

$\gamma\delta$ T-cells: basic features and potential role in vasculitis

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ABSTRACT

$\gamma\delta$ T-cells are a numerically small subset of T-cells with distinct features. They recognise antigens that are not seen by other immune cells. At the functional level, $\gamma\delta$ T-cells share some features with $\alpha\beta$ T-cells but also exert functions that are otherwise performed by specialised subsets of $\alpha\beta$ T-cells (e.g. IL-17 production, regulatory activity). We discuss the potential role of $\gamma\delta$ T-cells in various clinical forms of vasculitis.

Introduction

Most peripheral T-lymphocytes express a disulphide-bridged $\alpha\beta$ T-cell receptor (TCR) heterodimer. $\alpha\beta$ T-cells recognise short peptides presented on MHC class I molecules (to CD8 T-cells) or MHC class II molecules (to CD4 T-cells). Due to the large number of available variable (V) and joining (J) gene segments in the TCR α and β loci, the expressed TCR repertoire of $\alpha\beta$ T-cells is very diverse. A second T-cell subset expresses the alternative $\gamma\delta$ TCR, which is composed of a γ and δ chain heterodimer. In striking contrast to $\alpha\beta$ T-cells, the number of available V γ and V δ genes is small. $\gamma\delta$ T-cells account for approximately 5% of T-cells in the peripheral blood but constitute a major population in other anatomical localisations such as the small intestine and some epithelia (1,2). Two reasons might be envisioned why nature has afforded to maintain two separate sets of T-cells throughout evolution. First, the two T-cell subsets might be endowed with distinct functional capacities. Secondly, the TCR of the two subsets might be destined to recognise separate ranges of antigenic moieties. Accumulated evidence would speak in favour of the latter alternative. Over the years, many studies have analysed the functional activity of murine and human $\gamma\delta$ T-cells.

Depending on the cellular context and the activation signals, $\gamma\delta$ T-cells can produce a range of cytokines including interferon- γ (IFN- γ), tumour necrosis factor- α (TNF- α) and many chemokines (1,3-6). $\gamma\delta$ T-cells freshly isolated from human peripheral blood tend to be primed toward a Th1 phenotype but can be induced under appropriate *in vitro* culture conditions to produce the Th2 cytokine IL-4 (7). Strikingly, $\gamma\delta$ T-cells are poor IL-2 producers, which explains the T helper cell-dependency of $\gamma\delta$ T-cell activation *in vitro* (8). $\gamma\delta$ T-cells localised in specialised areas such as the mucosal tissues, epithelia and epidermis can produce additional cytokines with relevance for the local immune response. Thus, intraepithelial and epidermal $\gamma\delta$ T-cells contribute to epithelial cell growth and wound repair through the production of keratinocyte growth factor/fibroblast growth factor-7 and insulin-like growth factor-1 (9-11). Recently, $\gamma\delta$ T-cells have been recognised as an important cellular source of the pro-inflammatory cytokine IL-17 (see below). In addition to the production of soluble mediators, $\gamma\delta$ T-cells usually display potent cytotoxic effector activity. $\gamma\delta$ T-cells can kill susceptible target cells by the secretory pathway through the production of perforin and granzymes, but can also make use of the receptor-ligand (e.g. Fas/Fas-ligand) pathway (12-14). Taken together, it appears that the effector functions of $\gamma\delta$ T-cells are not fundamentally different from $\alpha\beta$ T-cells, with the possible exception of the production of particular mediators in selected areas of the body.

Ligands recognised by $\gamma\delta$ T-cells

It is by now well documented that the TCR of a particular subset of human $\gamma\delta$ T-cells recognises antigens that are not seen by any other immune receptor.

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The few expressed $V\gamma$ and $V\delta$ genes are not randomly used by human $\gamma\delta$ T-cells. Instead, there is a clear preponderance of cells expressing one particular $V\gamma V\delta$ combination, *i.e.* $V\gamma 9$ and $V\delta 2$, among circulating $\gamma\delta$ T-cells in the human blood. $V\gamma 9V\delta 2$ T-cells account for 50 to >90% of peripheral blood $\gamma\delta$ T-cells in most healthy adult individuals (15, 16). It has been shown that the $V\gamma 9V\delta 2$ TCR recognises non-peptidic phosphorylated molecules that are intermediates of the isoprenoid biosynthesis pathway required for the prenylation of proteins and the synthesis of cholesterol (17). There are two separate pathways leading to the synthesis of the isoprenoid metabolite isopentenyl pyrophosphate (IPP): While eukaryotic cells use the mevalonate-dependent pathway, most prokaryotic cells synthesise IPP *via* the non-mevalonate (or "Rohmer") pathway (18). IPP is recognised by the $V\gamma 9V\delta 2$ TCR but high concentrations (in the micromolar range) are required; while non-transformed normal cells produce too little IPP to activate $\gamma\delta$ T-cells, transformed tumour cells produce much higher amounts of IPP, which can then be sensed by $V\gamma 9V\delta 2$ T-cells, rendering many tumour cells susceptible to $\gamma\delta$ T-cell-mediated lysis (19). Interestingly, the intracellular levels of IPP production in eukaryotic cells can be easily modulated by clinically used drugs, *i.e.* aminobisphosphonates. Aminobisphosphonates inhibit the IPP-degrading enzyme farnesyl diphosphate synthase, resulting in the intracellular accumulation of IPP (19, 20). As a consequence, treatment with aminobisphosphonates or siRNA-mediated inhibition of farnesyl diphosphate synthase increases the susceptibility of tumour cells to $\gamma\delta$ T-cell-dependent lysis (21). In contrast to eukaryotic cells, bacteria and some parasites use the non-mevalonate pathway for the synthesis of IPP. Interestingly, intermediates of this pathway upstream of IPP have been found to be the most potent TCR-dependent activators of human $V\gamma 9V\delta 2$ T-cells. For example, only pico- to nanomolar concentrations of (E)-1-hydroxy-2-methylbut-2-enyl 4-diphosphate (HMB-PP) are required to stimulate human $V\gamma 9V\delta 2$ T-cells (22, 23). The recognition of a unique class of

pyrophosphate ligands that are secreted by bacteria or are overproduced by (some) malignant cells assigns an important function in immune surveillance to the human $V\gamma 9V\delta 2$ T-cell subset. The very same T-cell population thus plays an important role in immune defense against infection and in the elimination of transformed cells through the recognition of related molecules (17, 24, 25). In contrast to the peptide presentation to $CD4^+$ and $CD8^+$ $\alpha\beta$ T-cells, the presentation of such phosphoantigens is MHC-independent. As a consequence, the activation of $V\gamma 9V\delta 2$ T-cells is not MHC-restricted (1, 26). Moreover, all T-cells displaying the $V\gamma 9V\delta 2$ TCR (*i.e.* on average 2-5% of all peripheral blood T-cells) are activated by phosphoantigens, independently of CDR3 sequence variability. Thus, $V\gamma 9V\delta 2$ T-cells use their TCR as a pattern recognition receptor, comparable to well characterised pattern recognition receptors such as the Toll-like receptor family (25). This feature, together with the rapid and sustained activation of effector functions (27, 28), places $\gamma\delta$ T-cells as a potential link between the innate and adaptive immune system (29).

In view of the exclusive features of human $V\gamma 9V\delta 2$ $\gamma\delta$ T-cells, it is surprising to note that there is no homologous TCR in the murine $\gamma\delta$ TCR repertoire that would enable mouse $\gamma\delta$ T-cells to recognise microbial or eukaryotic phosphoantigens. However, the recognition of stress-induced ligands has emerged as a common theme of both murine and human $\gamma\delta$ T-cells. MHC class I-related molecules that are up-regulated upon cellular stress (*e.g.* heat-shock, DNA damage, infection), or are constitutively expressed on some normal and/or on transformed cells, have been shown to be recognised by specified murine or human $\gamma\delta$ TCRs. For instance, murine non-classical MHC class I molecules with constitutive (T22) or inducible expression (T10) have been identified as ligands for the murine G8 $\gamma\delta$ T-cell hybridoma, but some mouse $\gamma\delta$ T-cells can also recognise peptide-MHC class II I-E complexes (30). In humans, $\gamma\delta$ T-cells localised in mucosal tissue and expressing the TCR $V\delta 1$ chain (in contrast to the above mentioned blood $V\gamma 9V\delta 2$

T-cells), have been described to recognise stress-inducible MHC class I-related chain A (MICA) antigens that are inducibly expressed on epithelial cells and frequently constitutively expressed on tumour cells (31). Taken together, it seems reasonable to assume that $\gamma\delta$ T-cells use their TCR to control the integrity of tissue and to sense danger signals as generated by infection, inflammation, and cellular transformation.

Like conventional $\alpha\beta$ T-cells, $\gamma\delta$ T-cells express additional cell surface molecules with regulatory function on T-cell activation. An important activating receptor is the natural killer cell group 2, member D (NKG2D) receptor, a homodimeric type II transmembrane C-type lectin-like receptor which is expressed on all Natural Killer (NK) cells and $\gamma\delta$ T-cells, the majority of $CD8^+$ $\alpha\beta$ T-cells, and a minor population of $CD4^+$ $\alpha\beta$ T-cells with regulatory activity (32). Upon binding of corresponding NKG2D ligands (NKG2DL), human NKG2D transmits an activating signal *via* the associated transmembrane adapter protein DAP10 and the PI3-kinase pathway (33). NKG2D triggers cytolytic effector function in killer cells and co-stimulates cytokine production and proliferation in T-cells (34, 35). The ligands for NKG2D comprise an array of MHC class I-related molecules with two or three MHC class I-like domains. The human NKG2DL include MICA and MICB, and six members of the UL16-binding protein (ULBP) family that are also known as retinoic acid early transcript (RAET) proteins. In mice, the NKG2DL include five retinoic acid early transcript proteins (RAE-1 α - ϵ), three variants of the H60 minor histocompatibility antigen, and the murine UL-16-binding-protein-like transcript-1 (MULT1) (36, 37). With few exceptions, normal cells do not express NKG2D ligands, but expression can be induced in many cells by "stress" including heat shock, DNA damage, viral infection, or cellular activation (36, 37). Furthermore, the NKG2D/NKG2DL system is an integral part of local immunosurveillance, as evidenced in transgenic mouse models with inducible acute up-regulation of NKG2DL (38). Interestingly, human MICA molecules can be sensed

by the V δ 1 TCR (29), but can also directly activate $\gamma\delta$ T-cells *via* NKG2D (39). Integrated signals resulting from TCR-dependent antigen recognition and additional receptor-ligand interactions thus determine the overall $\gamma\delta$ T-cell response (25, 40).

Regulatory functions of $\gamma\delta$ T-cells

There is increasing evidence for multiple interactions between $\alpha\beta$ and $\gamma\delta$ T-cells. Activation of phosphoantigen-reactive human $\gamma\delta$ T-cells is inhibited by conventional TCR $\alpha\beta$ -expressing regulatory T-cells (Treg) (41, 42). Strikingly, however, it was observed that human $\gamma\delta$ T-cells themselves can display regulatory (*i.e.* suppressive) activity on $\alpha\beta$ T-cells (43, 44), and can antagonise the IL-2-driven expansion of Treg in mycobacteria-infected cynomolgus macaques (45). Using highly purified human CD4⁺ $\alpha\beta$ T-cells and $\gamma\delta$ T-cells, we observed under *in vitro* co-culture conditions that upon phosphoantigen stimulation, $\gamma\delta$ T-cells suppressed the expansion of antigen-reactive CD4⁺ T-cells (46). Depending on the experimental conditions, V γ 9V δ 2 T-cells can thus proliferate and differentiate into cytokine-secreting and cytotoxic effector cells in response to phosphoantigens, but can also regulate $\alpha\beta$ T-cell responses. However, $\gamma\delta$ T-cell subsets other than the dominant V γ 9V δ 2 population also seem to be endowed with regulatory functions. V δ 1 T-cells isolated from tumour-infiltrating lymphocytes in breast cancer patients suppress $\alpha\beta$ T-cell responses (47), and increased numbers of circulating V δ 1 T-cells were found to correlate with operational tolerance in liver allograft recipients (48). Similarly, regulation of CD4⁺ $\alpha\beta$ T-cell responses has been described for murine $\gamma\delta$ T-cells (49). Multiple additional evidence for suppressive functions of $\gamma\delta$ T-cells has been accumulated from studies in various mouse models, not the least in $\gamma\delta$ T-cell-deficient mice (50). Thus, it can be concluded that $\gamma\delta$ T-cells are not only capable of recognizing antigens that are not seen by other immune cells (*e.g.* phosphoantigens) but fulfil additional important regulatory functions. Recently, a new facet has been added by the demonstration that $\gamma\delta$ T-cells are

a major and early source of the pro-inflammatory cytokine IL-17. It has been recognised that Th17 cells represent a separate T helper cell lineage (in addition to Th1 and Th2) with an important role in the pathology of organ-specific autoimmune diseases and immunity to infection (51). Moreover, conventional (*i.e.* CD4⁺ TCR $\alpha\beta$ -expressing) Th17 cells are also discussed to contribute to the initiation and perpetuation of the autoimmune response in anti-neutrophil cytoplasmic autoantibody (ANCA)-associated systemic vasculitis (52). In mouse models of autoimmunity and infection, $\gamma\delta$ T-cells were found to share characteristic features with Th17 cells such as the expression of the chemokine receptor CCR6, the IL-23 receptor, the transcription factor retinoic orphan receptor γ t (ROR γ t), and the aryl hydrocarbon receptor, and to provide an early and innate source of IL-17 (53, 54). Taken together, it is thus obvious that $\gamma\delta$ T-cells despite being a numerically minor cell population in most localisations in the body, entertain multiple mechanisms to modulate innate and adaptive immune responses.

Role of $\gamma\delta$ T-cells in vasculitis

In view of the identification of stress-inducible ligands as relevant antigens and the documented regulatory function of $\gamma\delta$ T-cells, it is reasonable to assume that $\gamma\delta$ T-cells contribute to the immune response in vasculitis. So far, however, only few studies have addressed this issue.

Behçet's disease

Behçet's disease (BD) is a multisystemic disease, with vasculitic lesions occurring at multiple sites including eyes, skin, joints and brain. Early studies indicated that peripheral blood $\gamma\delta$ T-cells from BD patients responded specifically to peptides derived from the 65 kDa mycobacterial heat-shock protein (HSP) or the 60 kDa homologous human HSP; the V γ V δ TCR repertoire of the responding $\gamma\delta$ T-cells was not identified in these studies (55). Subsequently, increased proportions of $\gamma\delta$ T-cells in the peripheral blood of BD patients were reported, which displayed signs of activation and strong production of

IFN- γ and TNF- α (56, 57). Although the aetiology of BD is unknown, viral and bacterial infections have been repeatedly implicated. In addition to the described HSP reactivity of $\gamma\delta$ T-cells, proliferative responses of BD patient's $\gamma\delta$ T-cells to bacterial preparations have been reported (58, 59). While no specific bacterial antigen has been identified to date (with the exception of the mycobacterial HSP-65 (52)), V γ 9V δ 2 T-cells from BD patients can be activated (as expected) by microbial or synthetic phosphoantigens. Interestingly, Verjans and co-workers observed a high proportion (ranging from 15 to 29%) of $\gamma\delta$ T-cells among T-cells present in intraocular fluid (but not in the blood) in 3 of 6 affected BD patients (60). Among these $\gamma\delta$ T-cells, proliferation could be induced *in vitro* following stimulation with phosphoantigen IPP but not with recombinant HSP-65, pointing to the presence of V γ 9V δ 2 T-cells among the infiltrated $\gamma\delta$ T-cells (although this was not specifically proven in this study) (60). Furthermore, peripheral blood V γ 9V δ 2 T-cells from patients with active BD were found to secrete increased amounts of the serine protease granzyme A when stimulated with phosphoantigens (61). While the studies on ocular T-cells suggested the presence of V γ 9V δ 2 T-cells (60), no preferential expression of a particular V γ and/or V δ chain was detected in biopsies from ulcerated oral mucosa from BD patients (62). The latter study, however, confirmed the dominance of V γ 9V δ 2 T-cells among $\gamma\delta$ T-cells, similarly to healthy donors, also in BD patients. Furthermore, using the available anti-V γ /V δ mAb, Freysdottir *et al.* also analysed the presence of $\gamma\delta$ T-cells expressing other V γ /V δ elements. These studies revealed alterations in the relative proportion of several $\gamma\delta$ T-cell subsets among BD patients with different clinical conditions (62). In a recent study, Yasuoka and coworkers reported a preferential activation of peripheral blood V δ 1 as compared to V δ 2 $\gamma\delta$ T-cells in patients with active BD (63). In the absence of functional data, however, the significance of such alterations remains unclear. Taken together, it appears that activated $\gamma\delta$ T-cells are detected in the peripheral blood of BD

patients, and that $\gamma\delta$ T-cells are also present locally in the affected tissue, possibly with differential prevalence of $\gamma\delta$ T-cell subsets in oral mucosa *versus* ocular fluid. Further studies are needed to characterise the function of these $\gamma\delta$ T-cells in more detail.

Takayasu's arteritis

Takayasu's arteritis (TA) is a chronic inflammatory panarteritis characterised by intimal thickening, fibrosis and stenosis, presumably resulting from an immune-mediated dysfunction of the arterial endothelium (64). Based on the restricted TCR repertoire, the expression of HSP-65 and MHC class II molecules in the aortic tissue and the frequent infiltration with T-cells, the aetiology is thought to involve aberrant autoimmune responses (65, 66). A large proportion (on average 30%) of leukocytes infiltrating aortic tissue in TA patients are in fact $\gamma\delta$ T-cells (65). Moreover, increased proportions of $\gamma\delta$ T-cells are also present in the blood of patients with active but not with inactive TA (67). Interestingly, the TCR V δ 1 repertoire seems to be skewed toward an over-representation of V δ 1 when compared to healthy control donors (67). So far it is unknown whether the increased numbers of V δ 1 T-cells are functionally related to the pathogenesis and/or immune regulation of TA, *e.g.* by displaying regulatory activity as shown for V δ 1 T-cells in other clinical conditions (47). It was reported that peripheral blood $\gamma\delta$ T-cells from TA patients but not from healthy donors or SLE patients proliferate in response to recombinant human HSP-60 and kill aortic endothelial cells *in vitro* (68). Although not investigated in this study, it is possible that the HSP-60 reactivity resulted from the stimulation of V δ 1 $\gamma\delta$ T-cells, since the dominant V γ 9V δ 2 T-cells do not respond to HSP (69). Taken together, the numerically increased proportion in the blood, the associated alterations in the expressed TCR repertoire, and the significant accumulation at the arterial lesions point to an important role of $\gamma\delta$ T-cells in TA which deserves further investigation.

Wegener's granulomatosis

Wegener's granulomatosis (WG) is an immune-mediated disease with anti-neu-

trophil cytoplasmic antibody (ANCA)-associated systemic vasculitis (70). Clinical and experimental evidence supports a pathogenic role of ANCA in WG (71). Proteinase-3 has been identified as a target for humoral and cellular autoimmune responses in WG. A large range of phenotypic and/or functional alterations in the T-cell compartment has been described, which in essence all deal with $\alpha\beta$ T-cells (52, 72). Very little information regarding $\gamma\delta$ T-cells in WG is available. Using 5 WG patients as a control cohort for their studies on TA, Chauhan *et al.* did not observe appreciable alterations in the peripheral blood $\gamma\delta$ T-cell compartment (67). Furthermore, increased proportions of activated CD8 T-cells co-expressing CD57 have been reported, but the TCR phenotype was not identified in these studies (73). Interestingly, $\gamma\delta$ T-cells are present among the infiltrating cells around tubular and glomerular capillaries in the kidney of WG patients with active disease (74). There is also strong expression of MICA in the peritubular endothelium and in glomerular endothelial cells (74), pointing to a possible cross-talk of infiltrating $\gamma\delta$ T-cells with endothelial cells. Moreover, cluster-like formations of NKG2D⁺ and MIC⁺ cells are present in WG-granulomata from the upper respiratory tract (75). Although available data do not suggest a major role for $\gamma\delta$ T-cells in WG, it might still be informative to investigate their functional capacity in more detail.

Other types of vasculitis

In the leukoclastic form of cutaneous necrotising vasculitis (CNV) with presumed immune complex-mediated pathogenesis, a significant proportion of $\gamma\delta$ T-cells was present in skin lesions as analysed by immunohistochemistry (76). Even higher numbers of $\gamma\delta$ T-cells were detected in specimens from patients with documented infectious aetiology. The functions of the locally accumulated $\gamma\delta$ T-cells are, however, unknown.

Future perspectives for $\gamma\delta$ T-cell research in vasculitis

Apart from Behçet's disease, little attention has been paid to characterise $\gamma\delta$ T-cells in various forms of vasculitis.

In view of the expression of $\gamma\delta$ T-cell ligands in affected tissue (68) and the observed alterations in the $\gamma\delta$ TCR repertoire in some forms of vasculitis (64), it seems worthwhile to investigate the functions of $\gamma\delta$ T-cells more extensively in such diseases. This might include the regulatory/suppressive capacity of isolated V δ 1 $\gamma\delta$ T-cells in Takayasu's arteritis as well as the potential of $\gamma\delta$ T-cells for IL-17 production in Wegener's granulomatosis and Behçet's disease.

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