

Reduced Number and Impaired Function of Circulating $\gamma\delta$ T Cells in Patients with Cutaneous Primary Melanoma

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We studied the peripheral representation, *in vitro* expansion, cytokine production, and cytotoxicity of $\gamma\delta$ T lymphocytes from 23 patients with cutaneous primary melanoma and 28 healthy subjects. We demonstrated that the absolute number and the percentage of circulating $\gamma\delta^+$ T cells were significantly reduced in melanoma patients in comparison with healthy subjects. The decrease was due to a reduction of V δ 2 T cells, whereas the number of V δ 1 T cells was not affected. As a consequence, the V δ 2/V δ 1 ratio was inverted in melanoma patients. A lower percentage of $\gamma\delta^+$ T cells producing tumor necrosis factor- α or interferon- γ was found in melanoma patients. After a 10 d *in vitro* culture, both

the percentage and the expansion index of $\gamma\delta$ T cells, and in particular of V δ 2 subset, were significantly reduced in melanoma patients in comparison with healthy subjects. The cytotoxicity of sorted $\gamma\delta$ T cells against tumor cell lines and the percentage of $\gamma\delta$ T cells producing perforins were preserved in melanoma patients. The numerical and functional impairment of $\gamma\delta$ T cells could contribute to the inadequate immune response found in melanoma patients and offers the potentiality for the planning of new approaches of immune therapy of malignant melanoma. *Key words:* fluorescence-activated cell sorter/human/melanoma/tumor immunity/ $\gamma\delta$ T cells *J Invest Dermatol* 120:829–834, 2003

Various clinical and experimental observations point to the existence of an immunologic host defense in cutaneous malignant melanoma. CD3⁺ T cell receptor (TCR) α/β -expressing lymphocytes are considered the prevailing lymphocyte subset in primary as well as secondary malignant melanoma (Strohal *et al*, 1994). Immunomodulatory therapies with cytokines or adoptive transfer of T cells have accomplished complete or partial tumor regression in melanoma patients. Nevertheless, the immune response is in most cases inadequate to control tumor growth as tumor progression often occurs (Lee *et al*, 1999; Dudley *et al*, 2001). Hence, the coexistence of a cellular immune response in melanoma lesions (Kammala *et al*, 1999), demonstrated by the presence of clonally expanded T cells, and the inability of melanoma-specific killer T cells to arrest tumor growth, remains a major paradox of tumor immunology (Thor Straten *et al*, 1999; Nielsen *et al*, 2000).

T lymphocytes bearing the $\gamma\delta$ TCR represent a minor population of human peripheral lymphocytes (1–10%) the majority of them expressing the V δ 9/V δ 2 TCR and the CD3⁺CD4⁺CD8⁻ phenotype (Groh *et al*, 1989; Poccia *et al*, 1998). The second most frequent subset of peripheral blood $\gamma\delta$ T cells expresses V δ 1 in association with various V γ elements. The ability of $\gamma\delta$ T cells to respond to nonprocessed and nonpeptidic phosphoantigens in a major histocompatibility complex (MHC)-unrestricted manner

are important features distinguishing them from $\alpha\beta$ T cells (Brenner *et al*, 1987; Bukowski *et al*, 1995). $\gamma\delta$ T cells are activated by mycobacterial antigens, such as isopentenyl pyrophosphate (IPP) and related phenyl pyrophosphate derivatives (Costant *et al*, 1994; Tanaka *et al*, 1995; Burk *et al*, 1995), stress-associated heat shock proteins (Fisch *et al*, 1999), as well as by several cytokines, such as interleukin (IL)-2 (Kjeldsen-Kragh *et al*, 1993), IL-12 (Fujimiya *et al*, 1997), or tumor necrosis factor (TNF)- α (Ueta *et al*, 1996; Lahn *et al*, 1998). Other signals, such as MHC class I-related chains A and B, are able to engage the activating receptor, NKG2D, present on V δ 2 T cells, substantially enhancing the TCR-dependent V δ 2 T cell response to nonpeptide antigens (Das *et al*, 2001; Wu *et al*, 2002). Although little is known about the physiologic significance of $\gamma\delta$ T cells, their marked reactivity toward mycobacterial and parasitic antigens, as well as tumor cells, suggests that $\gamma\delta$ T cells play a part in the anti-infectious and anti-tumoral immune surveillance (Poccia *et al*, 1998; Zheng *et al*, 2001). $\gamma\delta$ T cells may contribute to the immune defense against cancer, having at least two important functions, i.e., reactivity to tumor cells, and regulatory interactions with $\alpha\beta$ T cells (Kabelitz *et al*, 1999). $\gamma\delta$ T cells strongly react against certain lymphoma cells, such as Daudi cells, suggesting a cross-reactivity between microbial and tumor-associated antigens (Kunzmann *et al*, 2000). Furthermore, $\gamma\delta$ T cells have been identified among tumor-infiltrating lymphocytes in various cancer types (Kabelitz *et al*, 2000). Once activated, $\gamma\delta$ T cells produce high levels of cytokines and, mainly, IFN- γ and TNF- α (Poccia *et al*, 1997). Mainly because of their cytokine production, $\gamma\delta$ T cells have been proposed to be involved in co-ordinating the interplay between innate and acquired immunity and, in particular, to guide the establishment of acquired immunity contributing to select appropriate antigens and the strategies for their elimination, and to the definition of $\alpha\beta$ T cell responses

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toward T helper (Th) 1 or Th2 phenotype (Fearon and Locksley, 1996; Mak and Ferrick, 1998). The strict relationship between $\gamma\delta$ and $\alpha\beta$ T cells is bidirectional, as demonstrated by the fact that $\gamma\delta$ T cell function may depend on $\alpha\beta$ T cells (Vila *et al*, 1995; Burns *et al*, 1996).

The role of $\gamma\delta$ T cells in the early anti-tumor defense against melanoma has been suggested by the evidence that $\gamma\delta$ T cells may constitute up to 25% of lymphocyte infiltrate in cutaneous primary melanomas, whereas they are not present in metastatic melanoma (Bachelez *et al*, 1992). These cells exert potent cytotoxic activity against autologous tumor cells (Bachelez *et al*, 1992; Nanno *et al*, 1992). In another study, the survival of patients with necrotizing choroidal melanomas was increased with evidence of V γ 1 and V δ 1 TCR⁺ cells (Bialasiewicz *et al*, 1999).

On the basis of the pivotal role that $\gamma\delta$ T cells may have in the immune response against melanoma acting directly on tumor cells and, secondarily, by modulating the phenotype of T cell responses, we evaluated the peripheral representation and the *in vitro* expansion, cytokine production, and cytotoxicity of $\gamma\delta$ T cells from melanoma patients, comparing the results with those obtained in healthy controls. This study demonstrated for the first time an alteration of circulating $\gamma\delta$ T cells in melanoma patients, with a significant decrease of the number of $\gamma\delta$ T cells in the peripheral blood, an altered pattern of cytokine production, and an impaired *in vitro* expansion of these cells. The knowledge about the deterioration of $\gamma\delta$ T cells could account for the melanoma-related alterations of T cell-mediated adoptive responses and may be helpful for the planning of new approaches of immune therapy in melanoma patients.

MATERIALS AND METHODS

Cell preparation and stimulation Human peripheral blood was obtained from 23 melanoma patients (mean age \pm SD 59.6 \pm 16.1 y; median: 64.5 y; range 32–80) and 28 healthy subjects (mean age \pm SD 57.5 \pm 18.9 y; median: 67.0 y; range 32–79). Healthy subjects were volunteers in good and stable clinical condition, and had laboratory parameters in the physiologic range. Melanoma patients have been admitted to the Dermatology Unit of the I.N.R.C.A. Hospital of Ancona (Italy). Melanoma patients were in good health other than for the existence of melanoma as checked on the basis of clinical and laboratory parameters. Melanoma patients and healthy subjects were equally distributed according to sex and the percentage of male and female inside each group was about the same. The investigations were performed after approval by a local institutional review board. A written informed consent was obtained from each subject. Diagnosis of melanoma was histologically confirmed. All patients brought cutaneous primary nonmetastatic melanoma and were staged according to the new American Joint Committee on Cancer staging system for cutaneous melanoma (Balch *et al*, 2000). In detail, the number of patients and the relative tumor thickness was as follows: \leq 1.0 mm (n = 15), 1.01–2.0 mm (n = 4), 2.01–4.0 mm (n = 2), and \leq 4.0 mm (n = 2). A blood drawing was taken before the surgical excision.

Peripheral blood mononuclear cells (PBMC) were fractionated on Ficoll-Paque (Pharmacia, Uppsala, Sweden) and separated by density gradient centrifugation (400 \times g, 30 min). Cells from the interface of the gradients were washed twice with Ca²⁺ and Mg²⁺-free phosphate-buffered saline (Gibco, Life Technologies, Gaithersburg, MD) and resuspended in RPMI 1640 supplemented with 10% heat-inactivated fetal bovine serum, penicillin (100 U per ml) and streptomycin (100 μ g per ml) (all from Life Technologies, complete medium) at a concentration of 1 \times 10⁶ per ml. The viability was always greater than 98% as determined by trypan blue exclusion. Lymphocytes from both melanoma patients and healthy donors were tested fresh. In each experiment a similar number of melanoma patients and healthy donors was examined. Each donor was tested once and all the tests were carried out with a single blood sample. Mononuclear cells were cultured in the complete medium supplemented with 100 U per ml of IL-2 (Chiron Milano, Italia). Phosphoantigen-specific stimulation of $\gamma\delta$ T cells was performed using the nonpeptidic antigen isopentenylpyrophosphate (30 μ g per ml, IPP; Sigma, St Louis, MO). After 1 wk of culture, the volume corresponding to half the culture medium was replaced by fresh medium. On day 10 of culture, viable cells were counted and used for fluorescence-activated cell sorter analysis and cytotoxicity. The expansion of $\gamma\delta$ T cells was followed by cytometric analysis through double staining of stimulated cells with anti-CD3

[phycoerythrin (PE)] and anti-pan $\gamma\delta$, anti-V δ 1, or anti-V δ 2 T cells [fluorescein isothiocyanate (FITC)] monoclonal antibody (MoAb). The absolute number of $\gamma\delta$ T cells in each culture was calculated as follows: (percentage of $\gamma\delta$ T cells among total cells) \times (total cell count)/100. The $\gamma\delta$ T cell expansion index was then calculated by dividing the absolute number of $\gamma\delta$ T cells in stimulated cultures by the absolute number of $\gamma\delta$ T cells before culture (Poccia *et al*, 1997).

MoAb and fluorescence-activated cell sorter analysis The PE-conjugated MoAb anti-CD3 was purchased from EuroClone (Devon, Wetherby, West York, UK). The FITC-conjugated anti-pan-TCR $\gamma\delta$, anti-TCR V δ 1, and anti-TCR V δ 2 were purchased from Endogen (Boston, MA). IgG1 (Becton Dickinson Milano, Italy) was used as isotype control.

PBMC (1 \times 10⁶) were labeled with 10 μ l of anti-CD3 or anti-TCR V δ 1 MoAb or 5 μ l of anti-pan-TCR $\gamma\delta$ or anti-TCR V δ 2 MoAb in a final volume of 150 μ l of RPMI 1640 with 10% fetal bovine serum for 30 min in ice. At the end of the incubation, cells were washed in PBS, resuspended in Isoton II (Coulter, Euro Diagnostics, Hialeah, FL) and immediately analyzed with a Coulter XL flow cytometer.

Intracellular detection of IFN- γ , TNF- α , and perforin Mononuclear cells were stimulated with IPP and IL-2 for 18 h, and GolgiPlug (a protein transport inhibitor containing brefeldin A; PharMingen, Milton Keynes, U.K.) was added during the last 12 h of culture to block intracellular transport processes and allow cytokine accumulation. Stimulated cells (1 \times 10⁶) were stained with the anti-pan-TCR $\gamma\delta$ MoAb for 30 min at 4°C. Fixation-permeabilization of cells was performed in PBS/2% paraformaldehyde for 15 min at 4°C, followed by incubation for 30 min at room temperature in the dark with PE-conjugated anti-human IFN- γ MoAb or anti-human TNF- α MoAb diluted in PBS, 1% bovine serum albumin, and 0.05% saponin. Cells were finally washed twice in PBS, 1% bovine serum albumin, and 0.01% saponin and analyzed on a XL flow cytometer (Coulter). Perforins were measured after 10 d of incubation using an anti-human perforin-PE (clone δ G9, BD PharMingen) at the same conditions above reported. The controls for unspecific staining included incubation with isotype-matched MoAb (PharMingen).

Isolation of $\gamma\delta$ T lymphocytes and cytotoxic assay $\gamma\delta$ T lymphocytes, *in vitro* expanded with IPP and IL-2 for 10 d, were isolated through cytofluorimetric cell sorting (Vantage, Becton Dickinson). The purity of $\gamma\delta$ T cells, assessed by cytofluorimetric analysis, was greater than 95%.

Cytotoxic assay was performed by a fluorimetric method as recently reported (Provinciali *et al*, 1992). The natural killer resistant cell line Daudi and the natural killer cell sensitive K562 cell line were used as target cells. Daudi is a human lymphoblastoid B cell line derived from a Burkitt lymphoma, which constitutively expresses antigens recognized by V δ 9 V δ 2 T cells. K562 is a human myeloid cell line derived from chronic myelogenous leukemia. The fluorescence was read with a 1420 VICTOR² multilabel counter (Wallac, Turku, Finland). The percentage of specific lysis was calculated as follows:

$$\% \text{ specific lysis} = [(F_{\text{med}} - F_{\text{exp}}) / F_{\text{med}}] \times 100$$

where F represent the fluorescence of the solubilized cells after the supernatant has been removed; med = F from target incubated in medium alone; exp = F from target incubated with effector cells.

Lytic units (LU₂₀/10⁷ cells) were calculated by using a computational method (Bryant *et al*, 1992). One LU corresponded to the number of effector cells required to produce 20% of specific lysis.

Statistical analysis Statistical analysis was performed using parametric (Student's t test) or nonparametric (Mann-Whitney rank sum test) tests on the basis of the distribution of the data. Difference between means were considered significant at p < 0.05. Correlation between $\gamma\delta$ T cell number or expansion and tumor thickness was compared by simple linear regression analysis. The statistical analysis was performed with SigmaStat software version 1.03 (Jandel Scientific, Erkrath, Germany).

RESULTS

Ex vivo analysis of $\gamma\delta$ T lymphocytes Peripheral blood lymphocytes from 23 melanoma patients and 28 healthy subjects were analyzed for the percentage and the absolute number of $\gamma\delta$ T cells through double staining with anti-CD3 and anti- $\gamma\delta$ MoAb. As shown in **Table I** and **Fig 1(A)**, the absolute number of $\gamma\delta$ T cells was significantly reduced in melanoma patients in comparison with healthy subjects (66.1 \pm 33.8 *vs* 92.9 \pm 32.6;

Table I. Absolute number of lymphocytes, $\gamma\delta$ T cells, V δ 1 T cells, and V δ 2 T cells, in healthy subjects and melanoma patients

Donors	Absolute no. ($n \times \text{mm}^3$)				
	Lymphocytes	$\gamma\delta$ T Cells	V δ 1 T Cells	V δ 2 T Cells	V δ 2/V δ 1 ratio
Healthy subjects	2159 \pm 118 ^a	92.9 \pm 32.6	38.5 \pm 13.1	68.2 \pm 35.8	1.8
Melanoma patients	2268 \pm 539	66.1 \pm 33.8*	33.1 \pm 19.4	31.7 \pm 13.7*	0.9

^aData are expressed as mean \pm SD.

* $p < 0.01$ vs healthy subjects.

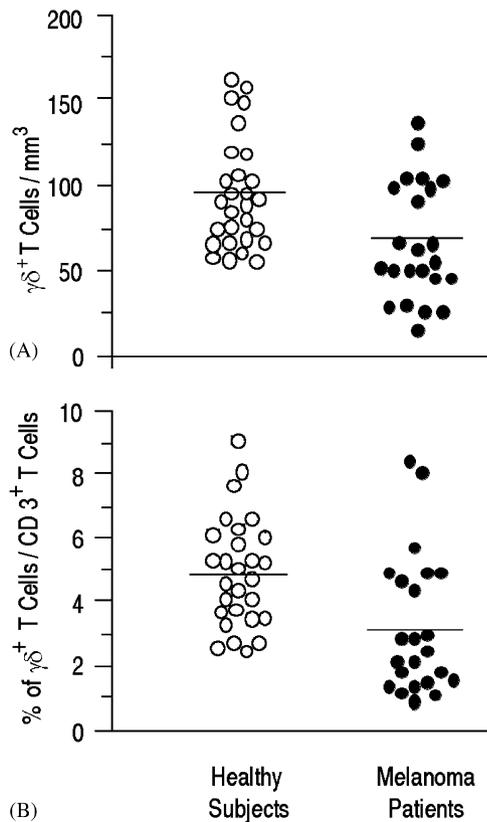


Figure 1. Absolute number and percentage of $\gamma\delta$ T cells in the peripheral blood of melanoma patients and healthy donors. Freshly isolated PBMC from patients with cutaneous primary melanomas ($n = 23$) or healthy subjects ($n = 28$) were double stained with MoAb anti-pan $\gamma\delta$ (FITC) and anti- $\text{CD}3$ (PE) and analyzed by flow cytometry. (A) The absolute number of $\gamma\delta$ T cells, and (B) the percentage of $\gamma\delta$ T cells among total $\text{CD}3^+$ T cells, were both significantly reduced in melanoma patients in comparison with healthy subjects ($p < 0.01$).

$p < 0.01$). The absolute number of peripheral lymphocytes was similar in the two groups of subjects (Table I). As shown in Fig 1(B), the percentage of $\text{CD}3^+$ $\gamma\delta$ T cells in peripheral blood was significantly lower in melanoma patients than in healthy donors (mean \pm SD, 3.2 ± 2.1 vs 4.8 ± 1.7 ; $p < 0.01$). As shown in Table I, the absolute number of V δ 1 T cells did not show significant difference in the two groups of donors (38.5 ± 13.1 and 33.1 ± 19.4 , in healthy subjects and melanoma patients, respectively). Differently, the absolute number of V δ 2 T cells was significantly reduced in melanoma patients in comparison with healthy subjects (31.7 ± 13.7 vs 68.2 ± 35.8 ; $p < 0.01$). The V δ 2 and V δ 1 subsets were differently represented in the two groups: in healthy controls the V δ 2 subset was predominant (V δ 2/V δ 1 ratio = 1.8), whereas in melanoma patients the proportion of V δ 2 and V δ 1 subsets was similar (V δ 2/V δ 1 ratio = 0.9) (Table I). No significant correlation was found plotting the number of $\gamma\delta$ T cells and the tumor thickness of the respective patient.

Cytokine production by $\gamma\delta$ T lymphocytes As it has been demonstrated that activated $\gamma\delta$ T cells produce IFN- γ and TNF- α , we studied the intracellular production of these lymphokines in 1 d stimulated $\gamma\delta$ T cells from healthy subjects and melanoma patients. As shown in Figs 2 and 3, the percentage of $\gamma\delta$ T cells producing IFN- γ was significantly reduced in melanoma patients in comparison with healthy controls (mean \pm SD, 9.3 ± 5.5 vs 18.3 ± 9.5 ; $p < 0.01$). In a similar way, the percentage of $\gamma\delta$ T cells producing TNF- α was lower in melanoma patients than in healthy controls (11.5 ± 3.8 vs 17.9 ± 7.7 for melanoma patients and healthy subjects, respectively; $p < 0.02$).

Expansion of $\gamma\delta$ T lymphocytes The expansion of $\gamma\delta$ T cells from healthy controls and melanoma patients was evaluated after 10 d of culture in the presence of IPP and low-dose IL-2. Both the proportion of $\gamma\delta$ T cells, evidenced by double staining fluorescence-activated cell sorter analysis, and their relative increase in comparison with the $\gamma\delta$ T cell number found on day 0 (expansion index) were evaluated. As shown in Fig 4, the proportion of $\gamma\delta$ T cells reached on day 10 was significantly lower in melanoma patients than in healthy donors (mean \pm SD, 11.9 ± 8.4 vs 36.5 ± 17.6 ; $p < 0.01$). In a similar way, the expansion index of $\gamma\delta$ T cells on day 10 vs day 0 was significantly reduced in melanoma patients in comparison with healthy controls (4.4 ± 3.8 vs 8.6 ± 6.7 ; $p < 0.02$) (Fig 4). As shown in Table II, the expansion index of the V δ 2 subset was significantly lower in melanoma patients than in healthy donors (3.3 ± 3.6 vs 5.8 ± 4.4 ; $p < 0.03$).

Cytotoxicity of $\gamma\delta$ T lymphocytes In order to evaluate the cytotoxic potential of $\gamma\delta$ T cell cultures after *in vitro* expansion, we tested the cell activity of $\gamma\delta$ T cells against Daudi and K562 tumor cell lines and their perforin content through flow cytometry. Purified cultures of activated $\gamma\delta$ T cells (10 d incubation), obtained through cytofluorimetric cell sorting, were cytotoxic against Daudi tumor cells. A great heterogeneity of cytotoxic activity was found in both groups considered. Mean levels of cytotoxicity were similar between healthy subjects and melanoma patients (438.2 ± 141.7 vs 402.5 ± 359.2 , data not shown). The cytotoxic activity of $\gamma\delta$ T cells was also tested against the K562 tumor cell line, with a similar distribution of cytotoxicity in normal donors and melanoma patients (data not shown).

The percentage of $\gamma\delta$ T cells producing perforins among total $\gamma\delta^+$ T cells was not significantly different in melanoma patients in comparison with healthy subjects (42.8 ± 37.6 vs 51.4 ± 8.8 , data not shown).

DISCUSSION

In this study we have demonstrated for the first time the existence of numerical and functional alterations of $\gamma\delta$ T cells from patients with cutaneous primary melanomas. The number of circulating $\gamma\delta$ T cells, the percentage of cells producing IFN- γ or TNF- α , and their *in vitro* expansion, were all decreased in melanoma patients when compared with healthy age-matched subjects.

Both the percentage and the absolute number of circulating $\gamma\delta$ T cells were significantly reduced in patients with cutaneous primary melanoma, thus suggesting a specific deficit for this lymphocyte population. The reduction of $\gamma\delta$ T cell number well

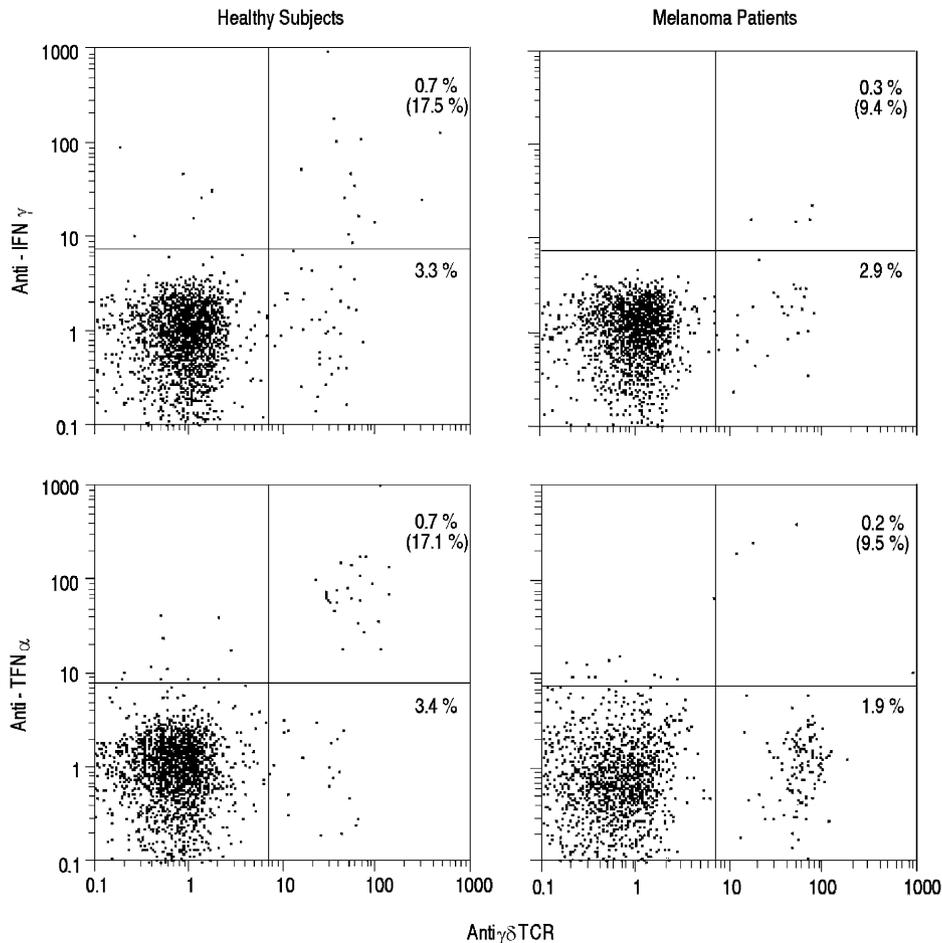


Figure 2. Analysis of cytokine production by $\gamma\delta$ T cells in melanoma patients and healthy donors. PBMC from patients with cutaneous primary melanomas or healthy subjects, were stimulated for 18 h in the presence of IPP (30 μ g per ml) and IL-2 (100 U per ml). The last 12 h of culture were performed in the presence of GolgiPlug, a protein transport inhibitor containing brefeldin. Single-cell analysis of cytokine synthesis in $\gamma\delta$ T cells from a representative healthy subject or melanoma patient was performed following dual staining with cell surface anti- $\gamma\delta$ (FITC) MoAb and intracellular anti-IFN- γ or anti-TNF- α (PE) MoAb. Numbers in brackets indicate the percentages of $\gamma\delta$ cells synthesizing a given cytokine among total $\gamma\delta$ T lymphocytes.

correlated with the decrease of the V δ 2 T cell subset, i.e., the most frequent subset of circulating $\gamma\delta$ T cells (Groh *et al*, 1989; Poccia *et al*, 1998). The V δ 2 population is involved in the reactivity toward microbial antigens and tumor cell antigens (Poccia *et al*, 1998; Zheng *et al*, 2001). The role of V δ 2 T cells in the immune defense against cancer has been demonstrated on the basis of their reactivity against certain lymphoma cells, such as Daudi cells (Kunzmann *et al*, 2000), and for their presence among tumor-infiltrating lymphocytes in various cancer types (Kabelitz *et al*, 2000). On this basis, our data strongly suggest that an impaired $\gamma\delta$ T cell potential may contribute to a lower immune defense against melanoma. Both V δ 1 and V δ 2 T cell populations have been reported to contribute to anti-tumor immunity because of their cytotoxic and Th1-type cytokines producing activities (Wu *et al*, 2002). Furthermore, both $\gamma\delta$ T cell subsets have been found able to respond to stress-induced expression of the MHC class I-related chains A and B (Wu *et al*, 2002). MHC class I-related chains function as ligands for NKG2D, an activating receptor complex that triggers natural killer cells, costimulates CD8 $\alpha\beta$ and V γ 9 δ 2 $\gamma\delta$ T cells, and is required for stimulation of V δ 1 T cells (Groh *et al*, 1999; Das *et al*, 2001; Girardi *et al*, 2001; Wu *et al*, 2002). The role of $\gamma\delta$ T cells in the early anti-tumor defense against melanoma has been previously suggested by the evidence that a significant accumulation of V δ 1 T cells was found in three of 11 primary cutaneous melanomas, with proportions of $\gamma\delta$ T cells ranging from 15 to 25% of lymphocyte infiltrate, whereas $\gamma\delta$ T cells were not present in eight metastatic melanomas

(Bachelez *et al*, 1992). In our study, we did not find numerical changes of the V δ 1 T cell population in melanoma patients in comparison with controls, whereas the V δ 2 subset was significantly reduced in melanoma patients. Almost half of our melanoma patients had an absolute number of circulating $\gamma\delta$ T cells similar or lower than half of the mean number found in controls (Fig 1). Phenotypic analysis of the tumor-infiltrating $\gamma\delta$ T cells revealed that they use the products of the V δ 9 and V δ 2 genes in a way similar to most of circulating TCR $\gamma\delta$. It is noteworthy that $\gamma\delta$ lines or clones developed from lymphocytes infiltrating primary melanoma display a potent autologous MHC-unrestricted tumor cell cytotoxicity (Bachelez *et al*, 1992; Nanno *et al*, 1992). In a more recent study, the presence of $\gamma\delta$ T cells well correlated with the survival of patients with necrotizing choroidal melanoma (Bialasiewicz *et al*, 1999). In our study, not only the number but also the function of $\gamma\delta$ T cells was altered in melanoma patients. The *in vitro* expansion of $\gamma\delta$ T cells, that represent one of the most relevant functional parameters for $\gamma\delta$ T cells, was significantly reduced in patients with cutaneous primary melanomas. This would imply that $\gamma\delta$ T cells from melanoma patients have a decreased proliferative capacity in comparison with healthy subjects. Under normal conditions, $\gamma\delta$ T cells respond to antigen challenge by secreting large quantities of TNF- α and IFN- γ (Poccia *et al*, 1997; Kabelitz *et al*, 2000), which contribute to the activation of both specific and aspecific immune responses. In this study, we show that the percentage of $\gamma\delta$ T cells producing either TNF- α or IFN- γ is significantly reduced in melanoma

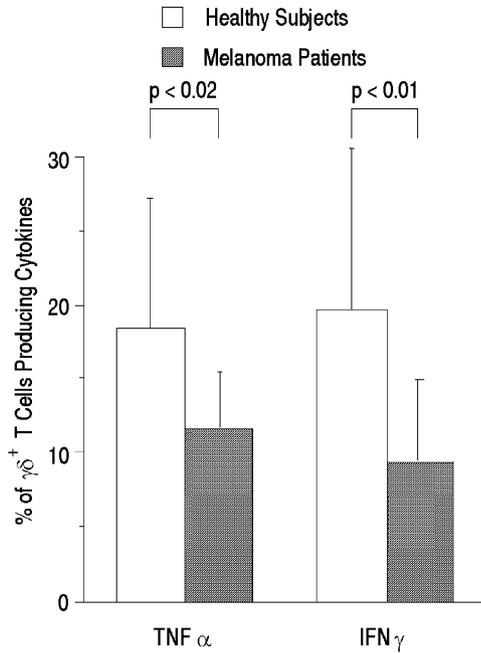


Figure 3. Percentage of $\gamma\delta$ T cells producing cytokines in melanoma patients and healthy donors. $\gamma\delta$ T cells from patients with cutaneous primary melanomas or healthy subjects were activated as reported in Fig 2 and analyzed for cytokine production after dual staining with cell surface anti-pan $\gamma\delta$ (FITC) MoAb and intracellular anti-IFN- γ or anti-TNF- α (PE) MoAb. The mean percentage of $\gamma\delta$ T cells producing either IFN- γ or TNF- α was significantly lower in melanoma patients in comparison with healthy subjects ($p < 0.01$ and $p < 0.02$ for IFN- γ or TNF- α , respectively).

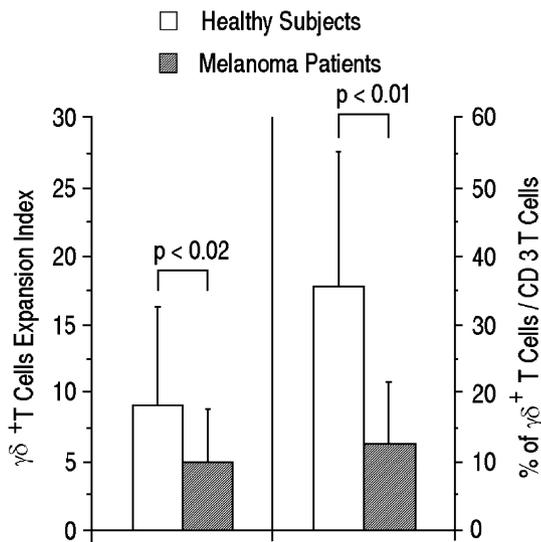


Figure 4. Evaluation of $\gamma\delta$ T cell expansion in melanoma patients and healthy donors. PBMC from patients with cutaneous primary melanomas or healthy subjects were stimulated for 10 d in the presence of IPP (30 μ g per ml) and IL-2 (100 U per ml). The expansion index (left panel) and the percentage (right panel) of *in vitro* expanded $\gamma\delta$ T cells among total CD3 $^+$ T cells were significantly lower in melanoma patients in comparison with healthy subjects ($p < 0.02$ and $p < 0.01$).

Table II. Expansion index of $\gamma\delta$ T cells and V δ 2 T cells in healthy subjects and melanoma patients.

Donors	Expansion Index ^a	
	$\gamma\delta$ T cells	V δ 2 T cells
Healthy subjects	8.6 \pm 6.7 ^b	5.8 \pm 4.4
Melanoma patients	4.4 \pm 3.8*	3.3 \pm 3.6**

^aThe expansion index was calculated by dividing the absolute number of $\gamma\delta$ T cells in stimulated cultures by the absolute number of $\gamma\delta$ T cells before culture.

^bData are expressed as mean \pm SD.

* $p < 0.02$ and ** $p < 0.03$ vs healthy subjects.

to the inefficient immune defense against melanoma with at least two different mechanisms. One of these is based on the lower $\gamma\delta$ T cell potential, determined by the reduced number and expansion of $\gamma\delta$ T cells, and particularly of V δ 2 subset, found in melanoma patients. In this case, the impairment of $\gamma\delta$ T cells may represent a new mechanism by which melanoma tumor cells escape an efficacious immune response. The second mechanism is based on the regulatory interactions between $\gamma\delta$ T cells and $\alpha\beta$ T cells. Several studies have demonstrated that the type and the functioning of the specific lymphocyte response is dependent on signals provided by the innate recognition system (Fearon and Locksley, 1996; Medzhitov and Janeway, 1997). Indeed, even if the recognition of foreign or nonself proteins is controlled by the rearranging genes that encode specific receptors, the functional outcome of most responses to pathogens and the initiation of the response itself is determined by the type of innate immune response they elicit (Fearon and Locksley, 1996; Medzhitov and Janeway, 1997). Recent evidence has suggested that one of the crucial players involved in the regulation of innate and acquired immunity is the $\gamma\delta$ T cell population (Mak and Ferrick, 1998). There is substantial cross-talk between $\gamma\delta$ and $\alpha\beta$ T cells. It has been demonstrated that the proliferative response of human peripheral blood $\gamma\delta$ T cells towards microbial antigens or Daudi tumor cells, depends on helper signals provided by CD4 $^+$ $\alpha\beta$ T cells (Burns *et al*, 1996a,b). Moreover, some $\gamma\delta$ T cell function may depend on $\alpha\beta$ T cells, as, for example, the $\gamma\delta$ T cell proliferation in response to activated CD4 $^+$ $\alpha\beta$ T cells (Burns *et al*, 1996a,b), and the antigen presentation by CD4 $^+$ $\alpha\beta$ T cells to $\gamma\delta$ T cells (Vila *et al*, 1995; Collins *et al*, 1998). On the other hand, substantial evidence has suggested that $\gamma\delta$ T cells regulate certain $\alpha\beta$ T cell-mediated immune responses, pointing the definition of $\alpha\beta$ T cell responses toward a Th1 or Th2 phenotype (Fearon and Locksley, 1996; Mak and Ferrick, 1998; Kabelitz *et al*, 2000). This fact is particularly related by the cytokines secreted by $\gamma\delta$ T cells that, in turn, are able to mediate both innate and acquired immunity (Ferrick *et al*, 1995). We propose that, in analogy with what has been recently demonstrated by us in aged people and centenarians (Argentati *et al*, 2002), the numerical and functional impairment of $\gamma\delta$ T cells demonstrated in this study could determine a derangement in the establishment of acquired immunity, with consequent difficulties in selecting appropriate antigens and the strategies for their elimination. Further studies will be performed to investigate the specific causes involved in the impairment of $\gamma\delta$ T cells in primary cutaneous melanoma patients. At present, we may only suggest that an impairment of $\gamma\delta$ T cells may determine a lower immune defense against melanoma which, in turn, may predispose to melanoma. The healthy and stable conditions of melanoma patients examined in this study make improbable that the $\gamma\delta$ T cell defect is secondary to the malignancy, even if this possibility may not be definitively excluded. Finally, the fact that approximately 10% of patients with melanoma have family histories of the disease, suggesting a genetic susceptibility (Platz *et al*, 1997), raises the question on whether the differences we have found might be attributable to the genetic background of the melanoma patients. We do not think this is the case, as melanoma patients and controls enrolled in our study were identified in the

patients. Instead, the lytic activity of $\gamma\delta$ T cells, evidenced by their cytotoxicity toward Daudi and K562 tumor cells and their production of perforins, seems to be well preserved in melanoma patients.

On the basis of the data reported in this paper, we suggest that the impairment of $\gamma\delta$ T cell number and function may contribute

same geographical area (Ancona area in Italy), and melanoma patients did not have family histories of melanoma.

In conclusion, the demonstration of a numerical and functional derangement of $\gamma\delta$ T cells, which we have found in patients with cutaneous primary melanomas, make them a potentially useful tool for the planning of new approaches of immune therapy in malignant melanoma.

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