Chemotherapy accelerates immune-senescence and functional impairments of \( V\delta^2 \)pos T cells in elderly patients affected by liver metastatic colorectal cancer

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**Abstract**

Human (gamma delta) \( \gamma\delta \) T cells are unconventional innate-like lymphocytes displaying a broad array of anti-tumor activities with promising perspectives in cancer immunotherapy. In this context, \( V\delta^2 \)pos T cells represent the preferential target of several immunotherapy protocols against solid tumors. However, the impact of both aging and chemotherapy (CHT) on \( V\delta^2 \)pos T cells is still unknown. The present study evaluates with multi-parametric flow cytometry the frequencies, terminal differentiation, senescence and effector-functions of peripheral blood and tumor infiltrating \( V\delta^2 \)pos T cells purified from liver metastases (CLM) of patients affected by colorectal cancer (CRC) compared to those of sex- and age-matched healthy donors. The peripheral blood of CLM patients underwent CHT is characterized by decreased amounts of \( V\delta^2 \)pos T cells showing a relative increase of terminally-differentiated \( CD27^{neg}/CD45RA^{pos} \) (TEMRA) cells. The enrichment of this latter subset is associated with an increased expression of the senescent marker \( CD57 \). The acquisition of \( CD57 \) on TEMRA \( V\delta^2 \)pos T cells is also coupled with impairments in cytotoxicity and production of TNF-\( \alpha \) and IFN-\( \gamma \). These features resemble the acquisition of an immune-senescence profile by \( V\delta^2 \)pos T cells from CLM patients that received CHT, a phenomenon that is also associated with the loss of the co-stimulatory marker CD28 and with the induced expression of CD16. The group of CLM patients underwent CHT and older than 60 years old showed higher frequencies of \( CD57^{pos} \) and TEMRA \( V\delta^2^{pos} \) T cells. Similar results were found for tumor infiltrating \( V\delta^2^{pos} \) T cell subset purified from CLM specimens of patients treated with CHT. The toxicity of CHT regimens also affects the homeostasis of \( V\delta^2^{pos} \) T cells by inducing higher frequencies of circulating \( CD57^{pos} \) TEMRA subset in CLM underwent CHT and younger than 60 years old. Taken together, our data demonstrate that the enrichment of senescent \( V\delta^2^{pos} \) T cells in CLM patients is not only induced by patients’ aging but also by the toxicity of CHT that further accelerates the accumulation of \( CD57^{pos} \) TEMRA cells highly dysfunctional in their anti-tumor activities. These results are important to both predict the clinical outcome of CLM and to optimize those protocols of cell cancer immunotherapy employing unconventional \( V\delta^2^{pos} \) T cells.

**Keywords:** \( \gamma\delta \) T cells, Immune-senescence/Aging, Cancer, Chemotherapy
**Introduction**

Human γδ T lymphocytes are divided in the two main Vδ1\textsuperscript{pos} and Vδ2\textsuperscript{pos} subsets on the basis of their TCRδ-chain repertoire. While Vδ1\textsuperscript{pos} cells preferentially localize in mucosal tissues and skin, Vδ2\textsuperscript{pos} T cells are mainly enriched in peripheral blood (PB) where they represent about 5% of all circulating T cells. The activation of Vδ2\textsuperscript{pos} T cells relies on the recognition of non-peptidic compounds (i.e. microbial or stress- or tumor-induced "phosphoantigens") in association with butyrophilin 3A1 (BTN3A1 also known as CD277). Besides the TCR interactions with phosphoantigens/ BTN3A1 complexes, several Natural Killer Receptors (NKRs) are involved in triggering the anti-tumor functions of Vδ2\textsuperscript{pos} T cells, with the C-lectin type NKG2D playing a major role [1, 2].

The differential expressions of CD27 and CD45RA surface markers identify different Vδ2\textsuperscript{pos} T cell subsets: CD27\textsuperscript{pos}/CD45RA\textsuperscript{neg} naïve cells (T\textsubscript{Naïve}), CD27\textsuperscript{pos}/CD45RA\textsuperscript{neg} central memory (T\textsubscript{CM}), CD27\textsuperscript{neg}/CD45RA\textsuperscript{neg} effector-memory (T\textsubscript{EM}) and the terminally-differentiated (T\textsubscript{EMRA}) CD27\textsuperscript{neg}/CD45RA\textsuperscript{pos} cells. These Vδ2\textsuperscript{pos} T cell subsets diverge not only for their maturation/differentiation status, but also for proliferative capacities, effector functions and resistance to cell death in response to antigens and/or cytokine stimulations [3].

Growing evidences highlighted the high impact of Vδ2\textsuperscript{pos} T cells in cancer immune-surveillance with promising perspectives in cancer immunotherapy [4, 5]. In this context, two main clinical approaches have been employed to boost anti-tumor activities of Vδ2\textsuperscript{pos} T cells. The first one activates them through the in vivo administration of either IL-2 or synthetic nitrogen-containing bisphosphonates (NBPs) drugs that, in turn, induce intracellular accumulation of phosphoantigens. A second strategy relies on adoptive transfers of Vδ2\textsuperscript{pos} T cells expanded in vitro with several methodologies such as the activation with zoledronate [5, 6]. However, these procedures showed both experimental and clinical limits and many efforts are currently being implemented to further improve the effector-functions and persistence in vivo of Vδ2\textsuperscript{pos} T cells. In this context, cellular senescence is certainly one of the main issues to solve considering that age-related changes of T cells greatly impair their capacity to expand and proliferate, thus leading to dysfunctional immune responses against tumors and pathogens [7].

The shift to senescence and accumulation of mature T cells physiologically occur after 60 years old when both αβ and γδ T lymphocytes lose their costimulatory molecules (i.e. CD27 and CD28), acquire terminally-differentiated T\textsubscript{EM} and T\textsubscript{EMRA} phenotypic profiles, express high constitutive levels of the senescence marker CD57 and shorten their telomerase lengths [8–11]. However, it is still controversial whether CD57 can be used as a single marker to identify senescent Vδ2\textsuperscript{pos} T cells regardless of differential expression of CD27 and CD45 [3, 11, 12].

Aging is certainly a major burden for social health systems in the industrialized countries as the populations are longer exposed to several pro-tumorigenic risk factors. This leads to a significant higher incidence of cancer onsets in the 6th, 7th and 8th decades of life [13]. Hence, there are rising numbers of elderly patients undergoing anti-cancer conventional chemotherapies (CHT), whose high toxicities greatly hamper both duration and quality of life. In this regard, several lines of clinical and experimental evidence pointed out that these anti-neoplastic treatments further accelerate immune-cell senescence, thus representing negative prognostic factors in aging and worsening the overall clinical outcomes of cancer patients [14, 15].

Since the use of Vδ2\textsuperscript{pos} T cells is currently considered one of the most promising tools in cancer immunotherapy [4, 5], understanding the exact impact of CHT on their immune-senescence is key to better predict the clinical outcomes of cancer in elderly and to optimize those therapeutic protocols targeting these highly cytotoxic unconventional T cell effectors. Colorectal cancer (CRC) represents the 3rd most frequent solid cancer and more than 50% of CRC patients undergo hepatic dissemination of the primary tumor. The gold-standard therapeutic approach of CRC patients with liver metastasis (CLM) is the surgical removal of hepatic secondary lesions after neoadjuvant combination CHT with or without biological therapy (BT) (Table 1) [16, 17]. Moreover, a higher infiltration of competent immune cells in tumor mass greatly improves the prognosis of CLM patients and increases their overall survival (OS) [18, 19]. Here, we analyze the impact of conventional CHT regimens on the homeostasis and effector-functions of Vδ2\textsuperscript{pos} T cells in a cohort of CLM elderly patients.

**Methods**

**Patients and specimen collections**

Biological specimens from CLM patients underwent CHT (n = 58), or from CHT naïve patients (n = 13) and aged- and sex-matched healthy donors (n = 40) (Table 1). Patients’ recruitment was performed according to the Declaration of Helsinki and the protocol had been approved by the Institutional Review Board (IRB) of Humanitas Research Hospital (HRH) (Approval N.168/18). All enrolled patients signed the related consent forms. Liver specimens and peripheral blood mononuclear cells (PBMCs) were isolated and stored as we previously described [19, 20].

**Flow cytometry**

Absolute γδ T cell counts were performed on 100 μl of fresh PB stained with following anti-human monoclonal...
antibodies (mAbs): CD3 (SK7; BV605) and CD45 (H130; AF700) (BioLegend) and Vδ2 (IMMU-389; FITC) (Beckman Coulter). We then used CountBright™ Absolute Counting Beads (Invitrogen) according to the manufacturer’s instructions.

For both regular and intracellular staining, γδ T cells were first screened for viability with Zombie Aqua™ Fixable Viability kit (BioLegend) and then processed as previously described [20]. The following mAbs were used: CD28 (CD28.2; PE-Cy7) (BioLegend); Vδ2 (B6; BUV395), CD3 (UCHT1; BUV661), CD45RA (H100; BUV737), CD16 (348; BUV496) (BD); CD57 (REA769; PE-Vio615) (Miltenyi); CD27 (0322; APCeFluor780) (eBioscience). The intracellular amounts of TNF-α (Mab11; PE) and IFN-γ (B27; Bv711) (BD) as well as the frequency of cytotoxic CD107a+ cells (H4A3, PE) (BD Biosciences) was evaluated after stimulating γδ T cells with Phorbol myristate acetate (PMA; 0.5 μg/mL) and L-glutamine (0.1 μg/mL) (Sigma Aldrich).

Flow cytometry experiments were performed on FACS Symphony™ (BD). All data and t-SNE® algorithm were analyzed with FlowJo Software (version 9.6) (FlowJo LLC) using single stained controls BD CompBeads™ (BD).

### Statistical analyses

The data were assessed by non-parametric Mann-Whitney U (unpaired) or Wilcoxon (matched-paired) tests by using GraphPad Prism version 7. For all correlation analysis Pearson’s coefficient was applied. Statistically significant p values were represented with GraphPad (GP) style and summarized with following number of asterisks (*): *P ≤ 0.05; **P ≤ 0.01; ***P ≤ 0.001; ****P ≤ 0.0001.

### Results

Vδ2+ T cells were gated within viable CD3+/CD45+ lymphocytes and their absolute counts are significantly lower in the PB of CLM patients underwent CHT compared to those of healthy donors (Fig. 1a-b). We then analyzed the surface expression of CD27 and CD45RA to track the differentiation and distribution of Vδ2+ T cell subsets. Our data showed a significant increase of Vδ2+ TEMRA in CLM patients underwent CHT (28.9 ± 20.6%) compared to healthy controls (9.4 ± 6.4%). This phenomenon is associated with the previous administration of CHT, as the frequency of circulating Vδ2+ TEMRA in those CLM patients naïve for CHT (16.7% ± 12.6) is similar to that of healthy donors and significantly lower to that of CLM patients undergone CHT (41.6% ± 19.6). The increased amounts of Vδ2+ TEMRA in CLM patients treated with CHT is counterbalanced by a significant decrease of Vδ2+ TCM in the same patients compared to their counterparts naïve for CHT (Fig. 1c-d-e). The great impact of neoadjuvant CHT in shaping the distribution of Vδ2+ T cell subsets in CLM patients is also confirmed by our findings showing that the number of CHT cycles (8.7 ± 2.7) inversely correlates with the percentages of PB Vδ2+ TCM, while not affecting at
all the overall frequencies of PB Vδ2pos T EMRA (Fig. 1f). This latter dichotomy reflects the different homeostatic status of Vδ2pos T CM compared to that of Vδ2pos T EMRA, as the first subset is composed of proliferating lymphocytes high susceptible to the toxicity of those chemotherapy compounds that kills all dividing cells without any specificity against tumor blasts. Instead, T EMRA Vδ2pos cells are terminally differentiated and not proliferating effectors resistant to CHT, thus explaining their high frequency even after several cycles of neoadjuvant anti-tumor chemotherapies.

The relative increased frequency of PB T EMRA Vδ2pos in CLM patients underwent CHT correlates with their higher expression of CD57. Notably, the expression of this latter marker of immune senescence follows the terminal differentiation of Vδ2pos T cells. Indeed, the frequency of PB CD57pos T EMRA Vδ2pos T cells resulted significantly higher compared to that of CD57pos T CM Vδ2pos T cells that, in turn, showed significantly higher amounts of CD57 when compared to T CM Vδ2pos T cells (Fig. 2c). Taken together, these data indicate that the PB of CLM patients underwent CHT is highly enriched of senescent CD57pos T EMRA Vδ2pos T cells dysfunctional in their anti-tumor effector functions.

To assess the impact of patients’ aging in the higher frequencies of CD57pos and T EMRA Vδ2pos T cells in

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**Fig. 1** Frequency and distributions of peripheral blood Vδ2pos T cell subsets in patients affected by liver metastasis of colorectal cancer and underwent chemotherapy. a) Representative dot plot flow cytometric graphs showing the gating strategy of viable CD45pos/CD3pos/Vδ2pos T lymphocytes. b) Statistical dot plot graph showing the absolute number of CD3pos (left) and Vδ2pos (right) T cells per 1 mL of blood in healthy donors (n = 12; mean age: 49.3 ± 9.5) and CLM patients underwent CHT regimens (n = 16; mean age: 51.5 ± 8.1). c) Representative dot plot flow cytometric graph (c) and pie charts (d and e) showing respectively the distribution and the percentages of CD27pos/CD45RApos T Naive (upper right in dot plot graph and light green in pie charts), CD27pos/CD45RApos central memory (T CM) (upper left in dot plot graph and gray in pie charts), CD27pos/CD45RApos effector-memory (T EM) (lower left in dot plot graph and purple in pie charts) and terminally-differentiated CD27neg/CD45RApos (T EMRA) (lower right in dot plot graph and orange in pie charts) Vδ2pos T cell subsets. Pie charts compare the frequencies of Vδ2pos T cell subsets between healthy donors (n = 34; mean age: 51.7 ± 10.8) with age-matched CLM patient underwent CHT (n = 33; mean age: 51.5 ± 8.1) as well as between CLM patients naïve for CHT (n = 13; mean age: 69.5 ± 8.1) and age-matched CLM patients underwent CHT (n = 41; mean age: 70.1 ± 6.5). f) Statistical analysis showing the Pearson correlations between the frequency (%) of either T CM (left) or T EMRA (right) Vδ2pos T cells with the number of CHT cycles (mean number: 8.7 ± 6.5) administered to patients affected by CLM (n = 40).
CLM patients underwent CHT, we divided this cohort in subjects younger or older than 60 years old. Our data confirmed that the age-induced immune-senescence significantly increases the percentages of both CD57$^{\text{pos}}$ and T$_{\text{EMRA}}$ V$\delta$2$^{\text{pos}}$ T cells in those patients > 60 years old (Fig. 2d). We also showed that CHT alone induces immune-senescence regardless of patients’ age. Indeed, the percentage of CD57$^{\text{pos}}$ T$_{\text{EMRA}}$ V$\delta$2$^{\text{pos}}$ cells resulted significantly higher in those CLM patients underwent CHT and younger than 60 years old compared to that of age-matched healthy donors (Fig. 2e). These data clearly indicate that both CHT and aging play synergic roles in the regulation of V$\delta$2$^{\text{pos}}$ T cell homeostasis in CLM patients with the final result of greatly accelerating their terminal differentiation towards a senescent CD57$^{\text{pos}}$/T$_{\text{EMRA}}$ subset highly impaired in its anti-tumor effector-functions. We also demonstrate here that the acquisition of CD57 inversely correlates with the surface expression of CD28 while being associated with increased surface amounts of CD16 (Fig. 2f).

Although the overall frequency of tumor infiltrating V$\delta$2$^{\text{pos}}$ T cells purified from CLM specimens is not
affected by the administration of CHT, we found a significant increase of CD57 expression on those Vδ2pos T cells from patients underwent CHT compared to naïve ones (Fig. 3c). Similar to their PB counterparts, the frequency of CD57pos/Vδ2pos T cells is significantly higher in elderly CHT patients ≥60 years old (Fig. 3d). Consistently with these data, t-SNE analysis showed also in CLM specimens of patient administered with CHT an increased frequency of age-related tumor infiltrating CD57pos/CD28neg/CD16pos TEMRA Vδ2pos T lymphocytes (Fig. 3e).

Discussion
The present study is aimed to measure the true impact of conventional CHT regimens on unconventional T cell senescence in elderly cancer patients, since the toxicity of conventional anti-tumor therapies greatly impairs their ability to clearance malignant cells [7, 12, 14, 15]. In particular, we focused our investigations on circulating Vδ2pos T cells that are endowed with great anti-tumor potentials currently being targeted by several protocols of cancer immunotherapies [4–6]. Our data showed that CLM patients underwent CHT, although showing lower absolute counts of circulating Vδ2pos T cells, retain high relative frequencies of terminally differentiated and senescent CD57pos/CD28neg/CD16pos TEMRA Vδ2pos T cells greatly impaired in their effector functions. This latter subset is resistant to the toxicity exerted by repeated CHT cycles administering DNA-damaging drugs that, in contrast, are highly toxic against less differentiated and still proliferating TCM Vδ2pos T cells.
The preferential accumulation in PB of senescent CD57\textsuperscript{pos} T\textsubscript{EMRA} V\textsuperscript{δ2pos} cells in CLM patients underwent CHT is associated with two major mechanisms. The first one is linked to natural immune-senescence of people aging as the incidence of many cancers is higher in patients $\geq$ 60 years old. In this context, liver metastatic CRC is one of the most common causes of cancer deaths worldwide with a higher incidence in elderly [16, 17]. Indeed, our cohort of recruited CLM subjects had a mean age of 61 $\pm$ 10.7 years old and both the frequencies of CD57\textsuperscript{pos} and T\textsubscript{EMRA} V\textsuperscript{δ2} T cell subsets resulted higher in that fraction of patients older than 60 years. The second mechanism is associated with a direct toxicity of CHT on immune cells, as also highlighted by several studies both in pediatric and geriatric cancer patients [14, 15, 22]. As a matter of fact, we show here that the expression of CD57 on T\textsubscript{EMRA} V\textsuperscript{δ2pos} cells is much higher on those CLM patients underwent CHT and younger than 60 years old compared to age-matched healthy donors. This demonstrates that neoadjuvant CHT induces immune senescence also on unconventional T cells regardless of CLM patients’ age. Notably, high frequencies of impaired CD57\textsuperscript{pos}/T\textsubscript{EMRA} V\textsuperscript{δ2pos} cells were able to persist in PB of CLM patients even after 6 weeks from the completion of the last CHT cycle and before surgical removal of liver metastases. Further prospective studies are required to assess how long senescent and functional impaired V\textsuperscript{δ2pos} T cells survive after CHT and what clinical impact they have on the OS of CLM patients. In this regard, it has been already reported that the enrichment of circulating subsets of CD57\textsuperscript{pos} $\alpha\beta$ T cells represents a negative prognostic factor in the clinical outcome of gastrointestinal cancers [23].

Our study also contributes to better characterize immune-senescence of V\textsuperscript{δ2pos} T cells, since it has been recently reported that expression of CD57 can define alone their senescent status without the need of also evaluating the expression of both CD27 and CD45RA [11]. This represents a key point that is currently being debated both in physiological and pathological conditions. We found that, at least in a human cancer setting, the expression of CD57 on senescent V\textsuperscript{δ2pos} T cell parallels their terminal differentiation towards T\textsubscript{EMRA} (CD27\textsuperscript{neg}/CD45RA\textsuperscript{pos}), a phenomenon associated with the loss of CD28 and the acquired expression of CD16. These results are in line with a previous study showing that V\textsuperscript{δ2pos} T\textsubscript{EMRA} are refractory to phosphoantigen stimulation, but rather respond to activation via FcγRIII [21].

The majority of cancer patients are older than 65 years old in line with population aging [14]. In this context, several clinical trials in the elderly are currently being implemented to optimize the anti-tumor activities of unconventional T cells. These therapeutic protocols are mostly aimed to expand V\textsuperscript{δ2pos} T cells both in vivo and in vitro [6]. Hence, a better understanding of the mechanisms accelerating immune-senescence in aging is fundamental to boost the effector-functions of these cytotoxic and cytokine-producer T lymphocytes. We show here that, neoadjuvant CHT regimens, although absolutely required to reduce tumor mass in CLM patients before surgery, greatly speed the senescence of V\textsuperscript{δ2pos} T cells in synergy with aging of cancer patients. This knowledge will allow us to better optimize immune-therapies against cancers in elderly. Indeed, senescence process can be reversed through the inhibition of p38 mitogen-activated protein kinase (MAPK) signaling [24]. This methodology could be then approached to develop new protocols implementing pre-treatment with MAPK inhibitors in elderly patients with CRC [25]. Alternatively, new methodology can be implemented in vitro to select and expand CD57\textsuperscript{neg}/V\textsuperscript{δ2pos} T cells that better resist to the terminal differentiations and senescence induced by CHT. Further studies are also required to better identify those CHT associated accumulation of impaired and senescent circulating V\textsuperscript{δ2pos} T cells.

**Abbreviations**
CHT: Chemotherapy; CLM: Colorectal liver metastatic cancer; CRC: Colorectal cancer; NBPs: Nitrogen-containing bisphosphonates; T\textsubscript{CM}: Central memory T; T\textsubscript{EM}: Effector memory T; T\textsubscript{EMRA}: Terminally differentiated T; T\textsubscript{Naïve}: Naïve T; t-SNE: t-Distributed Stochastic Neighbor Embedding; $\gamma\delta$: Gamma delta T

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**Authors’ contributions**
JM and DM developed the study. EB, VC, GS, and GL performed the experiments and statistical analyses. MC, MD, and GT recruited patients and collected biological specimens. JM, EB, FD, and DM contributed to the interpretation of the data and wrote article. All authors read and approved the final manuscript.

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**Availability of data and materials**
The dataset generated and analyzed in the current study are available from the corresponding authors on reasonable request.

**Ethics approval and consent to participate**
The collection of human samples for research purposes was ethically approved by the Institutional Review Board (IRB) of Humanitas Research Hospital (HRH) (Approval 168/18).

**Consent for publication**
Not applicable.

**Competing interests**
The authors declare that they have no competing interests.
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