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Adaptive immune response of $V\gamma 2V\delta 2$ T cells: a new paradigm

Zheng W. Chen^{1,2} and Norman L. Letvin²

¹ Tuberculosis Research Unit, Beth Israel Deaconess Medical Center, Harvard Medical School, 330 Brookline Ave, RE113/RE216, Boston, MA 02115, USA

² Division of Viral Pathogenesis, Beth Israel Deaconess Medical Center, Harvard Medical School, 330 Brookline Ave, RE113/RE216, Boston, MA 02115, USA

Abstract

The role of $\gamma\delta$ T cells in adaptive immunity remains uncertain. Recent studies have demonstrated that a unique subset of $\gamma\delta$ T cells in primates can mount adaptive immune responses during mycobacterial infections. This Review discusses notable similarities and differences in adaptive immune responses between non-peptide-specific $\gamma\delta$ T cells and peptide-specific $\alpha\beta$ T cells, and discusses both the molecular basis for $\gamma\delta$ T-cell responses and potential functions of these enigmatic cells.

$\gamma\delta$ T cells are a minor T-cell population that express T-cell receptors (TCRs) comprised of γ and δ heterodimers. The $\gamma\delta$ T-cell population can be separated into different subsets or subpopulations on the basis of their expression of particular $V\gamma$ or $V\delta$ elements. These subsets can be further divided into two major groups according to tissue distribution and TCR diversity: resident $\gamma\delta$ T cells with or without TCR diversity that take up residence in epithelia or mucosae of the organs, such as skin, lung, intestine, uterus, vagina and tongue; and circulating $\gamma\delta$ T cells with substantial TCR diversity that are found in the blood and lymphoid tissues. Murine $\gamma\delta$ T cells have important roles in immunity to infections, surveillance against tumors, immune regulation, modulating auto-immune responses and epithelial homeostasis [1–3]. By contrast, the roles of human $\gamma\delta$ T cells in immune responses are poorly understood. Although human $V\gamma 2V\delta 2^+$ T cells recognize non-peptide phosphoantigens from bacteria and plants [4], little is known about the nature and functions of immune responses of antigen-specific $\gamma\delta$ T cells. Recent studies have contributed considerably to our understanding of the immune biology and functions of resident and circulating $\gamma\delta$ T cells. This Review discusses new hypotheses and discoveries concerning immunological features of $\gamma\delta$ T cells and how they differ from $\alpha\beta$ T cells.

Major $\gamma\delta$ T-cell subsets in primates and their unique antigen specificity

The definition of antigens recognized by $\gamma\delta$ T cells and the availability of appropriate animal models for their study are of central importance for elucidating the immune functions of human $\gamma\delta$ T cells. $V\gamma 2V\delta 2^+$ T cells comprise the majority of circulating human $\gamma\delta$ T cells and recognize non-peptide antigens from microbes and plants [4]. By contrast, $V\delta 1^+$ T cells are enriched in tissue mucosae and recognize self and foreign lipids presented by CD1 [5] and the stress-inducible MHC class I-related chains A and B (MICA and MICB) [6,7]. The non-peptide antigens recognized by $V\gamma 2V\delta 2^+$ T cells are prenyl pyrophosphates [4], bisphosphonates [8] and alkylamines [9]. $V\gamma 2V\delta 2^+$ T cells also recognize *Staphylococcus* enterotoxin A and canarypox antigens [4,10]. Interestingly, the canarypox-specific and *Mycobacterium bovis*

Calmette–Guérin (BCG)-specific $V\gamma 2V\delta 2^+$ T cells have unique antigen specificity [4,10]. Despite the definition of these non-peptide antigens, studies of immune responses of $V\gamma 2V\delta 2^+$ T cells have been hindered by an absence of relevant animal models. Mice do not express a homologue of the $V\gamma 2V\delta 2^+$ TCR and there is no functional equivalent of these cells in rodents. Because non-human primates are genetically and biologically close to humans, macaques have been exploited as a model system in which to study the immune biology of $\gamma\delta$ T cells. Non-human primates express human TCR homologs, including $\gamma\delta$ TCR [11]. Macaque γ and δ genes encoding variable (V), joining (J) and constant (C) regions have been sequenced by us and others (Table 1) [12–15]. The macaque $V\gamma$ or $V\delta$ elements can share $\leq 91\%$ similarity in amino-acid sequences to their human counterparts, although the frequencies of $V\gamma^+$ and $V\delta^+$ T-cell subsets are not the same as their human counterparts. Given the remarkable similarity in sequences between human and macaque $\gamma\delta$ TCRs, it is predictable that macaque $V\gamma 2V\delta 2^+$ T cells can share antigen specificity with their human counterparts. In fact, macaque $V\gamma 2V\delta 2^+$ T cells can recognize prenyl pyrophosphonates, alkylamines and *M. bovis* BCG phosphoantigens as efficiently as human $V\gamma 2V\delta 2^+$ T cells [14–16]. It is important to note that only primate $V\gamma 2V\delta 2^+$ T cells, not $\gamma\delta$ T cells from non-primate species, recognize the non-peptide antigens identified to date. Given their structural and functional similarities to human $V\gamma 2V\delta 2^+$ T cells, non-human primates should provide invaluable models to explore the diverse roles of microbe-specific $\gamma\delta$ T cells in the immune response to infections as well as to test $V\gamma 2V\delta 2$ T-cell vaccines.

Adaptive immune responses of primate $V\gamma 2V\delta 2^+$ T cells

Unlike $\alpha\beta$ T cells, the roles of $\gamma\delta$ T cells in adaptive immunity remain unclear. It is attractive to suppose that $V\gamma 2V\delta 2^+$ T cells function as a bridge linking innate and adaptive immune responses, given that this subset of $\gamma\delta$ T cells can recognize a broad spectrum of non-peptide antigens. It is also conceivable that $V\gamma 2V\delta 2^+$ T cells can mount adaptive immune responses during microbial infections because these unique $\gamma\delta$ T cells resemble $\alpha\beta$ T cells in their expression of diverse TCRs and have a conserved capacity to proliferate and expand following their TCR recognition of non-peptide antigens in culture [15]. To test this hypothesis, we have examined the role of the TCR in $V\gamma 2V\delta 2^+$ T cells in adaptive immunity using a *Mycobacterium*-infected macaque model. Following *M. bovis* BCG infection, macaque $V\gamma 2V\delta 2^+$ T cells expanded ≤ 25 -fold in percentage and 200-fold in absolute numbers in the peripheral blood. Interestingly, a clear memory-type response of $V\gamma 2V\delta 2^+$ T cells was detected as early as 4–6 days after BCG re-infection and the magnitude of this expansion was 2–9-fold greater than that seen during primary BCG infection. The recall expansion of $V\gamma 2V\delta 2^+$ T cells persisted for as long as seven months after the second BCG inoculation. Importantly, some clonotypic $V\delta 2^+$ T cells exhibited clonal expansion during BCG infection and re-infection, suggesting that antigen-specific $V\gamma 2V\delta 2^+$ T cells participate in primary and memory immune responses [15]. Primary and recall expansions of $V\gamma 2V\delta 2^+$ T cells are also seen following *Mycobacterium tuberculosis* aerosol challenge of naïve and BCG-vaccinated macaques, respectively. The capacity to rapidly expand coincided with reduced *M. tuberculosis* burdens and immunity to fatal tuberculosis in BCG-vaccinated macaques [15]. These results provide evidence that $V\gamma 2V\delta 2^+$ T cells, as well as $\alpha\beta^+$ T cells, contribute to the adaptive immune response in BCG and *M. tuberculosis* infections. Conventionally, the adaptive (memory) immune response of T cells is characterized by their antigen-specific persistence following an initial infection and the rapid and prolonged recall responses on re-infection or re-exposure to the same categories of antigens. The findings in monkey models demonstrate that $V\gamma 2V\delta 2^+$ T cells possess the functional memory of immune responses and suggest that $V\gamma 2V\delta 2^+$ T cells are involved in the immune control of mycobacterial infections.

The adaptive immune responses of $V\gamma 2V\delta 2^+$ T cells following mycobacterial infection of monkeys is similar to the $\gamma\delta$ T-cell expansions detected during bacterial and parasitic infections

in humans [17–36]. Some humans can exhibit an early and prolonged $\gamma\delta$ T-cell expansion. It is difficult to determine in humans whether a $\gamma\delta$ T-cell expansion identified during an infection represents a primary or recall response because a history of previous exposure of $\gamma\delta$ T cells to non-peptide antigens or other infections is usually unknown. Given the typical course of primary and memory immune responses of macaque $V\gamma 2V\delta 2^+$ T cells in mycobacterial infections, an early and prolonged expansion of $\gamma\delta$ T cells in humans might indeed represent a recall or memory response that results from a second exposure to non-peptide antigens during a particular infection. The kinetics and magnitude of the *in vivo* expansion of $\gamma\delta$ T cells in primates with bacterial and parasitic infections appears to be similar to the enhanced ability of $V\gamma 2V\delta 2^+$ T cells of BCG-vaccinated humans to expand *in vitro* following stimulation with an *M. tuberculosis* lysate [37]. This marked *in vitro* expansion of $V\gamma 2V\delta 2^+$ T cells from BCG-vaccinated humans suggests that human $\gamma\delta$ T cells can mount a memory-like response on exposure to non-peptide antigens [37]. The adaptive immune response of macaque $V\gamma 2V\delta 2^+$ T cells appears to be consistent with post-birth selection of predominant human $V\gamma 2V\delta 2^+$ T cells in the circulation. The dominance of human $V\gamma 2V\delta 2^+$ T cells is probably adapted by selection pressure derived from endogenous and exogenous environments. Phenotypic analyses of human $\gamma\delta$ T cells also suggest that $V\gamma 2V\delta 2^+$ T cells can have effector or memory phenotypes based on their expression of the CD45RA and CD27 molecules. Central memory $V\gamma 2V\delta 2^+$ T cells are CD45RA⁻CD27⁺, whereas effector memory cells have lost the expression of CD27 costimulatory molecules and lack the proliferative potential of central memory cells [38]. Interestingly, the frequency of effector $V\gamma 2V\delta 2^+$ T cells are decreased in peripheral-blood mononuclear cells (PBMCs) of humans with pulmonary tuberculosis or active HIV-1 infection [38,39].

The magnitude of $V\gamma 2V\delta 2^+$ T-cell expansions appears to be greater than that of $\alpha\beta$ T cells during bacterial infections. The differences in the magnitudes of these expansions are particularly evident if the frequency of phosphoantigen-specific $V\gamma 2V\delta 2^+$ T cells is compared with those of CD4⁺ and CD8⁺ T cells specific for a single epitope or protein. During BCG infection or re-infection, interferon- γ (IFN- γ)-producing CD4⁺ or CD8⁺ T cells that are specific for a pool of overlapping 12mer peptides spanning Ag85B are usually less than 1–5/1000 T cells, shown by ELISpot or intracellular cytokine assays (Z.W. Chen *et al.*, unpublished). By contrast, $V\gamma 2V\delta 2^+$ T cells comprise $\leq 30\%$ of T cells in BCG-infected monkeys, $\leq 48\%$ in patients with tularemia, salmonellosis, and brucellosis [18–20] and 98% in those with ehrlichiosis [21]. In addition, the recall expansion of $V\gamma 2V\delta 2^+$ T cells can be seen as early as four days after BCG re-infection and lasts seven months to one year after resolution of active infection [15,40]. The longer duration of $V\gamma 2V\delta 2^+$ T-cell expansions compared with $\alpha\beta$ T-cell expansions in individuals with bacterial infections suggests that these cells can more readily undergo expansions in response to repeated infections. The diverse TCR repertoire and capacity of $V\gamma 2V\delta 2$ T cells to mount a memory response might indeed separate this $\gamma\delta$ -cell subset from conventional innate immune cells. Finally, because phospholigands have been demonstrated in mycobacteria, Gram-negative rods, some Gram-positive cocci and parasitic protozoa, $V\gamma 2V\delta 2^+$ T cells might mount memory responses to a variety of pathogens after an initial infection with one of them. This notion is supported by the observation that humans pre-sensitized with BCG can exhibit a $V\gamma 2V\delta 2$ T-cell expansion in response to *in vitro* stimulation with mycobacterial lysate containing phospholigands [37]. The remarkable breadth and magnitude of $V\gamma 2V\delta 2^+$ T-cell responses might, therefore, be unique in representing cross-reactive adaptive immunity to multiple microbes.

Migration of activated $\gamma\delta$ T cells during adaptive immune responses

Most natural infections occur as a result of pathogen invasion through mucosae resulting from airborne, oral or sexually associated transmission. Recruiting immune cells to infected tissues is therefore an important defense mechanism for immune control of infection. In murine models

of infectious diseases, the $\gamma\delta$ T cells involved have been identified at local sites of infections but not in draining lymph nodes or spleen [1]. This contrasts with what has been seen for the homing of peptide-specific $\alpha\beta$ T cells. Although the chemoattraction of leukocytes during inflammation has been studied, tissue trafficking and localization of antigen-specific $\gamma\delta$ T cells in immune responses to infecting microbes are poorly characterized. We have recently demonstrated that rapid recall expansion of $V\gamma 2V\delta 2^+$ T cells is seen in bronchial alveolar lavage (BAL) fluid following *M. tuberculosis* aerosol challenge of BCG-vaccinated monkeys [15]. This expansion of $V\gamma 2V\delta 2^+$ T cells is associated with an inflammatory-cell response characterized by increased numbers of neutrophils and macrophages in BAL fluid. The accumulation of $V\gamma 2V\delta 2^+$ T cells in the lung is probably a result of the recruitment of these cells from the circulation as well as local clonal expansion after *M. tuberculosis* challenge. Interestingly, increases in the number of $V\gamma 2V\delta 2^+$ T cells are also apparent in pulmonary and intestinal mucosae when an expansion of these cells is seen in the blood of monkeys inoculated intravenously with BCG [15]. This increased number of $V\gamma 2V\delta 2^+$ T cells is particularly marked in the lung despite the fact that BCG loads in the lung are extremely low. In addition, no apparent inflammation can be seen in the pulmonary compartment following intravenous BCG inoculation (Z.W. Chen *et al.*, unpublished). Surprisingly, larger increases in numbers of $V\gamma 2V\delta 2^+$ T cells than $\alpha\beta$ T cells are evident in the lungs of the monkeys intravenously inoculated with BCG (Fig. 1). These results suggest that there might be a preferential migration of activated $V\gamma 2V\delta 2^+$ T cells to the lung from the circulation or lymphoid tissues after mycobacterial infection. We cannot exclude the possibility that local expansions of $V\gamma 2V\delta 2^+$ T cells also occur in the pulmonary compartment. $V\gamma 2V\delta 2^+$ T cells in the lung and blood of monkeys infected with *M. tuberculosis* or BCG express remarkably high levels of chemokine receptors CXCR3 and CCR5 (Z.W. Chen *et al.*, unpublished). It is probable that the chemokine receptor-mediated trafficking in response to chemokines is responsible for transendothelial migration of immune cells to the lung from the circulation and/or lymphoid tissues. In fact, Glatzel *et al.* have recently shown that human $V\delta 2^+$ but not $V\delta 1^+$ or $\alpha\beta$ TCR⁺ T cells express high levels of CCR5 and CXCR3 [41]. Cipriani *et al.* have confirmed the high expression of CCR5 and related CC chemokines in $V\gamma 2V\delta 2^+$ T cells [42]. Furthermore, the ligands for CCR5, such as macrophage inflammatory protein-1 α (MIP-1 α), MIP-1 β and RANTES, have been shown to facilitate $\gamma\delta$ T-cell migration in an *in vitro* migration system [43]. Because inflammation after mycobacterial infection is inevitably associated with an increased production of chemokines, these chemokines might have roles in transendothelial chemotaxis of $V\gamma 2V\delta 2^+$ T cells into the lung or other organs. Although the mechanisms responsible for this transendothelial migration have not been characterized, $\gamma\delta$ T-cell-initiated migration pathways involving endothelial cells and/or stromal cells might regulate this process. $V\delta 2^+$ T cells can express the natural-killer (NK) receptor protein 1a (NKR1a or CD161) and engagement of NKR1a on $V\delta 2^+$ T cells results in activation of calcium calmodulin-dependent kinase II [44]. Interestingly, $V\delta 2^+$ T cells use NKR1a for transmigration across endothelial monolayers *in vitro* and this transmigration relies on the calcium calmodulin-dependent kinase II pathway [44,45]. Interleukin-12 (IL-12) appears to enhance CD161 expression and transendothelial migration of $V\gamma 2V\delta 2$ T cells [44]. The unique expression of chemokine receptors on $\gamma\delta$ T cells might explain differences between $\alpha\beta$ and $\gamma\delta$ T cells in tissue trafficking and homing. Thus, these inflammatory cytokines, including the chemotaxis-associated chemokines, probably have a role in the tissue trafficking of $\gamma\delta$ T cells during infections (Fig. 2).

Distinct molecular basis of $\gamma\delta$ T-cell responses

Differences between immune responses of $V\gamma 2V\delta 2^+$ T cells and $\alpha\beta$ T cells during bacterial infections have already been described (Table 2) and distinct molecular characteristics might determine these differences. As discussed, these cell populations express different chemokine receptors and use different receptor-related signaling pathways. Hayes and Love have recently

compared these cell populations through the components of their $\gamma\delta$ TCR and their TCR-induced signal transduction [46]. They found that most $\gamma\delta$ T cells but not $\alpha\beta$ T cells express a $\gamma\delta$ TCR complex that lacks a CD3 δ unit. Moreover, following activation of $\gamma\delta$ T cells, Fc ϵ R1 γ is expressed and included in the $\gamma\delta$ TCR complex. Moreover, TCR-mediated signal transduction and proliferation for $\gamma\delta$ T cells is superior to that seen in $\alpha\beta$ T cells [46]. The efficient signal transduction initiated by $\gamma\delta$ TCR engagement is consistent with the documented ability of V γ 2V δ 2⁺ T cells to expand rapidly in humans and monkeys following bacterial infection. Distinct signaling pathways of $\gamma\delta$ T cells have also been suggested in the studies of transcriptional profiles of intestinal $\gamma\delta$ intraepithelial lymphocytes (IELs) in mice.

Fahrer and colleagues [47] have recently reported that $\gamma\delta$ IELs do not express transcripts for certain key signaling proteins used by $\alpha\beta$ T cells. In addition, $\gamma\delta$ IELs can constitutively express cytotoxic effector genes, such as granzymes A and B, and the inflammatory chemokine RANTES. By contrast, lymph node CD8⁺ $\alpha\beta$ T cells must be activated to express those cytotoxic effector genes. Interestingly, NK inhibitory and activating genes are also constitutively expressed by $\gamma\delta$ IELs [47]. Studies reported by Shires *et al.* show the naïve expression of high levels of granzymes and Fas ligand, chemokines, anti-proliferative genes *Slfm2*, *Btg1* and *Btg2*, and some signaling-related genes, *Rgs1* and *Junb*, in $\gamma\delta$ IELs, although there appears to be no difference in expression of those genes between $\gamma\delta$ IELs and $\alpha\beta$ IELs in naïve mice [48]. These findings at both protein and transcriptional levels support the presumption that there are fundamental differences in some phenotypes and signal transduction machineries of $\gamma\delta$ T cells and $\alpha\beta$ T cells. Molecular differences might enable these two T-cell populations to function differently in innate and adaptive immune responses. In fact, engagement of NKG2D on V γ 2V δ 2⁺ T cells by MICA augments their effector function in response to antigens [49], although it is not known whether this also occurs in the setting of $\alpha\beta$ T cells. Killer Ig-like receptors (KIRs) might also be involved in the immune regulation of V γ 2V δ 2⁺ T cells because KIRs expressed on different $\gamma\delta$ or $\alpha\beta$ T-cell subsets can regulate innate and adaptive immune responses of these cells [50]. It would be useful to compare the molecular differences in TCR complexes, NK receptors, signaling kinases and chemokine and cytokine production between non-peptide-specific V γ 2V δ 2⁺ T cells and peptide-specific $\alpha\beta$ T cells during immune responses to microbial infections.

Potential roles of V γ 2V δ 2⁺ T cells in antimicrobial immunity

V γ 2V δ 2⁺ T cells that expand during bacterial or parasitic infections could contribute to adaptive immunity in various ways. Activated and expanded V γ 2V δ 2⁺ T cells might directly participate in antimicrobial immune responses. V γ 2V δ 2⁺ T cells kill bacteria-infected cells and bacteria [51,52], although this effector function detected *in vitro* might not reflect the *in vivo* function of these cells. On activation, V γ 2V δ 2⁺ T cells can produce a large amount of IFN- γ and tumor necrosis factor- α (TNF- α) [38,53,54], the cytokines important for controlling mycobacterial infection in mice. Through production of those Th1 cytokines, V γ 2V δ 2⁺ T cells could function as a link connecting innate and adaptive immune systems and facilitate the development of adaptive immune responses of antigen-specific $\alpha\beta$ T cells. In fact, V γ 2V δ 2⁺ T cells activated by non-peptide antigens are able to react quickly and protect SCID (severe combined immunodeficiency) mice from bacterial infections by reducing bacterial numbers [55]. The impact of $\gamma\delta$ T cells on innate immune cells, such as macrophages and NK cells, has also been reported in murine models [1]. Mice deficient in $\gamma\delta$ T cells develop enhanced inflammation characterized by disruption of macrophage homeostasis and liver necrosis. In the absence of $\gamma\delta$ T cells, IFN- γ production by NK cells is reduced, which leads to a delay in granuloma formation and an increase in bacterial growth [56]. The $\gamma\delta$ -deficient mice lack the whole population of $\gamma\delta$ T cells, including those subsets with 'innate' phenotypes and, therefore, the results observed in these animals might not represent the potential roles of V γ 2V δ 2⁺ T cells. However, human V γ 2V δ 2⁺ T cells might reserve some of those functions exerted by

murine $\gamma\delta$ T cells. $V\gamma 2V\delta 2^+$ T cells might also regulate immune functions of other immune cells, such as dendritic cells and B cells [57,58], given their ability to produce various cytokines. Finally, non-peptide antigen-specific $\gamma\delta$ T cells could contribute to anti-inflammatory function or tissue repair during infection and disease. This possibility is supported by the recent novel observation that resident $\gamma\delta$ T cells can have unique functions in epithelia or tissue homeostasis [2]. Murine $\gamma\delta$ T cells in the skin can be activated at wound sites and produce cytokines, including keratinocyte growth factors (KGFs) that participate in wound repair. In the absence of skin $\gamma\delta$ T cells, there are defects in keratinocyte proliferation and tissue re-epithelialization following tissue damage [2]. It is conceivable that $V\gamma 2V\delta 2^+$ T cells, once activated locally or recruited to tissue compartments, could participate in tissue repair or wound healing after tissue damage that occurs as a result of active infections and inflammation. The fact that the expansion of $V\gamma 2V\delta 2^+$ T cells is so prolonged after the resolution of bacterial infections in primates [15] suggests that these $\gamma\delta$ T cells might contribute to anti-inflammatory functions rather than simply contributing to the clearance of microbes.

Concluding remarks

In conclusion, primate $V\gamma 2V\delta 2^+$ T cells, as well as $\alpha\beta^+$ T cells, can contribute to adaptive immune responses in microbial infections, despite the fact that these two T-cell populations differ in many of their biological characteristics. The non-peptide phosphoantigen-specific $V\gamma 2V\delta 2^+$ T cells appear to differ from peptide-specific $\alpha\beta$ T cells in their antigen recognition and the magnitude of their immune responses. The unique ability of $V\gamma 2V\delta 2^+$ T cells to expand during mycobacterial infections suggests that vaccine-elicited $V\gamma 2V\delta 2^+$ T-cell immunity might prove beneficial. $V\gamma 2V\delta 2^+$ T cells might broadly contribute to both innate and acquired immunity against microbial infections.

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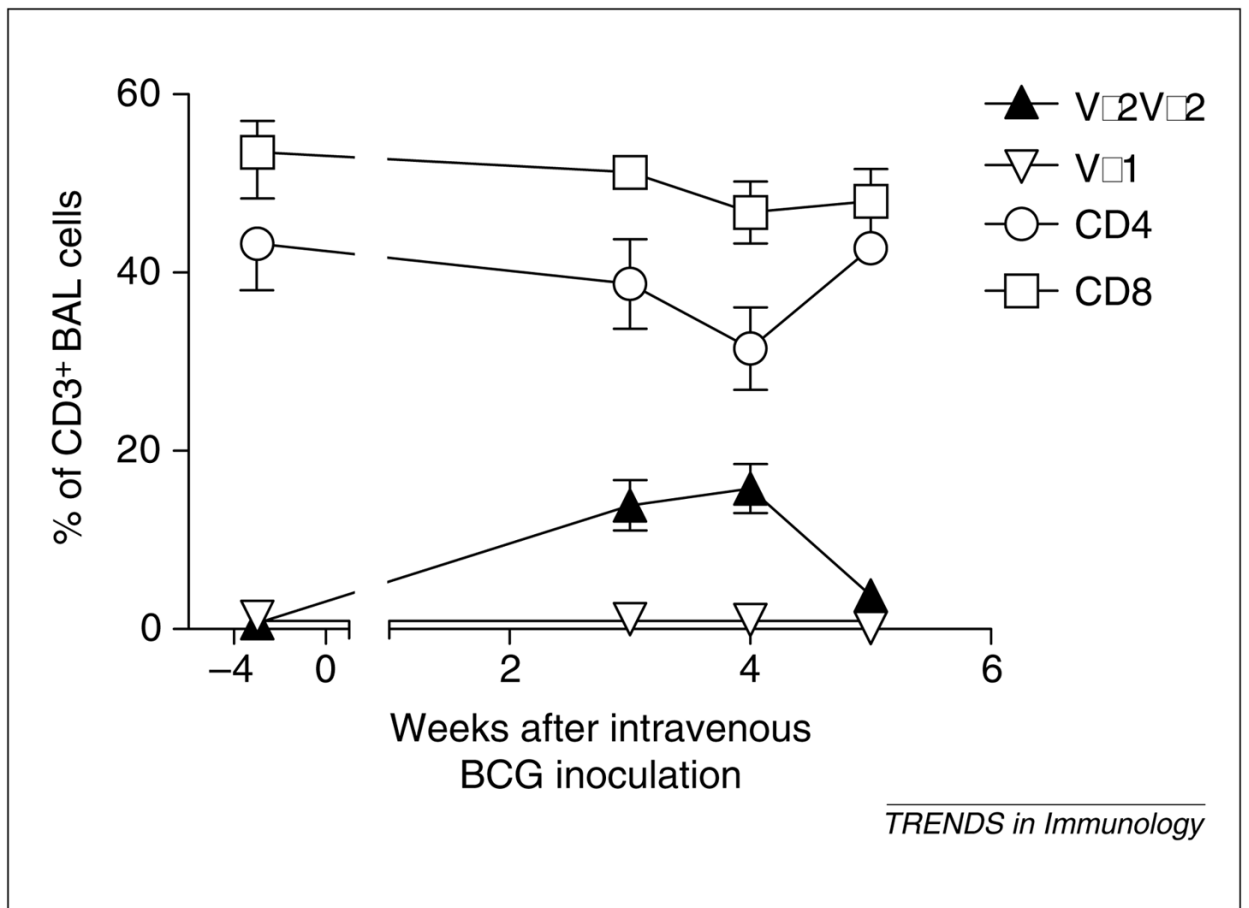
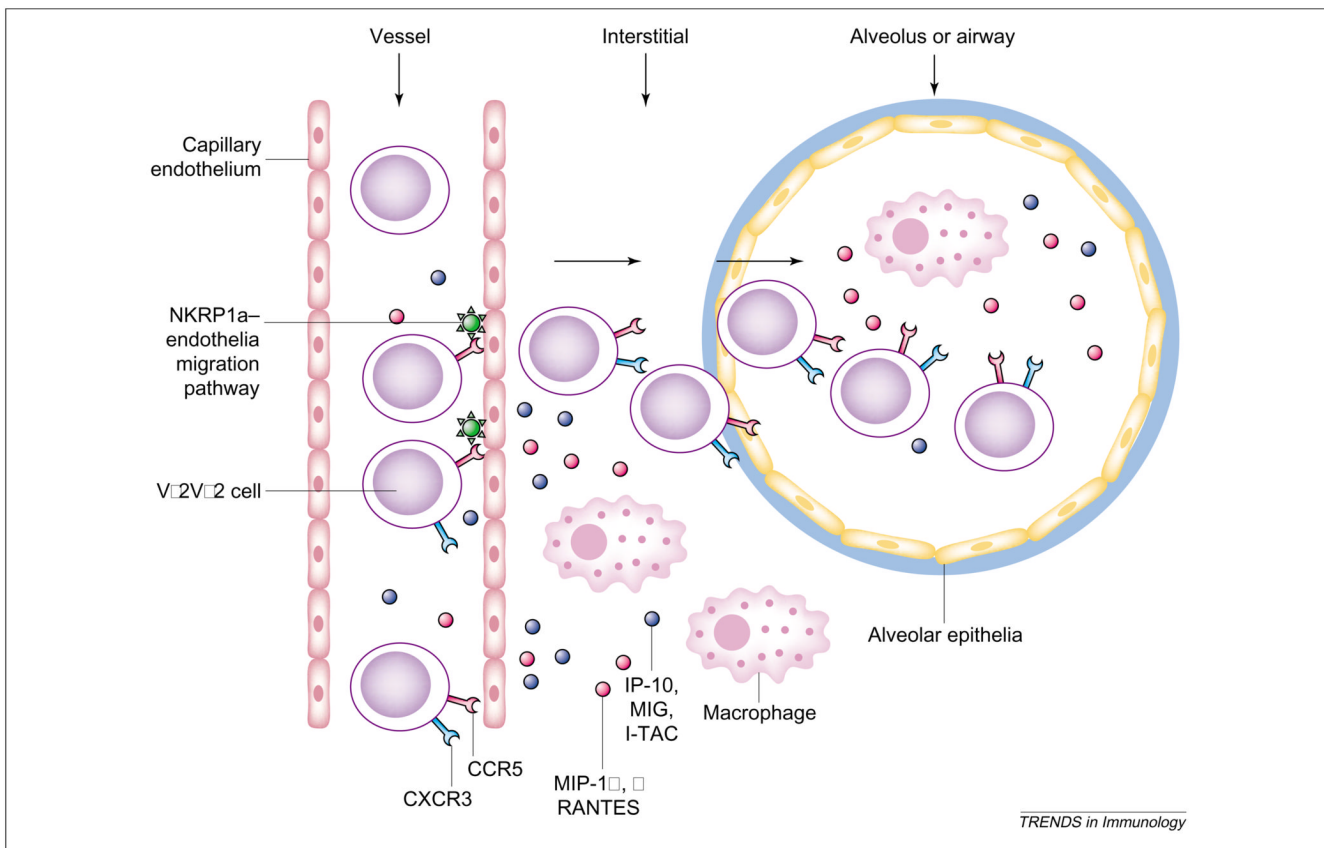


Fig. 1.

Systemic *Mycobacterium bovis* Calmette–Guérin (BCG) infection introduced by intravenous inoculation can result in a preferential increase in V γ 2V δ 2⁺ T cells in bronchial alveolar lavage (BAL) fluid. Note the lack of increase in numbers of CD4⁺ and CD8⁺ T cells at the time the increase in V γ 2V δ 2⁺ T cells is identified. Shown are mean values with the error bars of standard error of the mean (SEM) from four animals.

**Fig. 2.**

Proposed migration of $V\gamma 2V\delta 2$ T cells to lungs or other epithelial compartments during infections. MIP-1 α , MIP-1 β and RANTES (ligands for CCR5) produced after infections in the pulmonary compartment might have a role in chemoattracting CCR5⁺ $V\gamma 2V\delta 2$ T cells to the lung from the circulation or lymphoid tissues [41–43]. Because $V\gamma 2V\delta 2$ T cells also express CXCR3, these cells are probably attracted by chemokines, such as MIG, IP-10 and I-TAC. The chemoattracting process and IL-12 might also involve the NKRP1a-mediated endothelia migration pathway [44–45]. It is also probable that chemokines produced in the blood by activated $V\gamma 2V\delta 2$ T cells themselves or other immune cells initiate the transendothelial migration of these $\gamma\delta$ T cells to the lung from the circulation during systemic infection or immune activation. Abbreviations: IL-12, interleukin-12; IP-10, interferon (IFN)-inducible protein 10; I-TAC, IFN-inducible T-cell α -chemoattractant; MIG, monokine induced by IFN- γ ; MIP-1 α , macrophage inflammatory protein-1 α ; NKRP1a, NK receptor protein 1a (CD161).

Table 1 $\gamma\delta$ TCR genes in rhesus monkeys^a

$\gamma\delta$ TCR genes in rhesus monkeys	% similarity in a.a. sequences to human counterparts	Cross-reactive human TCR antibodies (clone name)
<i>Vδ1</i>	84	δ TCS1; TS8-1E12
<i>Vδ2</i>	87	15D
<i>Vδ3</i>	86	P11.5B
<i>Vδ4</i>	84 (monkey <i>Vδ4</i> is <i>Vα6.2</i>)	
<i>Vδ5</i>	83 (monkey <i>Vδ5</i> is <i>Vα21</i>)	
<i>Dδ^b</i>		
<i>Jδ1</i>	>90 to J1	
<i>Jδ2</i>	>90 to J2	
<i>Jδ3</i>	>90 to J3	
<i>Cδ</i>	86	anti-TCR δ 1
<i>Vγ1.1;1.4</i>	78 (<i>Vγ1.1</i>); 84 (<i>Vγ1.4</i>)	23D12; 4A11(<i>Vγ1.4</i>)
<i>Vγ2 (<i>Vγ9</i>)</i>	91	7A5
<i>VγIII (<i>Vγ10</i>)^c</i>	N/A	
<i>Jγ1</i>	>75 to <i>Jγ1.2</i>	
<i>Jγ2</i>	>90 to <i>Jγ2.3</i>	
<i>Cγ</i>	93	

^a Abbreviations: a.a., amino acid; TCR, T-cell receptor.

^b D and J gene segments are predicted from cDNA sequences.

^c *V γ 10* is expressed only in non-human primates; it is inactivated in humans.

Table 2Comparative biological features of V γ 2V δ 2, V δ 1–IELs and $\alpha\beta$ T cells

	V γ 2V δ 2 T cells (in the circulation)	Human V δ 1 or murine $\gamma\delta$ IEL ^a	$\alpha\beta$ T cells	Refs
Development	Thymus	Thymus or extra-thymus	Thymus	[1,59]
TCR repertoire	Diverse	Not diverse for DECTs	Diverse	[1,2,15]
CD3 complex	No CD3 δ for $\gamma\delta$ PBL			[46]
Phenotypes	Similar to $\alpha\beta$ T cells CD5 ⁺ , CD28 ⁺ , CD57 ⁻	Different from $\alpha\beta$ T cells CD5 ⁻ , CD28 ⁻ , CD57 ⁺	Typical T cells	[2,59,60]
Chemokine receptors (difference in expression)	CCR5 ⁺ , CXCR3 ⁺	CCR5 ⁻ , CXCR1 ⁺	CCR5 ^{-/+}	[41,42]
Migration-related molecules	CD161 ⁺ , CD31 ⁻	CD161 ⁻ , CD31 ⁺	CD161 \pm , CD31 \pm	[44,45]
Innate function		Naïve expression of CTL-related genes and others		[47,48]
Production of cytokines or chemokines on activation	IFN- γ ; TNF- α ; RANTES; MIP-1 α and β	Epithelial growth factor	IFN- γ , TNF- α , RANTES, MIP-1 α and β	[38,42,53,54]
Antigen recognition	Non-peptide phosphoantigens from bacteria and plants	Self-reactive; CD1, MICA; ICB; TL	Peptide or lipid antigens	[4–7,60,61]
Antigen presentation	MHC-independent, antigen processing not required		MHC- or CD1-restricted, antigen processing required	[4]
Precursor frequency for a single antigen	0.5–5/100 CD3 ⁺ T cells		1/2 \times 10 ⁵ CD8 ⁺ T cells	[62]
Immune responses	Adaptive responses	No evidence of adaptive responses	Adaptive responses	[15,63]
Expansion magnitude in bacterial infections	30–98% of CD3 ⁺ T cells		0.1–0.5% for Ag85B-specific CD4 ⁺ T cells	[15,21]
Expansion duration after resolution of infection	Can be longer than 7–12 months		Transient	[15,40]
CTL effector function	Yes		Yes	[1,51,52]
Non-immune function	?	Wound repair by murine DETCs		[2]

^a Abbreviations: CTL, cytotoxic T lymphocyte; CCR5, chemokine C receptor 5; DETCs, dendritic epidermal T cells; IELs, intraepithelial lymphocytes; IFN- γ , interferon- γ ; MICA, stress-inducible MHC class I-related chains A; MIP, macrophage inflammation protein; PBL, peripheral-blood leukocytes; TCR, T-cell receptor; TL, thymus leukemia antigen; TNF- α , tumor necrosis factor- α .