Human $\gamma\delta$ T-cell recognition of Yersinia enterocolitica

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SUMMARY

We have studied the human $\gamma\delta$ T-cell response to Yersinia enterocolitica, a facultative intracellular bacterium which causes gastroenteritis and, particularly in human leucocyte antigen (HLA)-B27⁺ individuals, reactive arthritis (ReA). A marked proliferation of that cytotoxic $\gamma\delta$ T cells is seen when Yersinia-infected lymphoblastoid cell lines or fixed intact Yersinia are added to cultures of mononuclear cells derived from the synovial fluid of ReA patients or from the peripheral blood of healthy donors. In contrast, heat-inactivated Yersinia fail to stimulate the $\gamma\delta$ T-cell response. The $\gamma\delta$ T-cell lines generated killed both autologous and allogeneic infected cell lines. Interestingly, a T-cell line generated from synovial fluid mononuclear cells (SFMC) killed infected autologous cell lines and a cell line matched for HLA-B27 less well than infected allogeneic target cells. $\gamma\delta$ T-cell clones isolated from this line were found to express $V\gamma9V\delta2$ T-cell receptor (TCR) and also killed infected mismatched cells more efficiently than autologous targets. Moreover, from experiments using major histocompatability complex (MHC)-deficient cell lines, it was apparent that target cell recognition was MHC independent. Our results suggest that $\gamma\delta$ T cells can be involved in immunity to Yersinia enterocolitica and should be taken into account when considering immunopathological mechanisms leading to reactive arthritis.

INTRODUCTION

A minor T-cell subset can be defined by the expression of the $\gamma\delta$ T-cell receptor (TCR). Although this receptor has structural similarities to the $\alpha\beta$ TCR, it is becoming clear that antigen recognition by $\gamma\delta$ T cells can be quite different from that exhibited by $\alpha\beta$ T cells.¹ Thus, while most $\alpha\beta$ T cells express either CD4 or CD8 and recognize peptidic antigen presented by an appropriate major histopcompatability complex (MHC) molecule, $\gamma\delta$ T cells usually lack CD4 or CD8 and recognize antigens independently of classic MHC molecules. Indeed, experiments with murine $\gamma\delta$ T cells have suggested that the $\gamma\delta$ TCR may function in an analogous fashion to immunoglobulin, recognizing intact polypeptide rather than small peptide fragments.^{2,3} Moreover, work with human $\gamma\delta$ T cells has revealed that antigens need not be derived from polypeptides; for example, mycobacteria-reactive $\gamma \delta$ T cells can recognize isopentenyl pyrophosphate and related prenyl pyrophosphate derivatives.⁴⁻⁶ The recognition of these non-peptidic ligands evidently proceeds via a novel MHC-independent extracellular pathway.⁷ However, other human $\gamma \delta$ T cells require the presence of members of the non-classic MHC-related molecules CD1,⁸⁻¹⁰ of which CD1b is known to present non-peptidic antigens such as lipoglycans and mycolic acid.¹¹⁻¹³

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Despite the delineation of some of the antigens recognized by $\gamma\delta$ T cells, the biological role of this T-cell subset is still under investigation. There is a consensus that $\gamma\delta$ T cells are involved in the immune response to infections.^{14,15} However, the precise function of the $\gamma\delta$ T-cell response is not clear. Experiments with gene knockout mice have indicated that $\gamma\delta$ T cells may complement $\alpha\beta$ T cells resulting in the efficient clearance of infective organisms.¹⁶ It has been suggested that the $\gamma\delta$ T cells act as a first line of defence linking the innate and the acquired immune responses.¹⁷ However, experiments in which $\gamma \delta$ deficient mice die when challenged with Mycobacterium tuberculosis indicate that $\gamma\delta$ T cells may be an absolute requirement for resistance to this infection.¹⁸ In addition to their role in resistance to infection, it has also been proposed that $\gamma\delta$ T cells may function as regulatory cells to limit immunopathology.¹⁶

In the present study we have examined the $\gamma\delta$ T-cell response to Yersinia enterocolitica. Encounter with this bacterium usually results in uncomplicated gastroenteritis. However, in some individuals reactive arthritis (ReA) can ensue (see 19 for review). ReA is an oligoarthritis which develops soon after infection of a mucosal site with a facultative or obligate intracellular pathogen such as Yersinia, Salmonella or Chlamydia. It is widely believed that an inappropriate or defective cellular immune response to the triggering bacteria may be responsible for this immunopathology. As ReA is associated with human leucocyte antigen (HLA)-B27, much interest has been focused on the CD8⁺ T cells. However, to date, HLA-B27-restricted CD8⁺ T-cell clones with

specificity for the inducing bacterium have only been isolated by one laboratory.²⁰ Most reports of cell-mediated immunity in patients have assayed CD4⁺ T cells and some bacterial antigens recognized by synovial CD4⁺ T-cell clones have been identified.²¹⁻²⁵ It would therefore appear that CD8⁺ T cells may not contribute solely to the immunopathology of ReA. The results of the present study suggest that, in addition to CD4⁺ and CD8⁺ T-cell activities, T cells expressing the $\gamma\delta$ TCR should not be ignored when considering ReA disease mechanisms. We report that a profound expansion of $\gamma \delta T$ cells is seen after culture of peripheral blood mononuclear cells (PBMC) from normal individuals or synovial fluid mononuclear cells (SFMC) from ReA patients with either Yersiniainfected and fixed autologous B lymphoblastoid cell line (BLCL) or with fixed bacteria. In contrast, heat-inactivated Yersinia did not support the proliferation of $\gamma\delta$ T cells. One Yersinia-reactive T-cell line from an arthritic individual was subjected to further testing and found to kill both infected autologous and allogeneic cell lines. Moreover, $\gamma \delta$ T-cell clones isolated from this line lysed infected targets in an MHCindependent fashion.

MATERIALS AND METHODS

Preparation of mononuclear cells from peripheral blood and synovial fluid

Synovial fluid was collected from three patients with a history of *Yersinia enterocolitica*-associated reactive arthritis. Peripheral blood was also donated by three normal individuals. The synovial fluids were treated with 10 U/ml hyaluronidase (Sigma Chemical Co. Ltd, Poole, UK) for 30-60 min at 37° . SFMC and PBMC were isolated on a Ficoll (Pharmacia, Milton Keynes, UK) gradient. The cells were either used fresh or cryopreserved until required.

Cell lines

Most of the B lymphoblastoid lines (BLCL) employed were generated from PBMC. The PBMC were incubated with supernatant from the Epstein-Barr virus (EBV)-producing cell line B95.8 at 37° for at least 1 hr. The cells were washed and cultured at $1-2 \times 10^6$ /ml in RPMI (Life Technologies Ltd, Renfrew, UK) supplemented with 5% heat inactivated fetal calf serum (HI FCS; Life Technologies Ltd), 10 mM HEPES (Sigma), 2 mM glutamine (Sigma), 100 μ g/ml streptomycin sulphate BP (Evans Medical Ltd, Greenford, UK) and 100 U/ml benzylpenicillin sodium BP (Glaxo, Greenford, UK) in the presence of 1 μ g/ml cyclosporin A. The cells were fed at weekly intervals with fresh medium supplemented with cyclosporin A was discontinued.

RML, HHKB, and PITOUT were given by S. Marsh (ICRF, London, UK). Daudi, C1R and K562 were provided by A. Rickinson (Department of Cancer Studies, University of Birmingham, UK).

Yersinia enterocolitica

Yersinia enterocolitica: O9 (Y9) was kindly donated by the Department of Infection, University of Birmingham, UK. The bacteria were grown overnight in 2YT broth in a shaking incubator at 37° . This late time of culture and temperature were chosen to generate stationary phase bacteria similar to those which may be encountered *in vivo*.

Infection and fixation of stimulatory and target cells

Cell suspensions were washed at least twice in antibiotic-free **RPMI.** An aliquot ($\approx 100 \ \mu l/1 \times 10^6$ cells) of the broth culture of Y9 was added to the cell suspension. The cells were then pelleted and incubated for 1 hr at 37°; this time being determined as sufficient to give infection without leading to premature lysis of cells. The infected cells were then washed three times in RPMI supplemented as described before except that 5% heat-inactivated male AB sera (West Midlands Blood Transfusion Service, Birmingham, UK) was used instead of FCS and 1 mm sodium pyruvate (Sigma), 1 × modified Eagle's medium (MEM) non-essential amino acid solution (Sigma), and 100 μ g/ml gentamycin (Sigma) were also added. The cells were used at this stage for cytotoxicity assays. For stimulation of SFMC or PBMC, the infected BLCL was fixed with 0.25% formaldehyde (Sigma) for 10 min, washed twice in 1.5 M glycine (Sigma), incubated for 30 min in RPMI supplemented as described, washed again and re-incubated prior to a last wash. The number of these cells used in stimulation and cloning assays was based on those cells which retained the ability to exclude trypan blue. Control uninfected BLCL were prepared in the same manner. For some experiments intact Yersinia enterocolitica were fixed as described above and added directly to cultures.

In vitro stimulation assays

The medium employed for all assays was RPMI supplemented as described in the previous section except that gentamycin was not routinely present. SFMC or PBMC were cultured in 2 ml volumes in 24-well plates at $1-2 \times 10^6$ /ml in the presence or absence of $3\cdot 3-6\cdot 7 \times 10^4$ /ml fixed infected or non-infected autologous BLCL or with fixed *Yersinia enterocolitica*.

T-cell cloning

Cells were cloned after 3–7 days of stimulation at limiting dilution in Terasaki plates (ICN Biomedicals Ltd, Thame, UK). The cells were cloned in the presence of irradiated PBMC obtained from three allogeneic donors (8×10^3 /well), infected autologous BLCL (1–5/well), 1 µg/ml phytohaemag-glutinin (PHA: Murex Diagnostics, Dartford, UK), 100 U/ml recombinant interleukin-2 (IL-2: Eurocetus UK Ltd, Harefield, UK) and 10% supernatant from concanavalin A stimulated PBMC. The isolated clones were maintained with 20–50 U/ml rIL-2 and restimulated, as required, with irradiated allogeneic PBMC from three donors and 1 µg/ml PHA.

Analysis of TCR gene usage

TCR γ and δ chain sequences were determined for a number of the T-cell clones using a procedure similar to that previously described for TCR α and β chain analysis.²⁶ Briefly, cDNA was prepared from RNA extracted from the T-cell clones, modified with a poly diguanosine triphosphate (dGTP) tail and subjected to anchor polymerase chain reaction (PCR). PCR for both γ and δ TCR used the anchor primer (5'CTATCTAGAGAGAGAGCTCGCGGCCGC₁₃3') and either the 3' C δ primer (5'CGCAGATCTCAGACAAGCGACAT TTG3') or the 3' C γ primer (5'CGCAGATCTGACAAAGG TATGTTCCAGCCTTCTGG3'). The amplification was performed in a Hybaid omnigene thermocycler (Hybaid Ltd. Teddington, UK) using 32 cycles of 93° for 45 s, 57° for 30 s and 72° for 1 min followed by extension at 72° for 4 min. The PCR products were digested with the restriction enzymes NotI and Bg/II and cloned into a M13mp18 cloning vector (kindly supplied by P. Moss and J. Bell, Oxford, UK). DNA sequencing was by the dideoxy chain termination method using a Sequenase version 2.0 reagent kit (Amersham International plc, Little Chalfont, UK).

Flow cytometry

Cells to be stained were washed and resuspended in phosphatebuffered saline (PBS) supplemented with 1% bovine serum albumin (BSA). The cells were stained with fluoroscein isothiocyanate (FITC)-conjugated monoclonal antibodies or control FITC-immunoglobulin G1 (IgG1) (Dako A/S, Glostrup, Denmark) in 96U-well floppy plates (Falcon Cell Culture Products, Cowley, UK) in 100 μ l volumes. The monoclonal antibodies employed were against CD3 (Dako), CD4 (Dako), CD8 (Dako), CD16 (Dako), TCR $\alpha\beta$ (Becton-Dickinson UK Ltd, Cowley, UK) and TCR $\gamma\delta$ (Becton-Dickinson). The cells were analysed with a FACSort® flow cytometer (Becton-Dickinson) and the data analysed using WinMDI version 1.3.1 (contact trotter@flosun.salk.edu).

Cytotoxicity assay

For most experiments the non-radioactive cytotoxicity kit, CytoTox-96TM, was used according to the manufacturer's directions (Promega Corporation, Southampton, UK). This is a colorimetric assay which measures the lactate dehydrogenase released from lysed cells. Target cells were added at 5000/well and effector cells were added at the indicated E:T ratio in a total of 100 μ l. The cells were incubated at 37° for 4 hr prior to harvesting 50 μ l supernatant. We found that the non-radioactive cytotoxicity kit gave comparable results to the traditional ⁵¹chromium release assay which was employed for

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the experiment shown in Fig. 6. In this case, aliquots of 1×10^6 target cells were labelled with 0·1 mCi/ml of ⁵¹Cr (Amersham International) in 100 μ l for 45 min, washed and then half the cells were infected with *Yersinia*, as described above. The assay was set up in triplicate in 200 μ l volumes with 5000 target cells and 50 000 effector cells. Maximum release was determined by adding Triton-X (Sigma) to cultures of target cells and minimum release from cultures of target cells in the absence of effector cells. After 4 hr, 50 μ l of supernatant was spotted onto a filter mat (Pharmacia) and radioactivity present assayed using a 1205 β -plate counter (LKB-Wallac, Turku, Finland). Percentage specific cytotoxicity was calculated in the standard manner.

RESULTS

A substantial proliferative reaction was noted when autologous BLCL infected with Yersinia were cultured with SFMC or PBMC. Flow cytometry analysis after 7-10 days of stimulation, revealed an expansion of TCR $\gamma\delta$ -bearing cells in the cultures (Fig. 1). Mononuclear cells from each of the individuals tested displayed less than 10% TCR $\gamma\delta^+$ cells prior to culture or after culture in the absence of infected BLCL (results not shown). The nature of the stimulus required for proliferation was further studied and the results from culture of SFMC from one individual, CM, are shown in Fig. 2. It is apparent that incubation of BLCL with Yersinia was required to promote expansion of $\gamma\delta$ T cells. Unfixed infected BLCL cultured with SFMC in the presence of gentamycin also stimulated $\gamma\delta$ T cell proliferation (results not shown). Interestingly, $\gamma\delta$ T-cell proliferation was also provoked by the addition of fixed intact Yersinia to SFMC. This stimulatory



Figure 1. Analysis of $\gamma\delta$ TCR expression after stimulation with fixed infected autologous BLCL. PBMC from normal individuals (top row) or SFMC from ReA patients (bottom row) were cultured for 7–10 days with fixed autologous BLCL infected with *Yersinia enterocolitica*. Expression of the $\gamma\delta$ TCR analysed by flow cytometry is shown. Each panel represents an independent experiment. In each case, TCR $\gamma\delta$ expression in unstimulated cultures was less than 6% and is not shown. No staining was apparent with an IgG1–FITC isotype control antibody.



Figure 2. TCR $\gamma\delta$ expression after culture of SFMC with various stimuli. TCR $\gamma\delta$ expression was determined by flow cytometry following culture of SFMC without added stimulators (a) or with fixed autologous BLCL (b), fixed infected BLCL (c), fixed *Yersinia* (d) or heat-inactivated *Yersinia* (e). No staining was detected with an isotype matched IgG1-FITC control (not shown).

capacity was almost completely ablated by heat inactivation of the Yersinia.

Cell lines generated from stimulation of CM SFMC were tested for their cytotoxic capacity. As shown in Table 1, when the line was tested against a panel of BLCL, although uninfected cell lines were killed to various degrees, enhanced killing was exhibited against the infected target cells irrespective of MHC allele expression. Cell lines derived from the PBMC of the three normal donors also killed infected target cells in an MHC-unrestricted fashion (results not shown).

T-cell clones were generated to study further the apparent MHC-unrestricted killing. Thirty-six of the 67 T-cell clones obtained were found to be TCR $\gamma\delta^+$. The remaining 31 clones

 Table 1. MHC-unrestricted killing of target cells by a Yersiniastimulated cell line

		Specific cytotoxicity		
Target cell	MHC matches	Uninfected	Infected	
СМ	Autologous	19.2	45.3	
GS	B27	11.8	47·2	
MR	B4 0	6.2	73.2	
ннкв	A3	26.6	60.5	
RML	Allogeneic	15.1	58.5	
PITOUT	Allogeneic	30.8	65.2	

A cell line generated by culturing SFMC from a patient, CM, with *Yersinia*-infected autologous BLCL was tested for its ability to kill various infected and uninfected target cell lines at E:T ratio of 10:1. The percentage specific cytotoxicity noted after 4 hr and MHC identities between autologous and other target cells are recorded.

expressed the $\alpha\beta$ TCR and CD4; no CD8⁺ T-cell clones were isolated. These CD4⁺ T cell clones were not analysed further. Twelve of the 36 TCR $\gamma \delta^+$ clones killed infected autologous targets to various degrees. Figure 3 shows cytotoxicity of eight representative clones. Flow cytometry was used to determine the expression of various surface markers and the results from a representative clone are depicted in Fig. 4. All of the clones had the T-cell markers CD2⁺ and CD3⁺ in conjunction with the $\gamma\delta$ TCR. The clones failed to express either of the co-stimulatory molecules CD4 or CD8. Staining for the natural killer (NK) marker, CD16 and the $\alpha\beta$ TCR was also negative. The $\gamma\delta$ TCR expressed by eight of the 12 Yersinia-reactive T-cell clones was examined by sequencing. As shown in Table 2, all of the clones were found to express $V\gamma 9$ in conjunction with V $\delta 2$. Moreover, with the exception of clone 5.32, all clones used the same J genes; JD1 and JGP. Two pairs of identical sequences (both DNA and deduced protein sequences) were found: 5.55 and 0.5.8, 5.23 and 5.4. The δ TCR CDR3 all exhibited a small hydrophobic amino acid (L, I or V) at position 97 and were of similar sequence length, being larger than those used for the γ chain. Analysis of 5.27 revelaed two in-frame sequences for δ gene usage, V δ 2 and V δ 1. However, only one in-frame sequence for γ gene products was found.

The T-cell clones were further tested for their ability to kill allogeneic infected BLCL. As shown in Fig. 5, BLCL were susceptible targets following their infection with *Yersinia*, irrespective of their MHC expression. Indeed, the allogeneic infected targets were killed more efficiently than the autologous infected cell line. An examination of the kinetics of the cytotoxic response revealed that killing could be detected as



Figure 4. Phenoype of a *Yersinia*-reactive T cell clone. Flow cytometry analysis of a T-cell clone stained with a control IgG1-FITC (a) or FITC-conjugated antibodies specific for CD3 (b), CD4 (c), CD8 (d), CD16 (e), TCR $\alpha\beta$ (f), or TCR $\gamma\delta$ (g) is shown.



Figure 3. TCR $\gamma\delta^+$ T cell clones kill *Yersinia*-infected but not uninfected autologous BLCL. Eight of the $\gamma\delta^+$ T-cell clones were tested for their ability to lyse uninfected and *Yersinia*-infected autologous BLCL at E:T=5:1. The percentage specific cytotoxic responses measured after 4 hr of incubation are shown.

early as 1 hr and was already maximal by 2.5 hr (results not shown). Cell lines deficient in MHC expression were also tested for their sensitivity to cytolysis by the T-cell clones (Fig. 6). Infection of Daudi cells (class I⁻), C1R cells (class I^{lo}) and K562 cells (class I and II^{lo}) enhanced their susceptibility to lysis by the T-cell clone. As for the allogeneic targets, cytolysis of these infected MHC-deficient cell lines was consistently greater than that observed for the autologous infected BLCL. Five of the $12 \gamma \delta$ T-cell clones were tested against MHC-deficient cell lines and all exhibited this MHC-independent killing of infected targets.

DISCUSSION

Our results indicate that cytotoxic T lymphocytes (CTL) with reactivity for Yersinia-infected targets can be readily generated from both PBMC of normal individuals as well as from SFMC of ReA patients. The CTL lines obtained were found to express predominantly the $\gamma\delta$ TCR. This profound expansion of TCR $\gamma\delta$ T cells in response to Yersinia is reminiscent of the reported stimulation of $V\gamma 9V\delta 2$ T cells in response to other bacteria such as mycobacteria, listeria and salmonella.²⁷⁻²⁹ Indeed, an analysis of the TCR $\gamma\delta$ gene usage from clones isolated from the Yersinia-stimulated T-cell lines revealed $V\gamma 9V\delta 2$ TCR products. Thus, Yersinia enterocolitica also has the potential to stimulate this T-cell subset. The extreme nature of the in vitro expansion (5-15-fold increase in percentage expressing TCR $\gamma\delta$) of the $\gamma\delta$ T cells was further reflected in the isolation of two sets of clones bearing identical TCR sequence; as the DNA sequences were invariant, these clones probably represent in vitro-derived sister clones. Although two δ gene products were found when 5.27 was analysed, only one in-frame γ gene product was found, despite analysis of multiple cloned PCR products. This suggests that either 5.27 is not clonal or allelic exclusion had failed, resulting in the generation of two δ gene products; such $\gamma\delta$ T-cell clones have been described.30

The Yersinia-reactive T-cell lines were found to kill infected target cells in an MHC-unrestricted fashion. Curiously, the T-cell line from the ReA patient, CM, exhibited poorest cytolysis of autologous or HLA-B27 matched infected BLCL. Results of recent experiments suggest that human $\gamma\delta$ T-cell clones, like NK cells, express receptors for self-MHC molecules which down-regulate killing of target cells expressing the

Table 2. T-cell receptor sequences of Yersinia-reactive T-cell clones

Clone	Vδ	n-D-n	Jδ		Vγ	n	Jγ	
5.55	CACD	ALGLNGOE	KLIFGKG	VD2/JD1	CALWEV		OELGKKIKVFGPG	VG9/JGP
0.5.8	CACD	ALGLNGOE	KLIFGKG	VD2/JD1	CALWEV		OELGKKIKVFGPG	VG9/JGP
5.23	CACD	TLLGDTY	KLIFGKGTRVT	VD2/JD1	CALW	DQK	ELGKKIKVFGPG	VG9/JGP
5.4	CACD	TLLGDTY	KLIFGKG	VD2/JD1	CALW	DÔK	ELGKKIKVFGPG	VG9/JGP
5.30	CAC	ESLGDTKS	TDKLIFGKG	VD2/JD1	CALWEV	QGGS	GKKIKVFGPG	VG9/JGP
0.5.6	CACD	PLGSGD	TDKLIFGKG	VD2/JD1	CALWE	EHR	ELGKKIKVFGPG	VG9/JGP
5.27	CACD	SILA	TDKLIFGKG	VD2/JD1	CALWEV	LG	LGKKIKVFGPG	VG9/JGP
	YFC	GSVVVGVLG	DKL	VD1/JD1				
5.32	CACD	PVEKRDTRQ	LIFGKGTRVT	VD2/JD1	CALWEV	QLARW	KLFGSGTTLVVT	VG9/JG1/2

The in-frame deduced amino acid sequences of TCR δ and γ chains are shown.



Figure 5. MHC-unrestricted cytotoxicity of infected target cells. Uninfected and infected autologous or allogeneic BLCL were incubated in a ⁵²Cr-release cytotoxicity assay with one of the $\gamma\delta$ T-cell clones (0.5.1) at an E:T ratio of 10:1 for 4 hr. The percentage specific cytotoxicity is shown.

relevant MHC molecules.³¹ Thus, the poor killing of HLA-B27 target cells may be due to a negative signal mediated via binding to a B27-reactive receptor on the TCR $\gamma\delta$ CTL. A further explanation for the poor killing of B27 expressing targets is suggested by the work of Kapasi and Inman^{32,33} who reported that cell lines expressing this MHC allele exhibited reduced rates of infectivity with gram-negative bacteria associated with reactive arthritis. Thus, infected cells expressing HLA-B27 may display a reduced number of T-cell ligands compared to infected target cells lacking this MHC molecule.



Figure 6. MHC-independent cytotoxicity of infected target cells. The $\gamma\delta$ T-cell clone was incubated at an E:T ratio of 10:1 with the indicated infected or uninfected target cells for 4 hr. The percentage specific cytotoxic response for each target is shown. (a) and (b) represent independent experiments.

To confirm that the MHC-unrestricted killing exhibited by the T-cell lines was not due to mixed populations we generated T-cell clones. The $\gamma\delta$ T-cell clones, like the cell lines, exhibited MHC-unrestricted killing of infected cell lines. Moreover, from the results with MHC-deficient cell lines, it is also apparent that the $v\delta$ T cells recognized infected target cells in an MHCindependent manner. As we did not analyse the cytotoxic capabilities of the CD4⁺ TCR $\alpha\beta$ T-cell clones, we cannot affirm that the pattern of killing documented is unique to the $\gamma\delta$ T-cell clones. Nevertheless, TCR $\gamma\delta^+$ T cells exhibiting such MHC-independent activities have been described.³⁴ It has been suggested that some $\gamma \delta$ T cells may, as has been demonstrated for a CD4⁻ CD8⁻ $\alpha\beta$ T cell,^{12,13} rely on antigen presented by CD1, a family of non-polymorphic β^2 microglobulinassociated glycoproteins. However, as the $\gamma\delta$ T cells reported here recognize infected Daudi cells (a cell line lacking β 2-microglobulin), they cannot require the expression of CD1. Non-peptidic, phosphorylated ligands for TCR $\gamma\delta$ T cells, particularly those expressing a $V\gamma 9V\delta 2$ TCR, have been identified from mycobacteria.⁴⁻⁶ Such compounds can stimulate the proliferation of $V\gamma 9V\delta 2$ T cells in the absence of classic MHC molecules or classic MHC-presentation pathways.⁷ Given the preponderance of $V\gamma 9V\delta 2$ TCR gene usage in the isolated T-cell clones, it is possible that Yersinia contain such nonpeptidic ligands or give rise to the expression of equivalent cellular derived compounds upon infection of cells. Precisely how the phosphorylated ligands are recognized by the $\gamma\delta$ T cells is, as yet, unknown. However, evidence is accumulating suggesting that at least some $\gamma\delta$ T cells may be stimulated by native unprocessed antigens, in an analogous fashion to the engagement of the immunoglobulin receptor on B cells.³⁴ The CDR3 regions of immunoglobulin light and heavy chains and TCR γ and δ chains are more similar in length distribution than the CDR3 regions of the α and β TCR molecules.³⁵ Thus the $\gamma\delta$ complex may be able to fold in an immunoglobulinlike manner to permit the recognition of antigenic shapes outside the constraints of the MHC-peptide complex used by the $\alpha\beta$ TCR. In agreement with this study, the CDR3 loop lengths reported here were found to be larger for the δ chains compared to the γ chains. However, the δ CDR3 regions showed remarkable consistency in length and were shorter than most of the sequences reported by Rock et al.³⁴ Moreover, our results indicate the occurrence of a small hydrophobic amino acid (L, I or V) at position 97 in the δ sequences. This sequence is poorly represented in fetal $\gamma\delta$ T cells, and it has been suggested that peripheral antigenic stimulation leads to the expansion of $\gamma\delta$ T cells with such a motif.^{36,37} We have not determined the nature of the stimulatory ligand for the Yersinia-reactive $\gamma\delta$ T cells reported here. However, formaldehyde-fixed, but not heat-inactivated, bacteria were able to stimulate the proliferation of $\gamma\delta$ T cells when added to mononuclear cells. Thus, the stimulation of $\gamma\delta$ T cells was not dependent upon the addition of B lymphoblastoid cell lines to the synovial fluid mononuclear cell cultures. Our results are compatible with a native, heat-labile ligand for $\gamma\delta$ T-cell stimulation. Moreover, since active infection was not required to provoke the expansion of the $\gamma\delta$ T cells, the stimulatory ligand is likely to be a Yersinia-derived antigen rather than a self antigen provoked by infection (e.g. heat-shock protein).

Regardless of the nature of the stimulatory ligand for the Yersinia-reactive $\gamma\delta$ T cells, the question arises as to the involvement of this T-cell subset in ReA. Previous studies of SFMC from ReA patients have also described a proliferation of $\gamma \delta$ T cells in response to live but not dead Yersinia enterocolitica.³⁸ Moreover, cytotoxicity of bacteria-infected target cells was shown for five of eight $V\gamma 9V\delta 2$ T-cell clones isolated by random cloning from SFMC, with four of those clones thought to recognize heat shock protein 65.39 These clones, like the ones described here, exhibited MHC-unrestricted cytotoxicity. From the present results, it is clear that $\gamma \delta$ T-cell proliferation can be induced by Yersinia not only from SFMC but also from PBMC donated by normal individuals. Whilst it is generally believed that $\gamma\delta$ T cells are important in immune responses to infectious agents, studies have also implicated a pathological role for this subset of T cells.¹⁴ It is clear that the $\gamma\delta$ T-cell response to *Yersinia* is not absent in ReA patients. However, we did not determine whether there was a quantitative or qualitative difference in the $\gamma\delta$ T-cell responses between normal and arthritic individuals. Because of the association of ReA with HLA-B27, one would not predict a pathological role for a T-cell population which displays MHC-independent cytolysis of infected target cells. However, HLA-B27, while not directly involved in $\gamma\delta$ T cell recognition of infected targets

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may, as discussed earlier,³¹⁻³³ indirectly modulate the activities of the $\gamma\delta$ T cells. Such interference with $\gamma\delta$ T cells may have consequences for the $\alpha\beta$ T-cell subset. For example, murine studies have demonstrated that, from the earliest stages of infection, $\gamma\delta$ T cells can influence the cytokine milieu determining whether a T helper 1 (Th1) or Th2 response predominates.^{40,41} It would be of interest to determine whether the *Yersinia*-responsive $\gamma\delta$ T cells from ReA individuals display a different cytokine response than those in healthy individuals.

In conclusion, the results of the present study indicate that both normal PBMC and SFMC can mount a potent *in vitro* response to *Yersinia enterocolitica*. The responsive cells express predominantly the $\gamma\delta$ TCR ($V\gamma9V\delta2$) and such cells can kill infected targets in an MHC-independent manner. We have yet to determine whether such $\gamma\delta$ T cells can play a pathological or beneficial role in ReA.

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