

Elevated Numbers of Gamma-Delta ($\gamma\delta$ +) T Lymphocytes in Children with Immune Thrombocytopenic Purpura

RUSSELL E. WARE^{1,2} and THAD A. HOWARD¹

Accepted: March 14, 1994

Immune thrombocytopenic purpura (ITP) in childhood is a heterogeneous clinical disorder characterized by immune-mediated platelet destruction. Although generally considered to involve autoreactive B lymphocytes which produce antiplatelet antibodies, there is increasing evidence that T lymphocytes also play an important role in this autoimmune process. We studied 11 children with acute ITP and 19 children with chronic ITP and observed elevated numbers of TCR $\gamma\delta$ + T lymphocytes in several patients. In the three children with the highest elevations (TCR $\gamma\delta$ + / CD3+ percentage ranging from 37.8 to 48.1% at initial evaluation), the expanded cell population exclusively expressed the surface V δ 2/V γ 9 heterodimer and had enhanced *in vitro* proliferation to mycobacterial extracts and IL-2. Analysis of the nucleotide sequences used by these TCR $\gamma\delta$ + cells demonstrated a diverse set of VDDJC gene rearrangements, indicating polyclonal expansion of cells reminiscent of a superantigen response. There was a close correlation between the number of TCR $\gamma\delta$ + T lymphocytes and the degree of thrombocytopenia in each patient. TCR $\gamma\delta$ + T lymphocytes may be important in the pathogenesis of immune-mediated platelet destruction in some children with ITP.

KEY WORDS: Immune thrombocytopenic purpura; T lymphocytes; autoimmunity; gamma-delta lymphocytes; heat shock protein.

INTRODUCTION

Immune thrombocytopenic purpura (ITP) is a hematologic disorder characterized by immune-mediated destruction of platelets (1). In over 90% of cases, there is measurable autoantibody on the

surface of circulating platelets (2, 3), which leads to clearance by the reticuloendothelial system and consequent thrombocytopenia (1, 4). In adults, ITP is typically characterized by persistent thrombocytopenia. ITP in childhood is a more heterogeneous disorder; most children have a self-limited form known as acute ITP, in which the thrombocytopenia resolves within 6 to 12 months. In contrast, 10–20% of children will have a prolonged clinical course termed chronic ITP, in which the thrombocytopenia persists for over 12 months (1, 5).

The pathogenesis of immune dysregulation which leads to antiplatelet autoantibody production in ITP is not well understood. Platelet autoantibodies typically are IgG (2, 6), although IgM and IgA antibodies can be detected on the platelets of some patients. The antibodies are usually polyclonal but may be oligoclonal (7) and have reactivity against a variety of platelet surface glycoproteins (8, 9). We and others have demonstrated that peripheral T lymphocytes from some patients with ITP have *in vitro* reactivity against platelets, which is measurable at both the polyclonal (10, 11) and the clonal level (11). It is therefore likely, at least in a subset of patients with ITP, that peripheral blood T lymphocytes play a direct role in the B lymphocyte-mediated autoimmune response (5, 12). However, the T lymphocyte subsets which are important in the pathophysiology of ITP and the mechanisms by which they contribute to immune platelet destruction have not been elucidated.

Recently, T lymphocytes which express the gamma/delta ($\gamma\delta$) T cell receptor (TCR $\gamma\delta$ + lymphocytes) have been implicated in a variety of autoimmune human diseases. Increased numbers of TCR $\gamma\delta$ + T lymphocytes have been reported in rheumatoid arthritis (13, 14), polymyositis (15), Sjogren's syndrome, and systemic lupus erythema-

¹Department of Pediatrics, Division of Hematology/Oncology, Duke University Medical Center, Durham, North Carolina 27710.

²To whom correspondence should be addressed at P.O. Box 2916, Duke University Medical Center, Durham, North Carolina 27710.

tosus (16). An adult patient with chronic neutropenia and an elevated number of T lymphocytes with a hybrid C α V δ 1 surface receptor has been reported (17). To date, however, there have been no descriptions of TCR $\gamma\delta$ + T lymphocytes in children or adults with ITP.

In this study, we report that increased numbers of peripheral blood TCR $\gamma\delta$ + T lymphocytes are present in some children with ITP. Three children with the highest elevations (37.8, 40.2, and 48.1% of CD3+ T lymphocytes, respectively) had additional investigation with the following observations: (i) predominant expression of the V δ 2/V γ 9 heterodimer, indicating a restricted TCR variable gene expression in the expanded cell population; (ii) enhanced *in vitro* proliferative responses to mycobacterial extracts and IL-2, but not against human heat shock protein or allogeneic platelets; (iii) polyclonal expansion of TCR $\gamma\delta$ + cells, with polymerase chain reaction (PCR) analysis demonstrating both V δ 2-J δ 1 and V δ 2-J δ 3 gene rearrangements and nucleotide analysis demonstrating unique V-D-D-J sequences; and (iv) a close correlation between the number of circulating TCR $\gamma\delta$ + T lymphocytes and the degree of thrombocytopenia. We hypothesize that TCR $\gamma\delta$ + T lymphocytes may be important in the pathogenesis of immune-mediated platelet destruction in some children with ITP.

METHODS

Patients. Patients were grouped into three categories: children with acute ITP, children with chronic ITP, and young adult normal volunteers. A total of 11 children with acute ITP (4 males and 7 females) were studied, with a mean age of 5.5 years, mean platelet count of $19 \times 10^9/L$, and median platelet count of $8 \times 10^9/L$. A total of 19 children with chronic ITP (11 males and 8 females) was studied, with a mean age of 10.7 years, mean platelet count of $49 \times 10^9/L$, and median platelet count of $40 \times 10^9/L$ excluding 2 patients with postsplenectomy normalization. Ten young adult volunteers (six males and four females) were also studied. Hematologic characteristics of some of these children have been previously described (11). All blood was obtained in accordance with a protocol approved by the Duke University Institutional Review Board.

Antibodies. Directly labeled monoclonal antibodies (mAbs) conjugated to either fluorescein isothiocyanate (FITC) or phycoerythrin (PE) were used

for single- and double-color analysis of lymphocyte subpopulations. The mAbs included CD2-PE, CD3-FITC, CD3-PE, CD4-FITC, CD4-PE, CD8-FITC, CD8-PE, CD20-FITC, and CD45RO-PE (Dako Corporation, Carpinteria, CA). Additional mAbs CD7, CD45RA (2H4), CD29 (4B4), and CD56 (NKH1), all conjugated to phycoerythrin, were kind gifts from Meryl Forman (Coulter Corporation, Hialeah, FL). Analysis of T cell receptor (TCR) antigens included Pan-TCR $\alpha\beta$, TCR δ 1, δ V1, δ V2, and γ V2 (T cell Diagnostics, Cambridge, MA). Control antibodies included IgG-FITC and IgG-PE from Dako. All antibodies were used according to the manufacturers' recommendations.

Immunophenotyping. Peripheral blood mononuclear cells (PBMC) were obtained from heparinized venous blood using Ficoll-Hypaque density centrifugation (18). Approximately 0.5×10^6 PBMC were incubated with saturating amounts of one or two mAbs for 30 min at 4°C as described previously (19). After washing with phosphate-buffered saline (PBS) with 2% bovine serum albumin (Sigma, St. Louis, MO), cells were fixed in paraformaldehyde and stored at 4°C in the dark until analysis. Samples were analyzed on a Profile II flow cytometer (Coulter) using forward- and side-scatter lymphocyte gating and color compensation.

In Vitro Proliferation Assays. PBMC from patients and controls were resuspended in RPMI with 15% human A serum (HAS; Worldwide Biologicals, Cincinnati, OH) at $1.0 \times 10^6/ml$. Final concentrations of stimuli included 0.5–2.0 μ g/ml phytohemagglutinin (PHA; Burroughs Wellcome, Research Triangle Park, NC), 0.5–2.0 μ g/ml concanavalin A (Con A; Sigma), 0.5–2.0 μ l/ml pokeweed mitogen (PWM; GIBCO, Grand Island, NY), tetanus toxoid (Connaught Laboratory, Toronto Canada), 0.125–1.0 mg/ml *Candida albicans* (Greer Lab, Boone, NC), 100 U/ml recombinant interleukin-2 (rIL-2; gift from Chiron Corporation, Emeryville, CA), 10 μ g/ml mycobacterial extract (mTB; gift from Dr. Drew Pardoll), 10 μ g/ml recombinant *Mycobacterium bovis* BCG 65-kD protein (gift from the World Health Organization), 10 μ g/ml 65-kD recombinant human heat shock protein (hsp65; gift from Satish Jindal of Procept, Cambridge, MA), and $100 \times 10^6/ml$ allogeneic platelets which were purified as described (20).

For blastogenesis assays, 100 μ l (10^5) cells were placed in triplicate into 96-well U-bottom plates (Costar, Cambridge, MA) along with 100 μ l stimuli and incubated at 37°C (3–4 days for PHA, 4–5 days

for Con A, 5–6 days for PWM, 6 days for *Candida* and tetanus, and 7 days for all other stimuli). Cells were pulsed with 0.4 μCi ^3H -thymidine (New England Nuclear, Boston, MA), harvested on a cell harvester (Cambridge Technologies, Watertown, MA), and counted on a beta scintillation counter. For bulk culture assays, 750 μl of cells was placed into 24-well plates (Costar) along with 750 μl of stimuli. After 7–9 days, 100- μl aliquots of cells were removed for immunophenotype analysis.

Reverse Transcriptase Reaction. Total RNA was harvested from PBMC of patients and controls using a cesium centrifugation method as described previously (19), then used in a reverse transcriptase (RT) reaction to generate cDNA. Total RNA (2.0 μg) was added to a 20- μl mixture containing 30 U of AMV RT (Stratagene, La Jolla, CA), 20 U of RNase inhibitor (Promega, Madison, WI), 1 μg of oligo(dT) (Pharmacia, Piscataway, NJ), 20 nmol (1 mM final concentration) of dNTPs (Pharmacia), 2 μl of 10 \times buffer (500 mM Tris pH 8.3, 200 mM KCl, 100 mM MgCl_2 , and 50 mM dithiothreitol), and ddH₂O. The RT reaction was incubated at room temperature for 10 min, then 42°C for 60 min, followed by 70°C for 10 min and quenching on ice.

Polymerase Chain Reaction. The V δ 2 primer from 5' to 3' was GGGGTCGACCCTCAGGTGCTCCATGAA, which contains a 5' *Sal*I restriction endonuclease site to allow subcloning of the PCR fragment. The J δ 1 primer, GGGGAATTCCACAGTCACACGGGTTC, and the J δ 3 primer, GGGGAATTCCACGAAGAGTTTGATGCC, each contain a 5' *Eco*RI cutting site. A 50- μl PCR reaction contained 1.0 μl of the RT reaction as a source of template cDNA, 1 μg of each primer, 2.5 units of *Taq* polymerase (Stratagene), 10 nmol of dNTPs, 5 μl of 10 \times reaction buffer (Stratagene), and ddH₂O. The reaction mixture was heated to 94°C for 3 min for initial denaturation, followed by 35 cycles at 94°C for 1 min, 51°C for 2 min, and 72°C for 2 min. Negative controls included the following reactions: (i) RT, no primers added; (ii) no RT, primers added; and (iii) no RT, no primers added.

Analysis of V δ 2 Gene Rearrangements. The PCR product was purified using Prep-a-gene (Bio-Rad) prior to restriction endonuclease digestion with *Eco*RI and *Sal*I, then purified again prior to ligation. Bluescript vector (Stratagene) was also cut with both restriction endonucleases, and ligation was performed in a 10- μl reaction containing T4 ligase (Pharmacia), dATP (Pharmacia), and 10 \times ligase buffer. The ligated product was transformed into

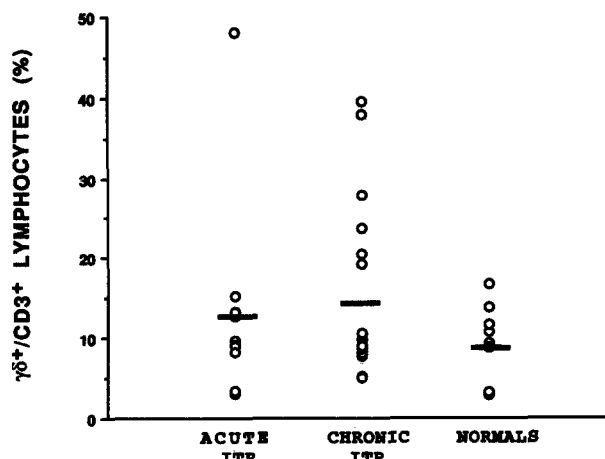


Fig. 1. Gamma-delta ($\text{TCR}\gamma\delta^+$) T lymphocytes in childhood ITP. Two-color analysis of PBMC from 11 children with acute ITP, 19 with chronic ITP, and 10 normal volunteers demonstrated elevated numbers of $\text{TCR}\gamma\delta^+$ T lymphocytes in several affected children. The horizontal bars represent the mean values, which were not significantly different among the three groups. The three children with the highest values (37.8, 40.2, and 48.1% of $\text{CD}3^+$ lymphocytes, respectively) underwent additional investigation.

TG-1 bacteria, and positive colonies were first identified using blue-white β -galactosidase selection and then confirmed by PCR using flanking T3 and T7 primers. Individual clones which contained inserts of the correct size were then grown overnight in a 10-ml culture, and the amplified plasmids were purified using Magicprep (Promega) and chemically denatured. Sequencing reactions were performed in both directions using the T3 and T7 primers and Sequenase (United States Biochemical, Cleveland, OH); sequencing gels were 6% acrylamide and run at 50°C.

Statistical Analysis. Statistical tests were performed using the Primer of Biostatistics (McGraw-Hill, New York) software package. The *t* test was used to compare values among the three groups.

RESULTS

Enumeration of $\text{TCR}\gamma\delta^+$ T Lymphocytes

The number of $\text{TCR}\gamma\delta^+$ lymphocytes as a percentage of peripheral blood $\text{CD}3^+$ T lymphocytes for each patient group is shown in Fig. 1. The 11 children with acute ITP had an average (mean \pm SE) of $13.1 \pm 3.7\%$ $\text{TCR}\gamma\delta^+$ lymphocytes, the 19 children with chronic ITP had $14.7 \pm 2.4\%$, and the normal volunteers had $8.3 \pm 1.6\%$ $\text{TCR}\gamma\delta^+$ lym-

Table I. Phenotypic and Functional Analysis of Peripheral Blood Lymphocytes at Initial Patient Evaluation^a

	Patient 1	Patient 2	Patient 3	Control
T lymphocyte enumeration (%)				
TCR $\alpha\beta$ /CD3	58.4	69.7	38.9	95.4 \pm 7.5
TCR $\gamma\delta$ /CD3	40.2	37.8	48.1	8.5 \pm 2.0
V δ 1/TCR $\gamma\delta$	7.7	4.7	9.8	36.7 \pm 10.0
V δ 2/TCR $\gamma\delta$	84.2	80.3	91.1	60.0 \pm 10.0
V γ 9/TCR $\gamma\delta$	94.7	92.0	93.3	66.7 \pm 10.0
Lymphocyte proliferation (cpm)				
Medium	2,292	1,713	2,445	3,116
PHA	214,368	169,828	225,100	207,383
Con A	261,533	211,120	218,812	218,275
PWM	183,273	168,933	115,315	125,294
Tetanus	7,351	50,126	157,785	120,067
Candida	2,471	89,275	81,737	41,774
mTB	139,244	10,618	51,547	56,252
BCG	12,769	4,038	151,941	9,748
rIL-2	286,744	151,941	364,519	216,753
rIL-2 + cyclosporin A	143,760	87,188	160,358	66,036
hsp65	2,494	2,640	1,887	3,360
Platelets	1,187	2,695	2,016	1,943

^aControl values (mean \pm SE) for lymphocyte enumeration are from nine normal volunteers, while control values for lymphocyte proliferation are the mean of three to five normal volunteers for each point.

phocytes. There were no statistically significant differences in the number of TCR $\gamma\delta$ + lymphocytes among the three groups, but several children with increased numbers of TCR $\gamma\delta$ + lymphocytes were identified (Fig. 1). Two children with chronic ITP and one with acute ITP had highly elevated numbers of TCR $\gamma\delta$ + lymphocytes on initial evaluation and were evaluated further.

Case Reports

Patient 1. A 23-month-old white male in good health developed a minor upper respiratory infection and, 2 weeks later, had bruising and petechiae; his peripheral blood counts were normal except for a platelet count of $3 \times 10^9/L$. He was diagnosed with ITP and treated with intravenous immunoglobulin (IVIG) with a good response, but required monthly treatments to maintain a platelet count above $30 \times 10^9/L$. Seven months later, the thrombocytopenia worsened and he became less responsive to IVIG therapy. Bone marrow aspirate showed slightly increased numbers of megakaryocytes and normal myeloerythroid maturation. Therapy with oral corticosteroids was effective, but thrombocytopenia recurred after steroid taper.

The patient was referred to Duke University Medical Center (DUMC) 12 months after diagnosis. Peripheral blood counts were normal except for a platelet count of $8 \times 10^9/L$. HIV ELISA was

negative, and bone marrow aspirate revealed a normal 46XY karyotype. Table I lists his initial PBMC immunophenotype, which included an abnormally high number of TCR $\gamma\delta$ + T lymphocytes (40.2% of CD3+ T lymphocytes; absolute count, 1266 cells/ μ l). Figure 2 depicts his platelet counts and TCR $\gamma\delta$ + lymphocyte percentages over time and demonstrates that the elevation was documented on seven occasions over the next 5 months. A trial of cyclosporin A (4–11 mg/kg/day orally divided b.i.d. to achieve a serum level of 200 μ g/ml) led to a slightly increased percentage of peripheral blood TCR $\gamma\delta$ + T lymphocytes. Following elective splenectomy, the platelet count and the number of TCR $\gamma\delta$ + T lymphocytes normalized.

Patient 2. A 10-year-old white male in good health presented with a 1- to 2-week history of epistaxis, bruising, petechiae, and oral mucosal bleeding. Peripheral blood count included a platelet count of $5 \times 10^9/L$, and he was referred to DUMC for further evaluation. Medical history and family history were negative for hematologic or autoimmune diseases, but the child had received a booster immunization of diphtheria/tetanus 12 days prior to presentation. Physical examination was normal except for bruising and petechiae, and peripheral blood counts were normal except for thrombocytopenia. Bone marrow aspirate was normal including 46XY karyotype, and HIV ELISA was negative. Table I shows the patient's initial PBMC

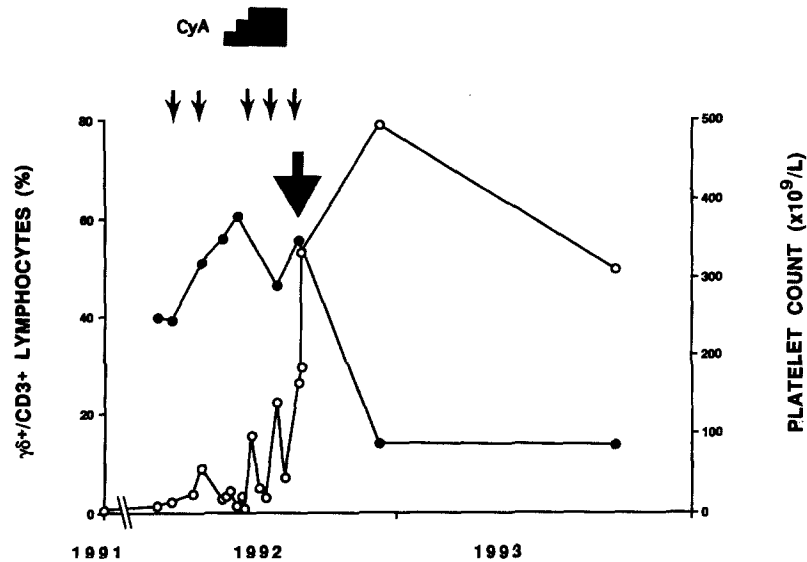


Fig. 2. Serial measurements of TCR $\gamma\delta$ + T lymphocytes and platelet counts in patient 1 with chronic ITP. The platelet counts (open circles) were below $100 \times 10^9/L$ from the time of diagnosis, despite therapy with IVIG (small vertical arrows) and cyclosporin A (CyA). Following elective splenectomy in late 1992 (large vertical arrow), the platelet count normalized. Measurement of TCR $\gamma\delta$ + T lymphocytes (filled circles) demonstrated 40–60% TCR $\gamma\delta$ + cells on multiple occasions despite therapy but normal values (<15%) following splenectomy.

immunophenotype, with an elevated number of TCR $\gamma\delta$ + T lymphocytes (37.8% of CD3+ T lymphocytes; absolute count, $539 \text{ cells}/\mu\text{l}$). He was initially treated with IVIG and has received periodic additional treatments for recurrent thrombocytopenia. He is currently 11 months from diagnosis (6 months since last IVIG), and his most recent platelet count was $97 \times 10^9/L$. Coincident with the amelioration of his thrombocytopenia, the number of circulating TCR $\gamma\delta$ + T lymphocytes diminished and, at last measurement, was 21.2% of CD3+ lymphocytes.

Patient 3. A healthy 13-year-old white male presented with a 10-day history of spontaneous bruises on his trunk and extremities. Peripheral blood counts included a platelet count of $45 \times 10^9/L$ but were otherwise normal. Medical history and family history were negative. At DUMC 3 days later, his platelet count had risen to $110 \times 10^9/L$ and his erythrocyte sedimentation rate was 10 mm/hr; fluorescent antinuclear antibody and HIV ELISA were negative. Bone marrow aspirate revealed increased numbers of megakaryocytes, normal myeloid maturation, and 46XY karyotype. Table I shows the PBMC immunophenotype at diagnosis and demonstrates an elevated number of TCR $\gamma\delta$ + T lymphocytes (48.1% of CD3+ T lymphocytes;

absolute count, $643 \text{ cells}/\mu\text{l}$). His platelet count rose spontaneously over the next month to $241 \times 10^9/L$ and, most recently (13 months from diagnosis), was $299 \times 10^9/L$. Figure 3 depicts the patient's platelet counts and percentage of TCR $\gamma\delta$ + T lymphocytes over time and shows that the number of TCR $\gamma\delta$ + T lymphocytes decreased toward the normal range (13–20%) coincident with resolution of his thrombocytopenia.

Phenotypic Subset Analysis of TCR $\gamma\delta$ + T Lymphocytes

Because TCR $\gamma\delta$ + T lymphocytes are heterogeneous and can be categorized on the basis of their variable gene usage, we performed additional immunophenotypic analysis. In each case, the TCR $\gamma\delta$ + T lymphocytes were almost exclusively of the V δ 2 subset, with few cells reacting with the V δ 1 mAb. In addition, the cells also were strongly positive with the V γ 9 mAb, demonstrating that the cells expressed the V δ 2/V γ 9 heterodimer (Table II).

Two-color analysis further demonstrated that the TCR $\gamma\delta$ + T lymphocytes of each patient were phenotypically similar to those of controls (Table II). Patient TCR $\gamma\delta$ + T lymphocytes expressed CD2, CD3, CD7, and CD29 antigens but were

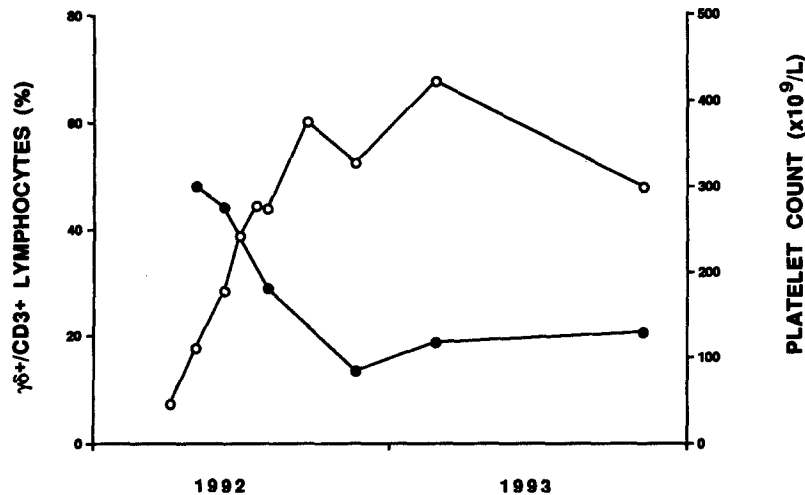


Fig. 3. Serial measurements of TCR- $\gamma\delta$ + T lymphocytes and platelet counts in patient 3 with acute ITP. The platelet counts (open circles) were initially low at the time of diagnosis, quickly exceeded $100 \times 10^9/L$, and have remained normal. Measurement of TCR- $\gamma\delta$ + T lymphocytes (filled circles) demonstrated >40% TCR- $\gamma\delta$ + cells on two occasions early in the disease but much lower values following the spontaneous resolution of the thrombocytopenia.

negative for CD4 and CD56 expression. CD8 expression was negative for patient 1 and low for patients 2 and 3, similar to CD8 expression in controls. Analysis of CD45 isoforms demonstrated that the majority of TCR- $\gamma\delta$ + cells was CD45RO+ in patients 1 and 2, while patient 3 (the child with acute ITP) had lower numbers of CD45RO+ TCR- $\gamma\delta$ + T lymphocytes at diagnosis, but predominantly CD45RO+ cells (82.6%), after resolution of thrombocytopenia.

Table II. Phenotypic Analysis of TCR- $\gamma\delta$ + T lymphocytes^a

mAb	Patient 1	Patient 2	Patient 3	Controls (mean \pm SE)
V δ 1	5.8	4.7	10.0	31.6 \pm 4.4
V δ 2	86.3	80.3	91.1	67.7 \pm 4.9
V γ 9	95.2	92.0	93.1	74.2 \pm 4.8
CD2	87.4	94.7	93.9	95.2 \pm 1.0
CD3	99.6	93.8	99.6	99.0 \pm 0.5
CD4	2.3	0.0	0.8	1.5 \pm 0.3
CD7	99.0	98.6	97.4	93.0 \pm 3.0
CD8	3.7	10.2	30.1	19.6 \pm 1.7
CD29	89.9	84.9	80.7	79.5 \pm 2.4
CD45RA	13.2	11.9	19.7	34.5 \pm 6.6
CD45RO	61.9	60.9	22.5	45.0 \pm 5.0
CD56	4.5	15.6	27.0	14.6 \pm 2.7

^aPBMC from each patient at initial examination were stained with TCR δ 1-FITC to identify all TCR- $\gamma\delta$ + T lymphocytes and with additional mAbs to analyze and calculate TCR- $\gamma\delta$ subsets. Results are the percentage of TCR- $\gamma\delta$ + T lymphocytes which were positive for each additional mAb. Control values derive from other children with ITP ($n = 16$) and normal volunteers ($n = 6$).

Lymphocyte Proliferative Responses

To determine the functional characteristics of the TCR- $\gamma\delta$ + T lymphocytes, PBMC from each patient were tested in blastogenesis assays for proliferation to various stimuli. Table I shows that at initial evaluation, each child demonstrated normal [³H]thymidine incorporation against mitogens including PHA, Con A, and pokeweed mitogen. Responses to recall antigens tetanus and candida were normal in patients 2 and 3 but low in patient 1, who was very young.

In vitro proliferative responses of patient PBMC to mycobacterial extract were positive as measured by [³H]thymidine incorporation, similar to values obtained with controls (Table I). However, the mTB stimulus induced an enhanced proliferation of patient TCR- $\gamma\delta$ + cells in bulk culture (Fig. 4); these results were much greater than those obtained from controls. Similarly, recombinant IL-2 led to strong [³H]thymidine incorporation in both patient and control PBMC but induced enhanced growth of patient TCR- $\gamma\delta$ + T lymphocytes compared with controls (Fig. 4).

In vitro blastogenesis using recombinant bovine BCG and human heat shock protein (hsp65) showed no proliferative responses (Table I) and no selective proliferation of TCR- $\gamma\delta$ + cells in bulk culture (data not shown). Stimulation with allogeneic platelets also induced no proliferation.

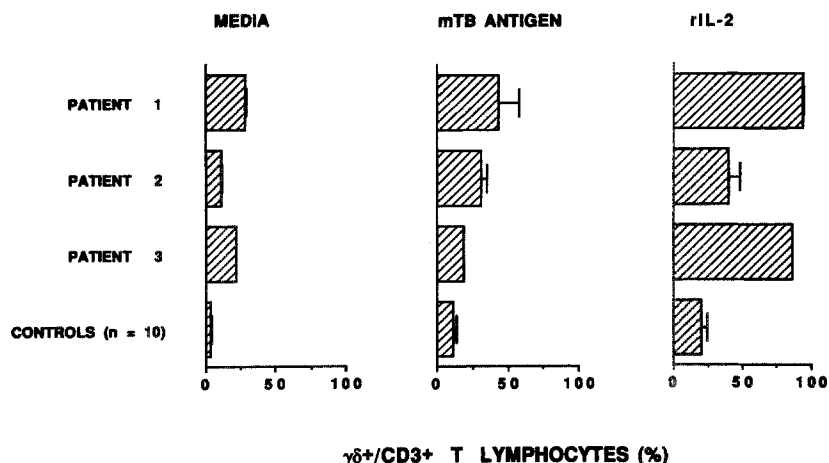


Fig. 4. Enhanced *in vitro* proliferation of TCR $\gamma\delta^+$ T lymphocytes in childhood ITP. PBMC from patients and controls were incubated in bulk culture with stimuli for 7–9 days, then immunophenotyped for the number of TCR $\gamma\delta^+$ cells as a percentage of CD3⁺ T lymphocytes. The three children with high values of TCR $\gamma\delta^+$ T lymphocytes demonstrated an enhanced TCR $\gamma\delta^+$ cell proliferation when stimulated with mTB antigen or rIL-2. The control values included both normal volunteers and other children with ITP. The data for patients 1 and 2 (with chronic ITP) were obtained from experiments on two different occasions, while data for patient 3 (with acute ITP) were obtained at the time of initial evaluation.

Analysis of V δ 2 Gene Rearrangements

To determine whether the elevated numbers of V δ 2⁺/V γ 9⁺ TCR $\gamma\delta^+$ T lymphocytes in our three patients represented a monoclonal, oligoclonal, or polyclonal expansion, V δ 2–J δ gene rearrangements were directly analyzed. RT-PCR using the V δ 2 primer with either the J δ 1 or the J δ 3 primer demonstrated that the patients utilized both J δ 1 and J δ 3 gene segments in combination with the V δ 2 gene (Fig. 5). These results showed that the TCR $\gamma\delta^+$ T lymphocytes did not represent a monoclonal population of cells. Next RT-PCR was performed using V δ 2 and C δ primers, followed by cloning of individual PCR products and sequencing of the VDDJ gene rearrangements. Figure 6 shows the nucleotide sequence of multiple clones from each patient and demonstrates that each V δ 2–D–D–J δ 1 nucleotide sequence was unique. Similar results were obtained for the V δ 2–D–D–J δ 3 nucleotide sequences (data not shown). These results demonstrate unequivocally that the elevated numbers of TCR $\gamma\delta^+$ T lymphocytes in the three children represented, in each case, a polyclonal expansion of cells.

DISCUSSION

Cells which express the gamma–delta T cell receptor (TCR $\gamma\delta^+$ lymphocytes) typically constitute

1–15% of the peripheral blood lymphocytes and are evenly distributed throughout the lymphoid system (21). TCR $\gamma\delta^+$ T lymphocytes accumulate in the granulomatous lesions of leprosy (22), proliferate in response to *Mycobacterium tuberculosis* (23) and *Plasmodium falciparum* (24), and are increased in HIV (25) and EBV (26) infection. Some TCR $\gamma\delta^+$ T lymphocytes recognize hsp (27, 28), while TCR $\gamma\delta^+$ clones may display cytolytic activity (29–31). It has been postulated, therefore, that TCR $\gamma\delta^+$ T lymphocytes have a distinct role in the primary immune response to infectious organisms (reviewed in Ref. 32). However, TCR $\gamma\delta^+$ T lymphocytes may proliferate *in vivo* against endogenous hsp and lead to autoimmune processes (33).

TCR $\gamma\delta^+$ T lymphocytes normally express CD2, CD3, and CD7, but surface CD8 is variable and CD4 is usually absent (21). Despite the presence of multiple human V δ genes, the V δ 1 and V δ 2 gene products form the two major subsets of TCR $\gamma\delta^+$ cells (34, 35), and the V δ 2 protein almost always forms a surface heterodimer with the V γ 9 gene product (36). V δ 2⁺/V γ 9⁺ T lymphocytes specifically proliferate against purified protein derivative (PPD) of *Mycobacterium tuberculosis* and, to a lesser extent, to recombinant purified hsp (37). It is likely that this limited peripheral blood repertoire is

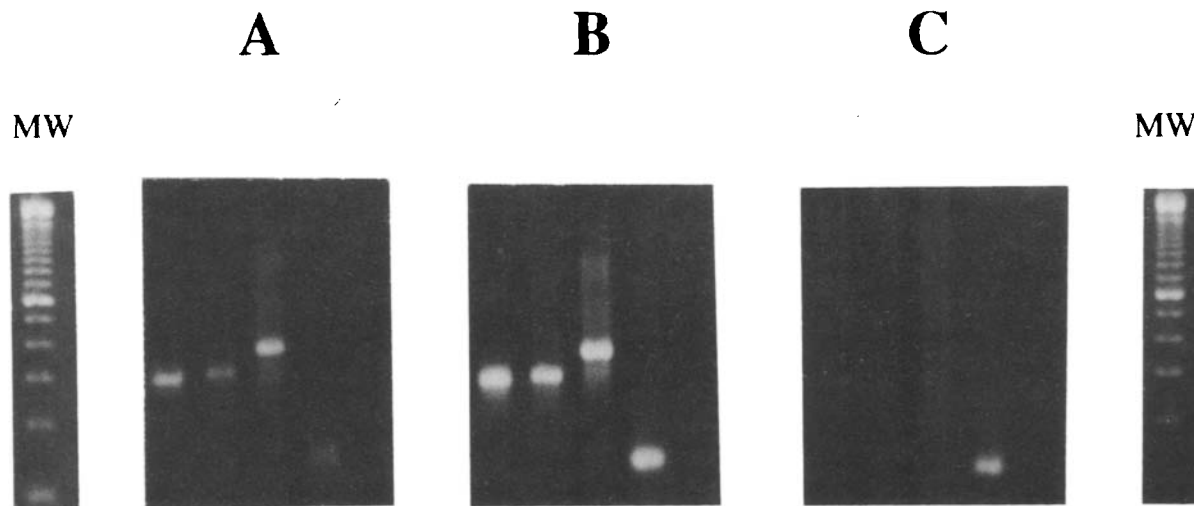


Fig. 5. Analysis of V-J gene rearrangements in the TCR $\gamma\delta$ + T lymphocytes of children with ITP. As the immunophenotype had demonstrated exclusive surface expression of the V δ 2 antigen, we performed RT-PCR using a V δ 2 primer with either a J δ 1 (lane 1), J δ 3 (lane 2), or C δ primer (lane 3). The positive control (lane 4) used primers for the CD7 gene, while a negative control is shown in lane 5. Molecular weight (MW) markers on each side have a bright band at 600 bp for size reference. (A) Patient 1, (B) patient 2, and (C) another child with chronic ITP who had normal numbers of circulating TCR $\gamma\delta$ + T lymphocytes. Both V δ 2-J δ 1 and V δ 2-J δ 3 gene rearrangements were present in each child, indicating that the expanded TCR $\gamma\delta$ + T lymphocyte population was not monoclonal.

due to postthymic selection and stimulation of this specific subset.

To understand T lymphocyte abnormalities in children with immune-mediated platelet destruction, we analyzed the peripheral blood of children with acute or chronic ITP. Three children had highly elevated numbers of TCR $\gamma\delta$ + T lymphocytes at initial evaluation, ranging from 37.8 to 48.1% of the total CD3+ T lymphocytes. These expanded cell populations had a common surface immunophenotype which was CD2+, CD3+, CD7+, CD29+, CD4-, and CD56- (Table II, Fig. 2). These TCR $\gamma\delta$ + cells also expressed the V δ 2+/V γ 9+ phenotype almost exclusively, consistent with an enhanced expansion of this subset. Analysis of CD45 isoforms in patients 1 and 2 (with chronic ITP) had predominantly CD45RO+ (memory) TCR $\gamma\delta$ + T lymphocytes. The TCR $\gamma\delta$ + cells at initial evaluation in patient 3 (with acute ITP) were primarily CD45RA+ (naive) cells but converted to the CD45RO+ phenotype. We hypothesize that the TCR $\gamma\delta$ + lymphocytes of patients 1 and 2 had previously encountered antigen and undergone *in vivo* proliferation and expansion. The TCR $\gamma\delta$ + T lymphocytes of patient 3 had recently encountered antigen at the time of diagnosis and initial evaluation and later acquired the memory phenotype.

There was no evidence of generalized immune deficiency which might account for the elevation of

TCR $\gamma\delta$ + T lymphocytes (38). The enhanced proliferative responses of the TCR $\gamma\delta$ + T lymphocytes to mycobacterial antigen and IL-2 are similar to those previously reported (39) and suggest that these cells were activated *in vivo*. As there was no *in vitro* proliferation to recombinant hsp, however, we were unable to identify potential antigens which might have induced *in vivo* proliferation of TCR $\gamma\delta$ + T lymphocytes.

These expanded cell populations do not represent a monoclonal malignant proliferation of TCR $\gamma\delta$ + T lymphocytes. PCR performed on TCR $\gamma\delta$ + T lymphocyte mRNA identified both V δ 2-J δ 1 and V δ 2-J δ 3 gene rearrangements (Fig. 5). Nucleotide sequencing of VDDJ sequences further identified diverse gene rearrangements (Fig. 6), confirming a polyclonal expansion of TCR $\gamma\delta$ + T lymphocytes. Selection and proliferation of lymphocytes which express a single variable gene region, but have extensive junctional diversity, are reminiscent of a superantigen response (40, 41). We speculate that a superantigen (possibly from an infectious organism) may have been the initiating immune stimulus and led to the tremendous polyclonal expansion of V δ 2+/V γ 9+ TCR $\gamma\delta$ + T lymphocytes.

There was a close correlation between the number of TCR $\gamma\delta$ + T lymphocytes and the degree of thrombocytopenia in each of the three children with ITP. Patient 1 had elevated TCR $\gamma\delta$ + T lympho-

	Vδ2	N	Dδ1	N	Dδ2	N	Dδ3	N	Jδ1	
GERMLINE:	TGTGACACC		GAAATAGT		CCTTCCTAC		ACTGGGGGATACG		ACACCGATAAACTC	FRAME
PT 1 CLONES										
23	TGTGAC		GGCG		TTC	G	GGGGA	GACTGAGT	ACACCGATAAACTC	+
30	TGTGACAC				GCTC		GGGGATAC	ACATC	CACCGATAAACTC	-
33	TGTGACAC	TG		CTAC	GG				ACACCGATAAACTC	+
35	TGTGACAC		TCT		TTC	TTC	GGGGAT	TT	ACACCGATAAACTC	+
54	TGTGACAC				A		ACTGGGGATACG	GGGG	ACTC	+
56	TGTGAC				CCGT		ACTGGGGAT	GGAGGGGCGCTTTTCAAGG	CCGATAAACTC	+
57	TGT	C	AAA	CC			CTGGGGATAC	CCC	CACCGATAAACTC	+
67	TGTGA				TCTGTC		ACTGG	AC	CCGATAAACTC	+
70	TGTGAC	TC	AGT				ACTGGGGATACG	CGC	AAACTC	+
71	TGTGACA				T		ACTGGGGAT		ACCGATAAACTC	+
110	TGTGACAC				T		CTGGGGAT	GACCCGA	ACACCGATAAACTC	+
111	TGTGAC				TCC	CT	ACTGGGGGA	CT	ACACCGATAAACTC	+
113	TGTGAC	CACG		TAG		CG	TGGG	CTTCGCCT	ACACCGATAAACTC	+
PT 2 CLONES										
36	TGTGACACC				TT	GCT	ACTGGGGGATA	TTGGGACAAACAAC	ACACCGATAAACTC	+
49	TGTGA	TCACCG		TAG	GG	TTC	T		ACACCGATAAACTC	+
50	TGTGAC					GC	CTGGGG	CCTC	CACCGATAAACTC	+
51	TGTGACAC		AC		TCCTAC		TGGGGATACG	GC	TAAACTC	+
91	TGTGACACC		GTTGGGGC		CCTA	AGCAGG	ACTGGGGAT	GCG	ACCGATAAACTC	+
103	TGTGAC		GAGCCCTCAAT		CTT	GACTAGCG	TGGG	ACCACGTTACA	CGATAAACTC	+
105	TGTGACACC				GAC		GGAT	GCGAGGAT	CCGATAAACTC	+
106	TGTGACAC	GG		TAG	ACGGTAA		CCT		CGATAAACTC	+
108	TGTGACACC				TT	GCT	ACTGGGGGATA	TTCCCAACAACAAC	CACCGATAAACTC	+
118	TGTGA	TGC		AAT	CGGGCA		TCCT		ACACCGATAAACTC	+
PT 3 CLONES										
74	TGTGACACC				GTGGT		ACTGGGGGAT	GCAAAGGGT	CCGATAAACTC	+
76	TGTGACAC				GGT		GGGA	CCCTGTAC	ACCGATAAACTC	+
79	TGTGA				TCG		ACTGGGGATACG	AGGC	TAAACTC	+
80	TGTG		TATC		CCTAC	GG	CTGGG	TTTGGGGG	ACCGATAAACTC	+
81	TGTGACACC				CT		CTGGGGATACG	AGGG	CGATAAACTC	+
84	TGTGACA				ACTT		ACTGGG	ACCTCCTCGGTTG	CCGATAAACTC	+
87	TGTGACAC	GTTAAGC		AGT			ACTGGGGGA	CGG	ACCGATAAACTC	+
97	TGTGAC				GT		ACTGGG	ATAGGGGT	ACCGATAAACTC	+
99	TGTGACAC						ACTGGG	AGGCCGTT	ACCGATAAACTC	+

Fig. 6. Analysis of VDDJC nucleotide sequences in the TCRγδ+ T lymphocytes of children with ITP. To determine if the elevated number of TCRγδ+ T lymphocytes represented an oligoclonal expansion of cells, RT-PCR followed by DNA sequencing was performed for each of the three children. Each sequence used in Vδ2-D-D-Jδ1 rearrangements was unique, indicating a polyclonal expansion of TCRγδ+ T lymphocytes. Similar results were obtained from sequence analysis of Vδ2-D-D-Jδ3 rearrangements.

cytes for several months, along with severe thrombocytopenia. However, following splenectomy, both his platelet count and the number of TCRγδ+ T lymphocytes normalized (Fig. 2). Patient 2, also with chronic ITP, had a definite diminution in the number of TCRγδ+ T lymphocytes as his disease process has ameliorated. Finally, patient 3, with acute ITP, had 48.1% TCRγδ+ T lymphocytes at initial evaluation, but the percentage of TCRγδ+ T cells has nearly normalized coincident with resolution of his thrombocytopenia (Fig. 3). These clinical observations support the hypothesis that the expanded polyclonal population of TCRγδ+ T lymphocytes played a direct role in the pathogenesis of ITP.

The therapeutic use of cyclosporin A (CyA) in Patient 1 was based in part on the *in vitro* observation that CyA inhibited his PBMC proliferation to IL-2 (Table I). Many immunosuppressive effects of CyA on T lymphocytes relate to inhibition of TCR-

mediated activation events (42). However, CyA has been shown not to affect the intrathymic development of TCRγδ+ cells (42, 43). Therapeutic CyA levels actually increased the number of TCRγδ+ T lymphocytes (perhaps by preferentially inhibiting TCRαβ+ lymphocytes) and had no effect on his thrombocytopenia.

The mechanisms by which TCRγδ+ T lymphocytes could mediate or participate in immune-mediated platelet destruction are only speculative. First, TCRγδ+ T lymphocytes may produce cytokines which are important in the pathogenesis of ITP (5). TCRγδ+ cells secrete a variety of cytokines following activation, including IL-2, GM-CSF, and IFN-γ (44-46). Alternatively, TCRγδ+ T lymphocytes could enhance antibody production by autoreactive B lymphocytes. TCRγδ+ cells can help B cells produce specific antibodies (47), even in the absence of nominal antigen (48). Finally, it is possible that TCRγδ+ T lymphocytes produce a

direct cytolytic effect on platelets. TCR $\gamma\delta$ + cells express killer activity against microbial organisms (45, 49), although the ligands which direct this effect are not known. In our study, no PBMC *in vitro* proliferative responses were observed against platelets, although the availability of autologous platelets might have provided a different result. The dramatic normalization of TCR $\gamma\delta$ + cells in patient 1 following splenectomy suggests that these cells were proliferating within the spleen, perhaps in response to platelet antigenic stimulation.

ACKNOWLEDGMENTS

R.E.W. is the recipient of a James S. McDonnell Scholarship in Molecular Medicine. The authors are grateful to Dr. Barton Haynes and Dr. Michael Krangel for helpful discussions and to Kathy Greenwell for secretarial assistance.

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