T10^b or T22^b-expressing background (**e**). Interestingly, this thymic phenotype (**c**) is shared by thymic DETC progenitors (**f**) during early ontogeny that are also thought to be TCR-agonist selected. Mature DETC (**g**), NKT-like, and TL-specific $\gamma\delta$ cells all secrete abundant IFN- γ . Progenitor (**b**) also likely generates CD27⁺CD44⁻CD122⁻ cells (**h**) through prolonged ligand-independent signalling, that seed the peripheral lymphoid organs (**i**) (and possibly the gut; not shown), where they generate abundant IFN- γ on activation. Progenitor (**b**) also appears to generate thymic IL-17A-secreting cells that are CD27⁻CD44⁺CD122⁻ (**j**). It is presently unclear how this transition proceeds; we suggest that TCR $\gamma\delta$ engagement may be required, although both ligand-independent TCR $\gamma\delta$ signalling and some element of pre-commitment have also been proposed. These cells (**j**) are likely precursors of

IL-17A-producing $\gamma\delta$ cells from the peritoneal cavity and female reproductive tract (**k**), dermis (**l**) and lymph nodes (**m**). Data is from references [7,9-17,20,25,47,48,63].

Figure 2. TCR signal strength and pre-commitment factors dictate $\gamma\delta$ versus $\alpha\beta$ lineage fate. Strength of TCR signalling rather than type of TCR complex (i.e. preTCR vs. TCR $\gamma\delta$) appears to dictate $\gamma\delta$ versus $\alpha\beta$ lineage fate; stronger TCR signals commit early thymocyte progenitors to a $\gamma\delta$ fate, whereas weaker signals promote an $\alpha\beta$ T cell fate. Operationally, this represents an instructional model of commitment, as TCR $\gamma\delta$ is known to signal more strongly than preTCR (predicted ranges of signal strength for preTCR and TCR $\gamma\delta$ are indicated). Nonetheless, factors unrelated to strength of TCR signalling, such as age (fetal vs. adult), developmental stage at which TCR signalling initiates (i.e. DN2 vs. DN3 vs. DN4), and expression of certain “pre-commitment” factors (e.g. CD127 or Sox-13), will impact on the likelihood that a particular signal strength will result in commitment to either lineage.

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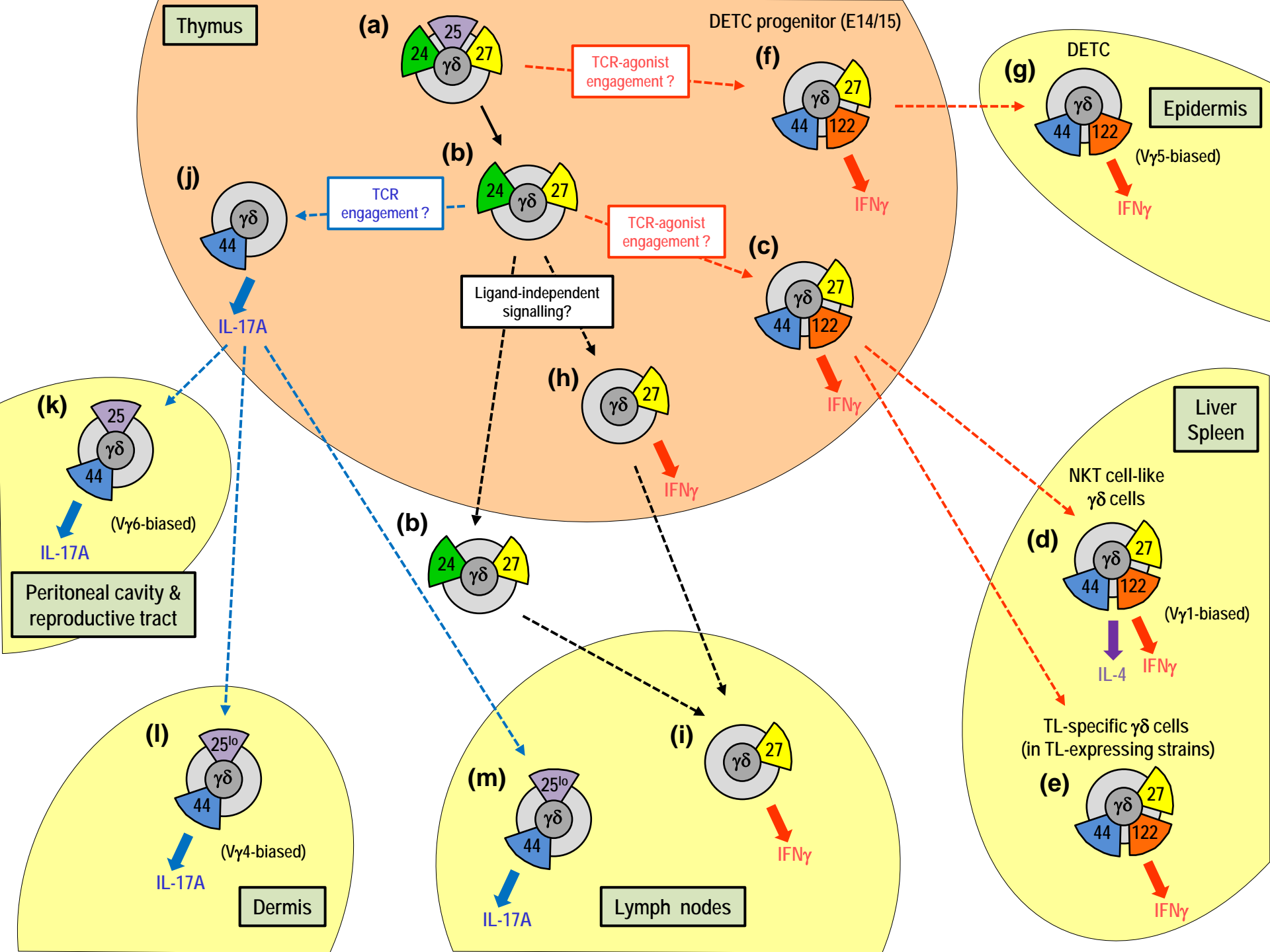
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	IFN- γ	IL-17A	CD24	CD25	CD27	CD44	CD62L	CD122	NK1.1	CCR6	CCR9	SCART2
DETC (V γ 5-biased)	+ ^[5] (Thy)	- ^[5,9] (Thy Sk)	- ^[48] (Thy)	- ^[5,9] (Thy Sk)	+ ^[5] (Thy)	+ ^[9] (Sk)	ND	+ ^[5,48] (Thy Sk)	+ ^[5] (Thy)	- ^[12] (Sk)	ND	- ^[5,13] (Thy Sk)
NKT-like $\gamma\delta$ cells (V γ 1-biased)	+ ^[11] (Thy)	ND	- ^[11] (Thy)	ND	- ^[16] (Thy)	+ ^[11] (Thy)	- ^[11] (Thy)	+ ^[47] (Thy)	Het ^[11] (Thy)	- ^[17] (Thy)	ND	ND
TL-specific $\gamma\delta$ cells (on TL ^b background)	+ ^[7] (Spl)	- ^[7]	ND	- ^[7] (Spl)	ND	+ ^[7] (Spl)	ND	+ ^[7] (Spl)	Het ^[7] (Spl)	ND	- ^[20] (Thy)	ND
Gut $\gamma\delta$ IELs (V γ 7-biased)	lo ^[16]	- ^[19,63]	ND	- ^[16]	+ ^[16]	lo ^[63]	ND	lo ^[19]	ND	- ^[17]	+ ^[19]	ND
Lymphoid $\gamma\delta^{27+}$ cells (CD122 ⁺ NK1.1 ⁻)	+ ^[16]	- ^[16]	- ^[16]	- ^[16]	+ ^[16]	- ^[16]	+ ^[16]	- ^[16]	- ^[16]	- ^[17]	ND	ND
Lymphoid $\gamma\delta^{27-}$ cells	- ^[16]	+ ^[16]	- ^[16]	lo ^[16]	- ^[16]	+ ^[16]	- ^[16]	- ^[16]	ND	Het ^[17] (LN)	Het ^[17]	Het ^[13] (LN)
Reproductive tract and peritoneal $\gamma\delta$ cells (V γ 6-biased)	- ^[14]	+ ^[14]	ND	lo ^[14]	- ^[15]	+ ^[14]	ND	- ^[14]	- ^[16]	ND	ND	ND
Dermal $\gamma\delta$ cells (V γ 4-biased)	ND	+ ^[9,12]	ND	lo ^[9]	ND	+ ^[9]	ND	ND	ND	+ ^[12]	ND	Het ^[13]

Expression of markers and cytokines by $\gamma\delta$ subsets

“+” indicates expression of marker by a subset; “-” indicates absence of marker; “lo” indicates low expression of marker; “Het” indicates heterogeneous expression of marker; “ND” indicates expression not determined. Brackets indicate organ where expression was specifically observed; “Thy” is thymus; “Spl” is spleen; “LN” is lymph nodes; “Sk” is skin. References are provided in square brackets.

