# Analysis of V $\delta$ 1 T cells in clinical grade melanoma-infiltrating lymphocytes

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Abbreviations: ACT, adoptive T-cell therapy; AJCC, American Joint Committee on Cancer; CTLA-4, cytotoxic T lymphocyte antigen 4; ICS, intracellular cytokine staining; LAC, leukocyte activation cocktail; RECIST, response evaluation criteria in solid tumors; REP, rapid expansion protocol; SEB, Staphylococcal enterotoxin B; TCR, T-cell receptor; TIL, tumor-infiltrating lymphocyte

 $\gamma\delta$  T cells, including V $\delta$ 1 and V $\delta$ 2 T cells, can recognize tumor-associated ligands neglected by conventional  $\alpha\beta$  T cells in a MHC-independent manner. Little is known regarding the anticancer potential and the possibility to isolate and expand V $\delta$ 1 T cells to therapeutically relevant numbers. In this study, we have detected low frequencies of V $\delta$ 1 T cells among tumor-infiltrating lymphocyte (TIL) products for adoptive cell transfer generated from melanoma metastases. An increased frequency of V $\delta$ 1 T cells was found among the cell products from patients with an advanced disease stage. V $\delta$ 1 T cells displayed in vitro antitumor activities and sufficient proliferative potential to generate over 1 × 10<sup>9</sup> cells using current protocols for T cell transfer. Infusion of V $\delta$ 1 T cells together with high numbers of  $\alpha\beta$  TILs in a clinical trial was safe and well tolerated. These data suggest that V $\delta$ 1 T cells should be further scrutinized as a potentially useful tool for the treatment of patients with metastatic melanoma.

## Introduction

Two major subsets of T cells carrying the  $\gamma\delta$  T-cell receptor (TCR) have been described. T cells expressing the Vδ2 gene usually account for more than 90% of the circulating  $\gamma\delta$  T-cell pool (representing about 1-10% of human peripheral lymphocytes), while intraepithelial  $\gamma\delta$  T cells more commonly express the V $\delta$ 1 gene.<sup>1</sup> Although no major differences exist relative to effector functions between T cells expressing the  $\alpha\beta$  TCR and  $\gamma\delta$  T cells, the latter are capable of recognizing tumor-associated ligands that are neglected by conventional  $\alpha\beta$  T cells in an MHC-independent manner.<sup>2</sup> Most  $\gamma\delta$  T cells lack the surface expression of CD4 and CD8, in agreement with their non-MHC restricted recognition of unconventional antigens. Many preclinical studies and clinical trials have focused their attention on  $V\gamma 9V\delta 2$  T cells, as they can be easily isolated from the peripheral blood of most individuals and activated with conventional drugs such as synthetic phosphoantigens and aminobisphosphonates.<sup>2</sup> However, little is known regarding the anticancer potential of  $\gamma\delta$ T cells expressing the V $\delta$ 1 gene.

It is currently believed that, due to their extremely limited diversity, V $\delta$ 1 T cells may not respond to a diversity of microbial antigens but rather to unique "stress antigens" that are markers

of cell infection or transformation. V $\delta$ 1 T cells reside mainly within epithelial tissues such as the intestinal epithelium and epidermis, where they might provide a first line of immunosurveillance against malignancy<sup>1,3</sup> by recognizing ligands such as MHC Class I-related molecules (i.e., MICA and MICB), whose expression can be induced in response to infection, injury or cellular transformation.<sup>4–6</sup> These molecules have no role in the presentation of peptide antigens, but may function themselves as tumorassociated antigens.<sup>7</sup> Of note, similarly to  $\alpha\beta$  T cells, V $\delta$ 1 T cells can inhibit tumor cell growth and recruit other immune cells by releasing a number of cytokines, including tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) and interferon  $\gamma$  (IFN $\gamma$ ).<sup>8–9</sup>

Because of the intraepidermal location of normal melanocytes, we considered melanoma an appropriate model to study the antitumor properties of V $\delta$ 1 T cells. In addition, previous studies have reported a variable infiltration of melanoma lesions with  $\gamma\delta$ T cells.<sup>10–12</sup> These observations prompted us to analyze clinical grade tumor-infiltrating lymphocyte (TIL) products from patients with metastatic melanoma. TILs were prepared according to current clinical protocols for adoptive T-cell transfer (ACT). Low frequencies of  $\gamma\delta$  T cells were found among TILs obtained from most patients, and V $\delta$ 1 T cells from patients with high  $\gamma\delta$  T-cell frequencies were further functionally characterized.

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Patient n°	Sex	AJCC stage	Previous systemic treatments	<b>Biopsy for TIL generation</b>	γδ+ (%)	Vδ1+ (%)
1	F	IV	None	LN	0.15	0.12
2	М	IIIC	None	SC	0.012	0.01
3	М	IIIC	None	LN	0.02	0.02
4	F	IV	IL-2/IFNα	LN	0.03	0.01
5	М	IV	IL-2/IFN $\alpha$ , DC vaccination	LN	0.84	0.37
6	М	IV	IL-2, DC vaccination	SC	0.12	0.11
7	F	IV	DC vaccination	SC	BLD	BLD
8	М	IV	IL-2/IFN $\alpha$ , anti-CD137, DC vaccination	LN	0.31	0.08
9	F	IIIB	None	LN	BLD	BLD
10	F	IIIC	None	SC/LN	0.02	BLD
11	F	IV	IL-2/IFN $\alpha$ , temozolomide, anti-CTLA4	SC	94.14	94
12	F	IIIC	None	LN	0.41	0.25
13	F	IIIC	None	LN	0.09	BLD
14	М	IIIC	None	LN	BLD	BLD
15	М	IV	IL-2/IFN $\alpha$ , anti-CTLA4, DC vaccination	LN	0.01	0.01
16	М	IV	IL-2/IFN $\alpha$ , anti-CTLA4	SC	3.08	2.98
17	F	IV	IL-2/IFN $\alpha$ , anti-CTLA4, DC vaccination	LN	0.25	0.15
18	F	IV	IL-2/IFNα	LN	10.67	7.8
19	М	IV	IL-2/IFN $\alpha$ , anti-CTLA4	SC	0.17	0.1
20	Μ	IV	IL-2/IFN $\alpha$ , anti-CTLA4	LN	2.47	2.03
21	Μ	IV	IL-2	LN	0.172	0.002
22	Μ	IV	IL-2/IFN $\alpha$ , anti-CTLA4	SC	3.57	2.86
23	F	IIIC	None	LN	BLD	BLD
24	F	IIIC	None	LN	0.11	0.03
25	М	IIIC	None	SC	BLD	BLD
26	М	IIIC	None	LN	2.37	0.48
27	F	IV	IL-2/IFN $\alpha$ , anti-CTLA4	SC	1.37	1.29

**Table 1.** Summary of patient characteristics and clinical grade TILs

AJCC, American Joint Comittee on Cancer; anti-CD137 (experimental treatment in protocol); BLD, below the limit of detection (< 0.001%); DC vaccination, dendritic cell vaccination (experimental protocol); IFN $\alpha$ , interferon  $\alpha$ ; IL-2, interlukin-2; LN, lymph node metastasis; SC, subcutaneous metastasis; TIL, tumor-infiltrating lymphocyte.

## **Results**

 $\gamma\delta$  T cells in clinical grade TIL products from melanoma. Twenty-seven clinical grade TIL products generated from an equal number of patients with AJCC melanoma Stage III (n = 11) or Stage IV (n = 16) were available for analysis. All such products contained > 97% of CD3<sup>+</sup> cells (data not shown). Patients and TIL characteristics are summarized in Table 1. Among TILs from 22 out of 27 patients a low frequency of CD4<sup>-</sup>CD8<sup>-</sup> (double negative, DN)  $\gamma\delta$  T cells was present. V $\delta$ 1 T cells were detected among TILs from 20 out of 27 patients (Table 1). The V $\delta$ 1<sup>+</sup> subset represented the majority of  $\gamma\delta$  T cells detected, accounting for 60 ± 35%.

TILs from patients with AJCC Stage IV disease contained significantly more V $\delta$ 1 T cells than those from Stage III patients (Fig. 1A). However, 15 out of 16 Stage IV patients included in this study had received prior systemic immune-based therapies (Table 1), while none of the patients with Stage III disease had received these treatments. The most common therapy

administered prior to tumor specimen collection was interleukin-2 (IL-2) (15/16 patients). Therefore, we cannot rule out a potential influence of immunotherapies on V $\delta$ 1 T cell infiltration.

No differences were observed between TIL products obtained from subcutaneous or lymph node metastases (Fig. 1B). However, stratification of patients with Stage IV disease based on their previous treatment with anti-CTLA4 antibodies showed a non-significant trend toward an increase of V $\delta$ 1 T cells in patients who had previously received this treatment (p = 0.09) (Fig. 1C). A typical plot showing the detection of V $\delta$ 1<sup>+</sup> T cells in a representative TIL culture is depicted in Figure 1D.

Antitumor activity of V $\delta$ 1 T cells. V $\delta$ 1 T cells from two patients selected for high frequency of V $\delta$ 1 T cells were stimulated with autologous (patient 11 and 18) or a panel of allogeneic HLA-A matched and HLA-A unmatched (patient 11) melanoma cell lines followed by intracellular cytokine staining (ICS) for type 1 cytokines, to assess whether they were able to recognize tumor cells and exert effector functions. V $\delta$ 1 T cells from both patients efficiently recognized and generated antitumor responses



**Figure 1.** Vô1 T cells in clinical grade tumor-infiltrating lymphocytes (TILs). (A–C) Frequency of detected Vô1 T cells in clinical grade TIL products grouped for AJCC disease stage (**A**), biopsy origin (**B**) and prior treatment with anti-CTLA4 antibodies (only patients with Stage IV disease are shown) (**C**). (**D**) FACS plot from a representative TIL product (from patient 17). An electonic gate was set on the CD3<sup>+</sup>CD4<sup>-</sup>CD8<sup>-</sup> live cell population. LN, lymph node metastatis; SC, subcutaneous metastasis.

against autologous target cells with production of Type I cytokines (Fig. 2A), although the frequency of responding cells was low (below 1% in both cases). Additionally, V $\delta$ 1 T cells from patient 11 recognized both HLA-A matched and unmatched allogeneic melanoma cell lines (Fig. 2A). Concomitant antitumor responses of  $\alpha\beta$  T cells (CD8<sup>+</sup> or CD4<sup>+</sup> T cells) were detected in the same products stimulated with autologous tumors (Fig. 4).

IFNγ production by cells from patient 11 (exhibiting > 90% of V $\delta$ 1<sup>+</sup> cells, see **Table 1**) was further confirmed by ELISPOT (Fig. 2B). Furthermore the melanoma cell line (from patient 11) inducing the highest V $\delta$ 1 T-cell reactivity, as assessed by ICS, was chosen for testing V $\delta$ 1 T cell killing abilities. Data confirmed that V $\delta$ 1 T cells possess cytotoxic capacity (Fig. 2C). Recognition of healthy non-transformed cells by V $\delta$ 1 T cells was excluded, because of the absent or very low (< 0.1%) cyto-kine production when the TILs from patient 11 were incubated with allogeneic irradiated peripheral blood mononuclear cells (PBMCs) from two different donors (data not shown).

Unspecific activation of V $\delta$ 1 T cells. Since the tumor-specific in vitro responses that we detected appeared to be quite low in frequency, we asked whether unspecific stimulating agents would induce a response in a larger fraction of V $\delta$ 1 cells. The Staphylococcal enterotoxin B (SEB) is a bacterial superantigen that is believed to stimulate T cells in an oligoclonal fashion depending on the expression of specific variable region gene elements in the  $\beta$  chain of the TCR ( $V_{\beta}$ ).<sup>15–17</sup> However, it has been reported that also cells bearing the  $\gamma\delta$  TCR are capable to respond to a number of bacterial superantigens including SEB, and may therefore be involved in local immune responses to such antigens as they mimic bacterial infections.<sup>18</sup> In addition, V $\delta$ 1 T cells isolated from patients with colorectal cancer or with multiple sclerosis have been proposed to efficiently respond to SEB stimulation.<sup>19–20</sup>

Based on these premises, we have tested the ability of SEB to induce Type 1 cytokine production in TIL products from the two patients displaying a high frequency of V $\delta$ 1<sup>+</sup> T cells (patients 11 and 18). In this setting, SEB failed induce any response, whereas  $\alpha\beta$  T cells in the same TIL products responded strongly. In contrast, stimulation with the leukocyte activation cocktail (LAC) led to the production of Type 1 cytokines on > 95% of V $\delta$ 1 T cells (**Fig. 2D**), demonstrating the intrinsic capacity of the vast majority of this cell population to generate meaningful T<sub>b</sub>1-like responses.

Phenotype and proliferative capacity of Vô1 T cells. Clinical grade Vô1 TILs from both patients analyzed (patients 11 and 18) displayed an activated (CD44<sup>+</sup>CD69<sup>+</sup>) phenotype (data not shown). In addition, they expressed the activating receptor NKG2D (Fig. 3A) and (about 70 and 100% of all



**Figure 2.** Antitumor activity. (**A**) Responses of V $\delta$ 1 T cells (from patient 11, > 90% of V $\delta$ 1 T cells) evaluated with the production of tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) and interferon  $\gamma$  (IFN $\gamma$ ) when unstimulated (control), or stimulated with autologous or an HLA-A-unmatched allogeneic melanoma cells. Both constitutive responses and responses upon stimulation of cancer cells with 100 IU/mL IFN $\gamma$  for 72 h (+ IFN $\gamma$ ) are shown. All the plots are gated on V $\delta$ 1+ T cells. (**B**) IFN $\gamma$  ELISPOT. Unstimulated tumor-infiltrating lymphocytes (TILs,control) or TILs stimulated with autologous tumor cells are shown. (**C**) Percentage of cytotoxicity at different effector:target ratios of unfractionated TILs vs. one HLA-A-unmatched allogeneic melanoma cell line. (**D**) Production of TNF $\alpha$  and IFN $\gamma$  by unstimulated (control) TILs or TILs unspecifically stimulated with the Staphylococcal enterotoxin B (SEB) or PMA/Ionomycin.

Vol T cells from patient 11 and 18, respectively) the neural cell adhesion molecule 1 CD56 (Fig. 3B). CD56 expression has previously been suggested as a specific signature of highly efficient

antitumor V $\delta$ 2 T cells.<sup>21</sup> Therefore, we examined whether this was also the case for V $\delta$ 1 T cells, testing TILs from patient 11 against autologous tumor cells. Surprisingly, cells endowed with

**Figure 3.** Phenotypic and proliferative characteristics. (**A**)  $V\delta 1$  T cells expressed NKG2D. (**B**) CD56 was expressed by a large fraction of  $V\delta 1$  T cells, and cells with in vitro anticancer activity were enriched in the CD56<sup>-</sup> compartment. Dotted light gray line: isotype control. (**C**) From day 8 to day 10 of the rapid expansion protocol (REP), undivided cells were 85% of  $\alpha\beta$  T cells vs. 45% of  $V\delta 1$  T cells. Light gray,  $\alpha\beta$ T cells; dark gray,  $V\delta 1$  T cells.

antitumor effector capacities were much enriched in the CD56<sup>-</sup> population (Fig. 3B). Moreover, we compared the proliferative capacity of V $\delta$ 1 T cells to that of  $\alpha\beta$  T cells during the rapid expansion protocol (REP) by means of the PKH dilution assay. Data indicate that V $\delta$ 1<sup>+</sup> T cells proliferate significantly more than  $\alpha\beta$  T cells in the same culture, with 45% undivided cells vs. 85% (Fig. 3C).

Involvement of MHC, IFNy treatment, NKG2D or MICA/B in tumor recognition. Additional experiments confirmed that both autologous and allogeneic tumor recognition of V $\delta$ 1 T cells from patient 11 were not dependent on MHC, as cancer cell preincubation with a combination of antibodies blocking MHC Class I and Class II molecules did not influence V $\delta$ 1 T cell recognition (Fig. 4A). As a positive control, these antibodies were shown to almost completely abrogate  $\alpha\beta$  T-cell responses (Fig. 4B). In line with these observations, the pretreatment of cancer cells with a known inducer of MHC expression as well as of the antigen-processing machinery such as IFN $\gamma$  (for 72 h at 100 IU/mL), which according to our recent data are able to increase the frequency of tumor-reactive  $\alpha\beta$  T cells in clinical grade TIL products,<sup>28</sup> was not able to foster anticancer responses by Vô1 T cells (Fig. 2A).



To get further insights into the elements underlying the recognition of cancer cells by V $\delta$ 1 T cells, experiments with antibodies blocking NKG2D on TILs or MICA/B on autologous and allogeneic target cells before stimulation were performed (on TILs from patient 11). Neither NKG2D nor MICA/B blockade did appear to significantly influence V $\delta$ 1 T-cell tumor recognition and responses (data not shown).

Association of V $\delta$ 1 T cells infusion and clinical response to ACT. Eleven samples analyzed in this study derived from cell

products infused into patients with Stage IV melanoma in the context of the ACT trial NCT00937625. The average number of infused V $\delta$ 1 T cells was 1.4 ± 2.5 × 10<sup>9</sup> (average of total T-cells infused > 50 × 10<sup>9</sup>). No major cell infusion-related side effects were observed.<sup>29</sup>

To date, ten patients were evaluated for clinical response. Three patients experienced an objective response (three complete responses, all ongoing to date). There were no differences between responding vs. non-responding patients for the frequency



**Figure 4.** MHC involvement in target recognition. (**A and B**) Tumor-infiltrating lymphocytes (TILs) were stimulated with autologous tumor cells preincubated with isotype control or a combination of MHC Class I and II-blocking antibodies. While MHC Class I and Class II-blocking antibodies significantly affected tumor cell recognition by  $\alpha\beta$  T cells (**B**), they did not modify tumor cell recognition by  $V\delta$ 1 T cells (**A**).

of V $\delta$ 1 T cells in infusion products, but the sample size was small (n = 10, p = 0.83). Notably, one patient achieving a complete response (patient no 18, see **Table 1**) was infused with 7.8% V $\delta$ 1 T cells (approximately 6.5 × 10<sup>9</sup> cells in total). However, the cell product contained also high numbers of tumor-specific CD8<sup>+</sup> T cells (data not shown).

## Discussion

The large-scale application of novel immunotherapies such as the adoptive transfer of ex vivo expanded TILs after host lymphodepletion has the potential to significantly improve the prognosis of patients with metastatic melanoma.<sup>22</sup> In this report, we showed that the products for infusion generated with current techniques are composed of a heterogeneous population of T cells, which contain not only CD4<sup>+</sup> and CD8<sup>+</sup> T lymphocytes (as previously reported) but also a small and in some cases significant fraction of  $\gamma\delta$  T cells, with a prevalence of the V $\delta$ 1 subset. Similar to a previous report,<sup>10</sup> we demonstrated that V $\delta$ 1 T cells are present in most infiltrates of human melanoma. In addition, we found that these cells exerted non MHC-restricted antitumor effector functions toward both autologous and allogeneic melanoma and that, importantly, they could be efficiently expanded with current clinical scale methods for ACT. Indeed, detectable amounts of Vô1 T cells in TILs after REP were found in 20 out of 27 patients analyzed, and ACT infusion products from ten patients contained on average more than  $1 \times 10^9$  of these cells. On the contrary, other melanoma-infiltrating immune cells such as natural killer (NK) cells do not efficiently

proliferate using these expansion methods (namely REP) and are not commonly detected among clinical grade TILs.<sup>13,23</sup>

We detected an increased frequency of Vô1 T cells in TIL products from patients with AJCC Stage IV compared with Stage III disease. This difference is in apparent contrast with the findings obtained by Bachelez et al.,10 who reported a higher frequency of Vol T cells in primary than in metastatic melanoma specimens. However, the settings considered are certainly different as the samples used in our study were all represented by regional or distant metastatic tumor sites. Thus, different expression of surface molecules on malignant melanocytes at subsequent stages of tumor progression, leading to the recruitment of different lymphocyte subsets, may underlie the observed discrepancy. Moreover, over 90% of patients with AJCC Stage IV disease in our study were treated with immunotherapies before specimen collection, namely, about 85% received IL-2 and 50% were also treated with anti-CTLA4 antibodies. Therefore, the possibility that immunotherapies would induce a relative increase of V $\delta$ 1 T-cell over  $\alpha\beta$  T-cell infiltration cannot be ruled out, in particular for IL-2, which represented the most common treatment administered to our group of patients.

Several reports pointed out that CD56 is an important effector marker of NK,  $\alpha\beta$  and  $V\gamma9V\delta2$  T cells.<sup>21,24–27</sup> Our data indicated that this may not apply to V $\delta1$  T cells, as tumor-reactive V $\delta1$  T cells were enriched in the CD56<sup>-</sup> compartment. This assumption needs to be validated as these results were obtained with TILs from a single patient.

In conclusion, the demonstration that V\delta1 T cells can exert antitumor effector functions in vitro coupled with data showing

that large numbers (up to  $6.5 \times 10^9$  cells) of these cells can be safely transferred into patients with metastatic melanoma suggest that V $\delta$ 1 T cells may represent a potentially useful therapeutic tool that should be further scrutinized. In addition, given their theoretical role in immunosurveillance, strategies for the stimulation of V $\delta$ 1 T-cell function may be translated into adjuvant treatments for patients at high risk of relapse.

# Materials and Methods

TIL products and melanoma cell lines. All the procedures were approved by the Scientific Ethics Committee for the Capital Region of Denmark. Written informed consent was obtained from patients before any procedure according to the Declaration of Helsinki. Tumor specimens of at least 1 cm<sup>3</sup> were obtained from patients with melanoma AJCC Stage III or IV undergoing standard-of-care surgical procedures or specimen collection for enrolment in a clinical trial (clinicaltrials.gov identifier: NCT 00937625). Clinical grade TIL products were generated from tumor fragments obtained from patients with metastatic melanoma with a 2 step protocol including a slow expansion IL-2 containing media (pre-REP phase), and a REP, as described previously.<sup>13</sup> Melanoma cell lines were generated from tumor fragments, as described.<sup>13</sup>

Flow cytometry and antitumor activity of TILs. The following fluorochrome-conjugated antibodies were used for flow cytometry: FITC-conjugated TCR  $\delta$  TCS1 (TCR1055, Thermo-Fisher), CD4 (345768), CD44, CD69 (347823); PE-conjugated CD56 (345812), pan- $\gamma\delta$  TCR (333141), PECy7-conjugated IFN- $\gamma$  (557643); PerCP-conjugated CD8 (345774), NKG2D (FAB139C, R&D Systems); APC-conjugated TNF $\alpha$  (554514). Unless otherwise specified, antibodies used for flow cytometry were from BD. Fixation/Permeabilization Buffer (00–5223–56 and 00–5123–43), Permeabilization Buffer (00–8333–56) and Fixable Viability Dye eFluor 450 (65–0863–14) were from Ebiosciences, GolgiPlug (555029) from BD and SEB (S4881) from Sigma-Aldrich. Leukocyte Activation Cocktail with GolgiPlug (550583, LAC; containing phorbol 12-myristate 13-acetate/ionomycin and brefeldin A) was from BD.

Evaluation of antitumor activity by ICS was performed as previously described.<sup>13</sup> For blocking experiments, target or

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effector (for NKG2D blockade) cells were incubated for 30 min at 37°C with blocking antibodies. Antibodies used were anti-HLA-ABC (M0736), anti-HLA-DR, DP, DQ (M0775, both from Dako), anti-NKG2D (MAB139, clone 149810, R&D Systems), anti-MICA/MICB (320909, clone 6D4, Biolegend). The number of IFN $\gamma$ -secreting TILs was quantified by IFN  $\gamma$  ELISPOT, as previously described.<sup>13</sup> To assess the cytotoxic abilities of V $\delta$ 1 T cells, a flow-cytometry based assay was applied as previously described.<sup>14</sup> A small modification of the assay was introduced by replacement of carboxyfluorescein succinimidyl ester with PKH-26 (PKH26GL-1KT, Sigma-Aldrich) and propidium iodide with 7-actinomycin D (559925, BD).

Analysis of cell proliferation. To assess the proliferation of V\delta1 T cells, TILs were stained with PKH26 Red Fluorescent Cell Linker at day 8 of REP according to the manufacturer's instructions. Red fluorescence from the CD4<sup>+</sup> or CD8<sup>+</sup> cell populations and from the CD4<sup>-</sup>CD8<sup>-</sup> cells was assessed after 48 h. As negative control (no proliferation) and to set the gate of the undivided cells, a PKH-stained cell sample was stored at 4°C for the entire period of incubation. Samples were analyzed using a BD FACSCanto II flow cytometer. Analysis was performed with BD FacsDiva Software. Dead cells were excluded from the analysis based on 7-actinomycin D positivity.

Assessment of clinical responses. Responses to ACT regimens were assessed with standard RECIST criteria in the context of a pilot clinical trial of T-cell therapy for patients with advanced melanoma (clinicaltrials.gov identifier NCT00937625).<sup>29</sup>

Statistical analyses. Data from different groups were compared with the non parametric two-tailed Mann-Whitney test. All the analyses were performed with Graph Pad Prism 5 software (Graph Pad Software Inc.).

## Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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