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## Direct presentation of non-peptide prenyl pyrophosphate antigens to human $\gamma\delta$ T cells

C.T. Morita<sup>(1)</sup>, Y. Tanaka<sup>(2)</sup>, B.R. Bloom<sup>(2)</sup>, and M.B. Brenner<sup>(1)</sup>

<sup>(1)</sup> *Lymphocyte Biology Section, Division of Rheumatology and Immunology, Department of Medicine, Brigham and Women's Hospital and Harvard Medical School, and*

<sup>(2)</sup> *Howard Hughes Medical Institute and Department of Microbiology, Albert Einstein College of Medicine, Bronx, New York 10461 (USA)*

### Introduction

T cells can be divided into two subsets based on their expression of  $\alpha\beta$  or  $\gamma\delta$  T-cell antigen receptors.

Although  $\gamma\delta$  T cells were discovered 10 years ago (Brenner *et al.*, 1986), relatively little is known about the types of antigens that stimulate  $\gamma\delta$  T cells or the presentation elements and presentation path-

ways that these antigens use. Few  $\gamma\delta$  T cells respond to conventional peptide antigens or superantigens presented by MHC class I or class II molecules that stimulate  $\alpha\beta$  T cells (Haas *et al.*, 1993; Porcelli *et al.*, 1991).

There were many indications that antigen recognition by  $\gamma\delta$  T cells might be very different from antigen recognition by  $\alpha\beta$  T cells. Although human  $\alpha\beta$  and  $\gamma\delta$  T cells share a similar structure for their antigen receptors, a variable heterodimer that associates with the invariant CD3 signalling complex, there are greater than 50 V $\alpha$  and 46 V $\beta$  gene segments that can pair to make several thousand receptor combinations. Any one pair constitutes only a small percentage of the blood T cells. In stark contrast, there are only 6 V $\gamma$  and 3-4 major V $\delta$  gene segments used by human  $\gamma\delta$  T cells (Porcelli *et al.*, 1991). Moreover, in normal adults, one of these pairs, the V $\gamma$ 2V $\delta$ 2 T-cell receptor, predominates, being expressed on approximately 50-75% of the  $\gamma\delta$  T cells. Thus, this one TCR pair is present on 3-5% of human T cells (Morita *et al.*, 1994; Parker *et al.*, 1990). A similar restricted repertoire of V $\gamma$  and V $\delta$  gene segments is seen in rodents (Haas *et al.*, 1993).

$\alpha\beta$  and  $\gamma\delta$  T cells also differ in their functional capabilities. In general,  $\alpha\beta$  T cells can be divided into helper cells that express CD4 or killer cells that express CD8. In contrast, the majority of  $\gamma\delta$  T cells are killer cells that express neither CD4 nor CD8 (Groh *et al.*, 1989; Morita *et al.*, 1991). Like  $\alpha\beta$  T cells,  $\gamma\delta$  T cells secrete a variety of cytokines, although they make somewhat lesser amounts of IL2 and GM-CSF (Morita *et al.*, 1991; Patel *et al.*, 1989; Spits *et al.*, 1991). Thus, given the differences in the TCR repertoire and functional capabilities, antigen recognition by  $\gamma\delta$  T cells might be distinct from antigen recognition by  $\alpha\beta$  T cells.

Recently, we have identified non-peptide prenyl pyrophosphate antigens that stimulate the major subset of human  $\gamma\delta$  T cells (Tanaka *et al.*, 1995). This has allowed us to study the presentation pathways used by these novel antigens. We find that  $\gamma\delta$  T cells recognize an entirely new class of antigens through an extracellular presentation pathway that does not require antigen uptake, antigen processing, or known antigen-presenting elements (Morita *et al.*, 1995).

### Human $\gamma\delta$ T cells and infections

One of the first suggestions that human  $\gamma\delta$  T cells could recognize foreign antigens was from studies examining the development of the  $\gamma\delta$  T-cell repertoire (Parker *et al.*, 1990).  $\gamma\delta$  T-cell numbers in the peripheral blood increase between ages

1-10. This peripheral expansion of  $\gamma\delta$  T cells is restricted to the V $\gamma$ 2V $\delta$ 2 subset and leads to the predominance of this subset in the  $\gamma\delta$  T-cell repertoire of most adults and the high level of V $\gamma$ 2V $\delta$ 2<sup>+</sup> T cells found among peripheral blood T cells. Identical twins were found that were disparate for the level of expression of the V $\gamma$ 2V $\delta$ 2 T-cell subset in that one twin had high levels of V $\gamma$ 2V $\delta$ 2 T cells while the other did not. As identical twins are genetically identical, environmental factors appear to influence the development of the  $\gamma\delta$  T-cell repertoire.

Infections with microorganisms are likely candidates for environmental factors that influence  $\gamma\delta$  T-cell development.  $\gamma\delta$  T cells were found to increase dramatically in the peripheral blood in response to a variety of infections, both bacterial (Balbi *et al.*, 1993; Bertotto *et al.*, 1993; Caldwell *et al.*, 1995; Hara *et al.*, 1992; Sumida *et al.*, 1992; Ueta *et al.*, 1994) and protozoal (Ho *et al.*, 1990; Perera *et al.*, 1994; Scalise *et al.*, 1992) (table I). In patients with infections with *Ehrlichia*, a rickettsial organism, extremely high levels of  $\gamma\delta$  T cells were observed (up to 97% of peripheral blood T cells) (Caldwell *et al.*, 1995). Moreover, these  $\gamma\delta$  expansions are limited to cells expressing the major V $\gamma$ 2V $\delta$ 2 T-cell receptor.

These *in vivo* expansions of  $\gamma\delta$  T cells can be reproduced *in vitro* by exposure of normal peripheral blood mononuclear cells to extracts from mycobacteria (De Libero *et al.*, 1991; Kabelitz *et al.*, 1990; Panchamoorthy *et al.*, 1991). Like the *in vivo* expansions, the *in vitro* expansions were limited to cells bearing the V $\gamma$ 2V $\delta$ 2 T-cell receptor (De Libero *et al.*, 1991; Kabelitz *et al.*, 1991). Upon sequencing of the V $\gamma$ 2V $\delta$ 2 T-cell antigen receptors, these expanded populations were found to be polyclonal (Ohmen *et al.*, 1991; Panchamoorthy *et al.*, 1991). This suggested that germ-line-encoded segments of the T-cell receptor were involved in the recognition of soluble antigens from mycobacteria.

$\gamma\delta$  T-cell clones expressing the V $\gamma$ 2V $\delta$ 2 T-cell receptor also responded to soluble extracts from *Mycobacterium tuberculosis in vitro* (De Libero *et al.*, 1991; Holoshitz *et al.*, 1989). This allowed us to test whether one  $\gamma\delta$  T cell could respond to different pathogens or whether distinct populations of V $\gamma$ 2V $\delta$ 2 T cells were responding to different pathogens. V $\gamma$ 2V $\delta$ 2 T-cell clones responded to extracts from a number of different bacteria including mycobacteria, Gram-positive and -negative rods and Gram-positive cocci (data not shown). Therefore, individual V $\gamma$ 2V $\delta$ 2 T cells can respond to a number of different microorganisms, suggesting that V $\gamma$ 2V $\delta$ 2 T cells may recognize common structural features of antigens from the microorganisms.

**Table I.** Expansion of human  $\gamma\delta$  T cells in response to infection.

Disease	$\gamma\delta$ T cells (mean % of CD3 <sup>+</sup> )		Reference	
	Normals	Patients (max)		
Bacterial infections	Salmonellosis	5	18 (48)	(Hara <i>et al.</i> , 1992)
	Tularaemia	3	33 (33)	(Sumida <i>et al.</i> , 1992)
	Brucellosis	4	29 (48)	(Bertotto <i>et al.</i> , 1993)
	Tuberculosis	6	14 (35)	(Balbi <i>et al.</i> , 1993)
	Tuberculosis contacts	5	10 (18)	(Ueta <i>et al.</i> , 1994)
	Ehrlichiosis	5	57 (97)	(Caldwell <i>et al.</i> , 1995)
Protozoal infections	Visceral leishmaniasis	8	44	(Raziuddin <i>et al.</i> , 1992)
	Acute malaria	4	16 (27)	(Ho <i>et al.</i> , 1990)
	Malarial paroxysms	4	11 (27)	(Perera <i>et al.</i> , 1994)
	Toxoplasmosis	4	9 (15)	(Scalise <i>et al.</i> , 1992)

#### Isopentenyl pyrophosphate and related non-peptide compounds from mycobacteria are antigens for human V $\gamma$ 2V $\delta$ 2<sup>+</sup> T cells

To help determine the molecular basis for the recognition of mycobacterial antigens, we isolated and identified two antigens from mycobacteria that stimulate  $\gamma\delta$  T cells. Preliminary studies by Pfeffer *et al.* demonstrated that these antigens were protease-resistant and extremely small, suggesting that the antigens for  $\gamma\delta$  T cells are non-proteinaceous compounds (Pfeffer *et al.*, 1990; 1992). Subsequently, these antigens were shown to contain critical phosphate residues on the basis of their sensitivity to alkaline phosphatase and nucleotide pyrophosphatase (Constant *et al.*, 1994; Schoel *et al.*, 1994; Tanaka *et al.*, 1994).

Based on these and other experiments, we devised a purification protocol to isolate these antigens from *Mycobacteria smegmatis*. Two antigens were purified and identified by their fragmentation spectra on tandem mass spectroscopy (Tanaka *et al.*, 1995). Isopentenyl pyrophosphate (fig. 1) and its related hydroxymethyl derivative were able to stimulate V $\gamma$ 2V $\delta$ 2 T-cell clones and to expand the V $\gamma$ 2V $\delta$ 2 subset from peripheral blood cells. These are among the first, natural, non-peptide antigens described for T cells and constitute a new class of antigens distinct from peptide antigens, superantigens, and the recently described lipid antigens.

Isopentenyl pyrophosphate has two major structural features; a carbon isoprenyl chain and a pyrophosphate moiety. Studies with structural analogs of isopentenyl pyrophosphate demonstrate that V $\gamma$ 2V $\delta$ 2 T cells show specificity in their recognition

of phosphate-containing compounds and can only recognize non-bulky carbon chains that do not contain amino groups (Bürk *et al.*, 1995; Tanaka *et al.*, 1995; 1994) and data not shown). Similarly, changes in the pyrophosphate moiety also affect recognition of isopentenyl pyrophosphate. Multiple substitutions of the phosphate group were not permitted. Removal of a phosphate group from isopentenyl pyrophosphate decreased specific biological activity by 230-fold. The addition of a uridine triphosphate molecule had minimal effects on recognition.

The addition of a nucleotide triphosphate group was suggested from the results of Constant *et al.* (Constant *et al.*, 1994) who have described a TTP-containing antigen (TUBag4) for V $\gamma$ 2V $\delta$ 2 T cells from the cytoplasm of *M. tuberculosis*, although they have not as yet determined the structure of the carbon chain. They also find two pyrophosphate-containing compounds, TUBag1 and TUBag2, but have not yet determined the structure of the carbon chain. Thus, there may be a number of pyrophosphate-containing compounds produced by different microorganisms.

The identification of isopentenyl pyrophosphate and related compounds as natural antigens from mycobacteria for V $\gamma$ 2V $\delta$ 2<sup>+</sup> T cells suggests an explanation as to why  $\gamma\delta$  T cells respond to so many different types of pathogens. Isopentenyl pyrophosphate is an essential precursor for a variety of biological molecules such as cholesterol and its derivatives, vitamins, dolicol phosphates and ubiquinones, and is required for the membrane anchoring of a number of important signal transduction molecules such as ras. As both prokaryotic and eukaryotic

### Isopentenyl pyrophosphate

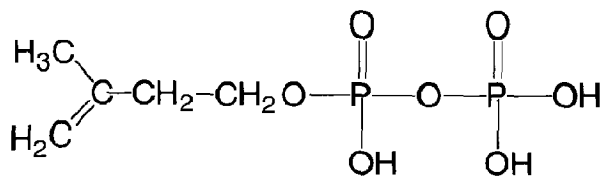


Fig. 1. Structure of isopentenyl pyrophosphate.

There are two main structural features, the isopentenyl carbon chain (on left) and the pyrophosphate moiety (on right).

### Extracellular Presentation Pathway

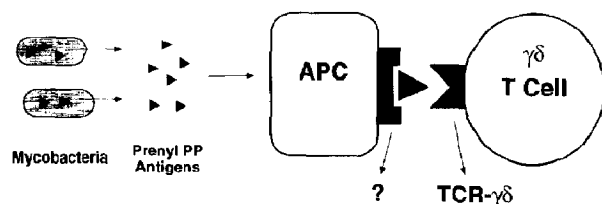


Fig. 2. Proposed extracellular presentation pathway for the prenyl pyrophosphate antigens.

organisms use isopentenyl pyrophosphate and related compounds as essential precursors,  $V\gamma 2V\delta 2$ -bearing T cells can respond to infection by both prokaryotic and eukaryotic pathogens. Given the variety of different microorganisms that stimulate  $V\gamma 2V\delta 2$ -bearing T cells, the number of such prenyl pyrophosphate derivatives recognized by  $\gamma\delta$  T cells may be expected to grow in coming years.

### Extracellular presentation pathway for the prenyl pyrophosphate antigens

Our identification of prenyl pyrophosphate compounds as antigens for  $V\gamma 2V\delta 2$  T cells allowed us to examine the characteristics of the presentation of these antigens and to investigate the involvement of known antigen presenting molecules in recognition (Morita *et al.*, 1995).

Recognition of the non-peptide prenyl pyrophosphate antigens is restricted to  $\gamma\delta$  T-cell clones expressing  $V\gamma 2V\delta 2$  T-cell receptors (Davodeau *et al.*, 1993; Tanaka *et al.*, 1994). This recognition is TCR-dependent as cDNAs encoding  $V\gamma 2$  and  $V\delta 2$  chains confer non-peptide antigen recognition when

transfected into a TCR-negative Jurkat T-cell tumour mutant line (Bukowski *et al.*, 1995). The  $V\gamma 2V\delta 2$  transfectant, DBS43, released IL2 in response to isopentenyl pyrophosphate but not in response to the biologically inactive compound, phenethylpyrophosphate (Tanaka *et al.*, 1995). A control  $V\gamma 1V\delta 1$  TCR transfectant did not respond to either compound, although it did respond to mAbs directed to the TCR. These results establish the TCR dependency of the response to the prenyl pyrophosphate antigens.

Presentation of the prenyl pyrophosphate antigens has unique features in comparison to the presentation of protein antigens (Morita *et al.*, 1995). Mycobacteria are able to secrete non-peptide antigens, suggesting that  $\gamma\delta$  T cells may encounter free antigen *in vivo*. Professional antigen-presenting cells are not required for the presentation of prenyl pyrophosphate antigens (Morita *et al.*, 1995; Vila *et al.*, 1995). The majority of  $V\gamma 2V\delta 2$ -bearing T-cell clones and the  $V\gamma 2V\delta 2$  TCR transfectant could respond to non-peptide antigens in the absence of additional APCs. Moreover, recognition of non-peptide antigens is rapid, occurring as soon as 2-3 min. after their addition (Morita *et al.*, 1995). This suggests that antigen uptake and processing are not required and consistent with this hypothesis, both non-fixed and fixed APC can present non-peptide prenyl pyrophosphate antigens to an APC-dependent  $\gamma\delta$  T-cell clone (Fisch *et al.*, 1990; Morita *et al.*, 1995). Thus, fixed APC which are unable to take up antigens or process them could still present the prenyl pyrophosphate antigens.

The presentation of the prenyl pyrophosphate antigens most resembles the presentation of preprocessed peptide antigens to human  $\alpha\beta$  T cells. Peptide antigens can stimulate most reactive  $\alpha\beta$  T cells in the absence of APC and antigen uptake and processing because the peptides can bind directly to MHC class I or class II molecules on the T-cell surface. The resulting MHC/peptide complexes are then presented to neighbouring T cells (Pichler and Wyss-Coray, 1994). This presentation of peptides by one T cell to another requires cell-cell contact. A similar cell-cell contact requirement is observed for the recognition of the prenyl pyrophosphate antigens, as was determined by increases in calcium flux (Morita *et al.*, 1995) and in  $TNF\alpha$  production (Lang *et al.*, 1995) in response to exposure to the non-peptide antigens.

Even though the presentation of the prenyl pyrophosphate antigens required cell-cell contact, known antigen-presenting molecules were not required for their recognition (Fisch *et al.*, 1990; Holoshitz *et al.*, 1993; Morita *et al.*, 1995). Monoclonal antibodies directed to class II, class I, and CD1 molecules did not block the recognition of the prenyl pyrophosphate antigens (Morita *et al.*, 1995). Moreover, the

V $\gamma$ 2V $\delta$ 2 TCR transfectant lacked MHC class II expression but was able to respond to prenyl pyrophosphate antigens in the absence of APC (Morita *et al.*, 1995). Lastly, mutant APC lacking classical class I,  $\beta_2m$ , or class II molecules were as efficient as wild type APC in presenting mycobacterial non-peptide antigens to an APC-dependent CD4<sup>+</sup>  $\gamma\delta$  T-cell clone and class II-negative APC could function as targets for  $\gamma\delta$  T cells in the presence of antigens from *M. tuberculosis* (Morita *et al.*, 1995). Therefore, the presentation of the prenyl pyrophosphate antigens does not involve known antigen-presenting molecules.

It is not clear if a novel antigen-presenting element exists for the non-peptide prenyl pyrophosphate antigens. Unlike most peptide antigens and superantigens, culturing antigen-presenting cells with the prenyl pyrophosphate antigens does not render the "pulsed" antigen presenting cells stimulatory for V $\gamma$ 2V $\delta$ 2 bearing T cells (Lang *et al.*, 1995; Morita *et al.*, 1995). One interpretation of this result is that non-peptide antigens do not stably associate with the antigen-presenting element on the surface of the APC. A similar lack of stable association between peptides and the murine MHC class II allele, I-A<sup>g7</sup>, has been reported whereby immunogenic peptides cannot be pulsed onto the APC surface although the peptides stimulate I-A<sup>g7</sup>-restricted T cells when cultured together (Carrasco-Marin *et al.*, 1996). The putative presenting element would be predicted to be non-polymorphic and to be widely expressed on many types of cells.

Thus, we propose the following model for the extracellular presentation of prenyl pyrophosphate antigens. The natural antigens for V $\gamma$ 2V $\delta$ 2-bearing T cells are isopentenyl pyrophosphate and related prenyl pyrophosphate compounds. These compounds are synthesized by mycobacteria and a variety of other bacteria, as they are essential metabolites for all living things. However, mycobacteria can secrete these compounds extracellularly where they can associate with the surface of an antigen-presenting cell, perhaps by binding to a novel antigen-presenting element. The T-cell receptor of V $\gamma$ 2V $\delta$ 2 T cells can recognize these compounds. The T cells then become activated to secrete cytokines, proliferate, and kill neighbouring cells. This process requires T cell/APC cell-cell contact but does not require that the APC be a professional APC, nor does it require antigen uptake or processing. In this respect, the presentation of the non-peptide prenyl pyrophosphate antigens is quite distinct from the presentation of conventional protein antigens and superantigens by MHC molecules or lipid antigens by CD1 molecules.

This novel extracellular pathway allows for the direct and rapid recognition of prenyl pyrophosphate antigens by  $\gamma\delta$  T cells. This suggests that  $\gamma\delta$  T-cell

recognition non-peptide antigen might complement  $\alpha\beta$  T-cell antigen recognition of protein antigens that are presented by intracellular pathways. In this way,  $\gamma\delta$  T cells might function as a bridge between the innate immunity of myeloid cells, NK cells and complement and the adaptive immunity of  $\alpha\beta$  T cells and B cells. By recognizing a new class of non-peptide antigens,  $\gamma\delta$  T cells probably play important and unique roles in human immunity and autoimmunity.

### Acknowledgments

This work was supported by grants from the NIH, the Arthritis Foundation (American College of Rheumatology), and the Howard Hughes Medical Institute.

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*Note added in proofs:*

#### Response of Kasahara *et al.*

We would like to thank Drs. Ahouse and Simister for writing the commentary on our article on the original evolution of the class I genes. We would like to address some of the issues they raised and clarify potential misunderstandings.

First, Ahouse and Simister suggest that structural variations among class I heavy chains are likely to be constrained by the need to always interact with the same partner,  $\beta_2m$ . We agree with this notion. However, we do not agree with the idea that class II molecules can vary more easily than class I molecules because they do need to interact with  $\beta_2m$ . Germain and colleagues (1985) showed that even minor allelic variations can affect the pairing of class II A and B chains. This is in contrast to the fact that the same  $\beta_2m$  molecule can form a heterodimer with structurally divergent heavy chains ranging from those of class Ia to even CD1 molecules. Thus,  $\beta_2m$  is less discriminating than class II A or B chains in choosing its partner, and we believe that, for this reason, class I genes can be more plastic than class II genes.

Second, we are not proposing the existence of a genetic mechanism that selectively translocates

class I genes equipped with specialized functions to the outside of the MHC. Because class I genes located outside the MHC class I region are likely to enjoy a longer life span than their MHC-encoded counterparts, they have a better chance of acquiring non-typical functions. We think, therefore, that class I molecules encoded outside the MHC presumably acquired specialized functions as a *consequence* of the translocation.

Finally, the very recent structural analysis of the HSP70 peptide-binding region (Zhu *et al.*, 1996) makes it difficult to justify an evolutionary relationship between class I and HSP70. In order to maintain this hypothesis, one must propose intermediate stages between HSP70 and class I; while these stages may exist (or have existed), we must admit that a gradual evolution of class I from an HSP70 ancestor is not in the spirit of our original model (Flajnik *et al.*, 1991b). Of course the HSP70 structure does not rule out the possibility that it is a fore-runner of class I, but the same can be said of other gene families as well — e.g., see Hashimoto and Kurosawa (1991) for a discussion of immunoglobulin domains as MHC ancestors. As suggested by Ahouse and Simister, molecules ancestral to class I may be present in jawless fish and protochordates, taxa in which no evidence exists for *bona fide* MHC. We (and others) will continue to mine for such gems.

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Key-words: MHC, CD1, Antigen presentation; Non-classical MHC, MHC class Ib, MHC evolution; Forum.