# $\gamma\delta$ T lymphocytes in the diagnosis of human T cell receptor immunodeficiencies

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#### **INTRODUCTION**

Human T cell receptor (TCR) immunodeficiencies (TCRID) are rare autosomal recessive disorders caused by mutations affecting TCR, CD3, or CD247 chains, which share developmental, functional, and TCR expression defects (1). Their rapid diagnosis is fundamental for patient survival and early hematopoietic stem cell transplantation. Here, we propose that studying  $\gamma\delta$  T cells, which are often neglected, can be helpful for a timely diagnosis. We thus offer a diagnostic flowchart and some lab tricks based on published cases.

## γδ T CELL AND TCR PHYSIOPATHOLOGY

γδ T lymphocytes are a minor subset (1-10%) of human peripheral blood T cells. Most (>70%) are CD4<sup>-</sup>CD8<sup>-</sup> [double negative (DN)], some (30%) are CD8<sup>+</sup>CD4<sup>-</sup> and very few (<1%) are CD4<sup>+</sup>CD8<sup>-</sup> [CD8<sup>+</sup> or CD4<sup>+</sup> single positive (SP), respectively]. Most γδ T cells in adults express Vδ2/Vγ9 TCR variable regions (65-90%), the rest being mostly V $\delta$ 1<sup>+</sup>, some V $\delta$ 3<sup>+</sup> or V $\delta$ 5<sup>+</sup>, all with different Vy chains (2). As peripheral blood  $\gamma\delta$  T cells are scarce, their over-representation is more conspicuous than their under-representation, which is very rarely reported and normally associated to a single subset, such as  $V\delta 2^+$  in granulomatosis (3) or aging (4). Indeed, no selective  $\gamma\delta$  T cell immunodeficiency (ID) has been reported to date, although absence of  $\gamma\delta$  T cells has been described together with other lymphocyte derangements in rare primary ID (5). The clinical significance of increased  $\gamma\delta$  T cells, defined as >10% of peripheral blood T lymphocytes (6), requires clarification in several diseases including infection, autoimmunity, cancer, and primary ID.

The human  $\gamma\delta$ TCR (**Figure 1A** inset) is an octameric protein complex composed of three heterodimers (TCR $\gamma$ /TCR $\delta$ ,  $CD3\gamma/CD3\epsilon$ , and  $CD3\delta/CD3\epsilon$ ) and a single CD247 homodimer (also termed  $\zeta/\zeta$ ). The complex can be abbreviated as γδTCR/γεδεζζ. The TCRγ/TCRδ heterodimer contains variable regions, which allow for antigen recognition, while the other three dimers are invariant and are required for surface TCR expression and for intracellular propagation of the recognition signal (7). Therefore, defects in any chain would expectedly impact γδTCR expression and  $\gamma\delta$  T cell selection and function.

### γδ T CELLS IN TCRID

 $\alpha\beta$  T cells have been extensively studied in TCRID. In contrast,  $\gamma\delta$  T cells have been frequently ignored, in part due to their scarcity but also to the lack of markers

other than the TCR to identify them when TCR expression is reduced, as is the case in TCRID. Although their functions are still debated, we believe that their accurate study (relative numbers, **Figure 1A**, TCR expression, **Figure 1B**, and main subsets) can help to diagnose TCRID, as reviewed below and summarized in a practical diagnostic flowchart in **Figure 1C**.

TCRa deficient patients showed combined ID and autoimmune features due to a selective block in  $\alpha\beta$  T cell development, as TCRa is part of the TCRa/TCRB (αβTCR, Figure 1A inset) antigen-binding heterodimer (9). In contrast, the  $\gamma\delta TCR$ was unaffected, as demonstrated by normal surface expression (Figure 1B), which allowed for normal absolute but increased relative numbers of  $\gamma \delta$  T cells (Figure 1A). This is unique among TCRID and thus a useful feature in the diagnostic flowchart (**Figure 1C**). Such  $\gamma \delta$  T cells were proposed to be in part protective from infections in the two reported patients. Indeed,  $\gamma\delta$ T cells are involved in immune responses against a variety of pathogens including virus, bacteria, and parasites, whereas still other act as antigen-presenting cells (10) or B cell helper cells (11). Their beneficial effects in vivo have found recent unexpected recognition in haploidentical allogeneic hematopoietic cell transplantation after depletion of  $\alpha\beta$  T and B cells (12),

Abbreviations: DN, double negative; DP, double positive; ID, immunodeficiency; mAb, monoclonal antibody; SCID, severe combined immunodeficiency; SP, single positive; TCR, T cell receptor; TCRID, T cell receptor immunodeficiencies.



**FIGURE 1 |**  $\gamma\delta$ **T cells in TCRID**. (**A**) Proportion of  $\gamma\delta$ T cells within the T cell compartment. The percentage of  $\gamma\delta$ T cells (mean ± SEM) was defined as  $\gamma\delta$ TCR<sup>+</sup> using 11F2, IMMU510, or anti-TCR $\delta$ -1 monoclonal antibody (mAb) within the T lymphocyte gate (defined as CD3<sup>+</sup>) and ordered from left to right in decreasing values. The gray band indicates the normal range for infants (8). Inset: human TCR isotypes. NA: not analyzable (no T cells); \*: leaky mutations (partial defects); *n*: number of patients for which data was available. (**B**) % TCR surface expression (mean ± SEM) in  $\gamma\delta$  or  $\alpha\beta$ T cells relative to healthy donors. TCR surface expression was determined by flow cytometry using different anti-CD3 mAb,  $\gamma\delta$ T cells were identified as in (**A**) and  $\alpha\beta$ T cells as  $\gamma\delta$ TCR<sup>-</sup>CD3<sup>+</sup> or CD4<sup>+</sup> cells. ND: not determined. (**C**) Our suggested TCRID diagnostic flow cytometry data (top) to point to the most likely culprit TCR chain (bottom). TCR chains are represented by black boxes arranged according to the proportion of  $\gamma\delta$ T cells from (**A**) and their surface TCR expression relative to normal controls from (**B**). The white box indicates normal value. \*: as in (**A**). Brackets represent expected defects, as  $\gamma\delta$ T cells values were not available in these TCRID.

which showed that  $\gamma\delta$  T cells did not cause graft vs. host disease and may have helped with host immune maintenance and recovery. The fact is that, compared to other complete TCRID, symptoms in both TCR $\alpha$ deficient patients appeared rather late (6 and 15 months of age) and transplantation took place very late (6–7 years of age).

Similar to TCRa deficient patients, patients with partial CD38 deficiency  $(CD3\delta^*$  in **Figure 1**) due to a leaky splicing mutation showed strongly reduced a BT cell numbers and normal absolute but high relative numbers of  $\gamma \delta$  T cells (Figure 1A), although with low surface TCR expression [(13) and Figure 1B]. In contrast to TCRa deficiency, partial CD38 deficiency showed early severe combined ID (SCID) features and required very early transplantation (before 2 years of age), thus their  $\gamma \delta T$  cells were not protective, perhaps as a consequence of their impaired TCR expression and function (13). Unexpectedly, partial CD38 deficiency caused a stronger impact in  $\gamma\delta$  (Figure 1B) than in  $\alpha\beta$ TCR surface expression (25 vs. 55% relative to controls (13). A detailed study of their  $\gamma\delta$  T cells showed an enrichment in a subset of otherwise rare CD4<sup>+</sup>  $\gamma\delta$  T cells, which exhibited an activated phenotype and were refractory to further TCR stimulation (14). This CD4-expressing y8 T cell subset seems to be pathognomonic for partial CD38 deficiency, since: (i) it has been ascertained in three of three tested patients with this condition and (ii) it was not found in other TCRID (14, 15). Its developmental origin deserves further comment.  $\alpha\beta$  and  $\gamma\delta$  T cells differentiate within the thymus from a late DN common progenitor (16). In humans, development of most γδ T cells seems to mimic that of  $\alpha\beta$  T cells: from DN progenitors through a CD4+CD8+ double positive (DP) pathway (17, 18), to DN and either CD4<sup>+</sup> or CD8<sup>+</sup> SP populations. DN and CD8<sup>+</sup> SP γδ T cells are minor intrathymic subsets but become the major  $\gamma\delta$  T cell subsets in the periphery, while CD4<sup>+</sup> SP are the main intrathymic subset, followed by DP (19). Notably, the last two subsets can be found in peripheral blood in pathological conditions, and most bone marrow and peripheral blood  $\gamma\delta$  T cells from patients with  $\gamma\delta$  T cell acute lymphoblastic leukemia are either CD4<sup>+</sup> SP or DP (20). Thus, we believe that the 10-fold enrichment of CD4<sup>+</sup> SP γδ T cells observed

in patients with partial CD3 $\delta$  deficiency is due to low TCR-dependence for positive selection of CD4<sup>-</sup>  $\gamma\delta$  T cells and disrupted negative selection of CD4<sup>+</sup>  $\gamma\delta$  T cells (14).

CD3 $\gamma$  deficient patients, most of which showed mild ID (21), had normal numbers of polyclonal peripheral blood  $\gamma\delta$  T cells [absolute and relative, (22) and **Figure 1A**] with low surface TCR [around 30% of control levels (22) and **Figure 1B**], similarly to their  $\alpha\beta$  T cell counterparts (23), likely with an abnormal  $\gamma\delta$ TCR/ $\delta\epsilon\delta\epsilon\zeta$  stoichiometry. Despite their high homology, the invariant CD3 $\gamma$  and  $\delta$  chains show different roles in human vs. mouse  $\gamma\delta$  T cell development. Indeed, CD3 $\gamma$ -deficient mice exhibited a severe  $\gamma\delta$  T cell developmental block (24).

 $\gamma\delta$  T cells were studied in only two of three reported CD247 deficient patients (21). The patients showed SCID features and reduced absolute and relative  $\gamma\delta$  T cells numbers (**Figure 1A**). Surface  $\gamma\delta$ TCR expression was also reduced (4% vs. healthy controls, **Figure 1B**). The number of  $\alpha\beta$ T cells was only slightly reduced despite their similarly reduced surface TCR, with all reported cases showing reduced numbers of CD4<sup>+</sup> T cells but normal or high numbers of CD8<sup>+</sup> T cells (21, 25–27).

A single patient with partial CD3 $\epsilon$  deficiency (28–30) showed very low surface  $\alpha\beta$ TCR expression (10% of normal levels, **Figure 1B**, CD3 $\epsilon^*$ ), mild ID, normal CD8<sup>+</sup>, and reduced CD4<sup>+</sup> ( $\alpha\beta$ ) T cells, but no  $\gamma\delta$  T cells as determined with the anti-TCR $\delta$ -1 monoclonal antibody (mAb) (**Figure 1A**). We have however considered for **Figure 1C** that surface  $\gamma\delta$ TCR expression might have been similar to  $\alpha\beta$ TCR expression.

Three studied CD38 deficient patients out of 16 reported, all with severe T cell lymphopenia and SCID (31, 32)], showed a few circulating CD3<sup>+</sup> T cells, which were DN but  $\gamma\delta TCR^{-}$  (33).  $\gamma\delta TCR^{+}$  cells were indeed undetectable by flow cytometry in peripheral blood or by immunohistochemistry in the thymus, lymph nodes, spleen, or gut. However, gene microarray analysis and protein expression of patient thymocytes showed increased levels of TCRy and TCR $\delta$  transcripts and proteins (33), which could be interpreted as presence and thus significant selection of y8 T cells unable however to leave the thymus, perhaps due to insufficient surface TCR compared to partial CD38 deficiency.

Finally,  $\gamma\delta$  T cells have not been studied in SCID patients with complete CD3 $\epsilon$  deficiency (31). Nevertheless, given their severe T cell lymphopenia, we can safely presume for **Figure 1C** that they were absent.

In summary, the proportion of  $\gamma\delta$  T cells within total T lymphocytes (Figure 1A) and the level of surface  $\gamma\delta$  vs.  $\alpha\beta TCR$ expression (Figure 1B), as well as the severity of lymphopenia (T<sup>-</sup>B<sup>+</sup>NK<sup>+</sup> or  $T^{\pm}B^+NK^+$  phenotype), can be used to generate a practical TCRID diagnostic flowchart (Figure 1C). For instance, if an infant has SCID and no T cells but normal B and NK cell numbers (T<sup>-</sup>B<sup>+</sup>NK<sup>+</sup> phenotype) and other causes have been ruled out, CD38 or CD3e deficiency should be considered (Figure 1C). In contrast, if some T cells are present  $(T^{\pm}B^+NK^+$  phenotype) and yo TCR expression is low, TCRa deficiency can be ruled out. If CD16 expression by NK cells is normal, CD247 deficiency can be excluded, and the presence or absence of high absolute numbers of CD4<sup>+</sup>  $\gamma\delta$  T cells will rule out CD3 $\gamma$  or partial CD38 deficiency, respectively.

# LAB TRICKS TO IDENTIFY $\alpha\beta$ AND $\gamma\delta$ T CELLS IN TCRID

When surface TCR expression is low,  $\alpha\beta$  T cells can be identified by the expression of CD4 or CD8 $\alpha\beta$  (i.e., CD8<sup>bright</sup>) within the lymphoid subset (23), whereas  $\gamma\delta$  T cells are identified only by expression of the  $\gamma\delta$ TCR. We have reported that most CD3<sup>+</sup> cells within normal DN lymphocytes are  $\gamma\delta$  T cells (34), and this may also help in certain TCRID.

Despite their reduced numbers and surface TCR expression, an appropriate multicolor flow cytometry approach can help to identify vo T cells in TCRID. To avoid underestimation due to low TCR surface expression, we recommend: (i) the use of bright fluorochromes such as PE, PE-Cy5.5, PE-Cy7, or APC, rather than FITC, (ii) an appropriate choice of CD3 mAb such as UCHT-1, F101.01, or S4.1 due to their high signal-to-noise ratio in TCRID, (iii) twocolor stainings with CD3 and γδTCR mAb, which can also help to single out  $\gamma\delta$  T cells as a DP subset, and (iv) to avoid mixing  $\alpha\beta$ TCR and CD3 mAb, as they sometimes compete (UCHT-1, for instance).

CD4 and CD8 expression by  $\gamma\delta$  T cells should also be tested to rule out partial CD3 $\delta$  deficiency (see above). CD4, CD8,

 $\gamma\delta$ TCR (IMMU510 or 11F2), and CD3 (UCHT-1 or S4.1) is a useful combination, to this end. Lastly, intracellular stainings for invariant TCR chains has been shown to be useful to identify T cells expressing very low surface TCR, such as those with CD247 (21) or partial CD38 deficiency (14).

# **CONCLUSION AND PERSPECTIVES**

Human  $\gamma\delta$  T lymphocytes are still puzzling in terms of development, function, and TCR stoichiometry in ways that mouse models do not wholly recapitulate. Human TCRID share defects in T cell development and function and in TCR expression. While their  $\alpha\beta$  T cells have been studied in detail.  $\gamma\delta$  T cells have been frequently ignored, in part due to their scarcity and to the lack of appropriate markers to identify them when TCR expression is reduced. Here, pooling published studies, we proposed some technical tricks to identify yo T cells in TCRID patients and made the point that their careful analysis can help to inform a rapid differential diagnosis using a flowchart, with clinical benefit.

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