# CD137+ T-Cells

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The CD137 receptor is expressed by activated antigen-specific T-cells. CD137<sup>+</sup> T-cells were identified inside TILs and PBMCs of different tumor types and have proven to be the naturally occurring antitumor effector cells, capable of expressing a wide variability in terms of TCR specificity against both shared and neoantigenic tumor-derived peptides. The aim of this review is thus summarizing and highlighting their role as drivers of patients' immune responses in anticancer therapies as well as their potential role in future and current strategies of immunotherapy.

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### **1. Introduction**

Immunotherapy aims to re-educate the patient's immune system to recognize and fight cancer cells. The existence of T-cells with a potential antitumor effect has laid the foundation for most of the current approaches of immunotherapy. In fact, the use of therapies such as immune checkpoint inhibitors (ICIs), DC vaccines, and adoptive T-cell transfer (ACT) finally relies on the presence of a population of effector T-cells that is capable of killing tumor cells. These immune-based drugs thus aim to unleash this population from different regulatory constraints such as T-cell exhaustion or the impossibility of reaching cancer cells, to subsequently limit tumor growth and progression. As a confirmation, the accumulation of tumor-infiltrating lymphocytes (TILs) correlates with a better clinical outcome and an improved survival in most tumor models <sup>[1][2][3][4][5][6][7][8][9][10][11]</sup>, indicating their importance in predicting patients response to anticancer therapies. Nevertheless, the composition of TILs is heterogeneous <sup>[12]</sup> and it still remains challenging to identify the real population of naturally occurring antitumor T-cells <sup>[13]</sup>. Therefore, this review will discuss the emerging role of the CD137<sup>+</sup> T-cells population as the main effector population activated against cancer cells with all the possible implications for the future of immunotherapy.

# 2. CD137: The Receptor

The CD137 receptor (4-1BB, TNFRSF9) is a member of the tumor necrosis factor receptors (TNFR) family and was characterized as an inducible costimulatory receptor on T-cells, together with its ligand (CD137L, 4-1BBL), both in human and mice <sup>[14]</sup>. CD137 was initially described as a surface marker expressed by activated T-cells, with an in vitro peak expression 48 h after the primary T-cell activation signal and a decline starting from day 4–5 <sup>[14][15]</sup> <sup>[16][17]</sup>. In vivo, its expression upon activation turned out to happen even earlier, starting indeed at 12h post-immunization <sup>[18][19]</sup>. Both CD4<sup>+</sup> and CD8<sup>+</sup> T-cells are able to upregulate CD137, even if its expression on CD8<sup>+</sup> T-cells is earlier and higher <sup>[20][21][22]</sup>.

However, CD137 receptor is not a specific marker for T-cells, since it can be expressed, even if to a smaller extent, also by dendritic cells (DCs), monocytes, natural killer (NK) cells, eosinophils, and microglia <sup>[23]</sup>. On the other side, CD137L is expressed by activated antigen presenting cells (APC) as macrophages, DCs, and B-cells <sup>[20][23][24][25]</sup>. Therefore, it is reliable to suppose that the engagement between CD137 and its ligand is part of the complex pathways of interactions between APCs and T-cells.

Similarly to other members of the TNFR family, the CD137 receptor relies on TRAFs proteins to build its signaling <sup>[27]</sup>. The binding of both CD137L <sup>[28][29]</sup> and agonistic antibodies <sup>[30]</sup> results in a quick recruitment of TRAF1 and TRAF2 to the receptor. The consequent TRAF-mediated activation of NF-kB and MAPK intracellular signaling, leads to T-cell division and proliferation, an increased cell survival and enhanced effector functions in both CD4<sup>+</sup> and CD8<sup>+</sup> T-cells <sup>[15]</sup>. As for CD137 expression, also TRAF1 expression is induced by T-cell activation, confirming that the CD137-induced signalosome is required for cytotoxic T-cells (CTL) expansion and for the boosting of effector functions <sup>[27]</sup>. In fact, a number of mice experiments proved that CD137 stimulation is able to increase Tcell proliferation and cytokine production [14][23]. Consistently, in absence of the CD28 signal, T-cells treated with an anti-CD3 and CD137L can proliferate and produce interleukin 2 (IL-2) to a similar extent of those treated with the combination of anti-CD3 and anti-CD28, but just in the presence of a consistent antigen stimulation [31][32][33][34]. This evidence was one of the starting points to get to the notion that CD137 identifies those T-cells that are activated against a specific antigen. The CD137L stimulation of human CD8<sup>+</sup> T-cells leads to the expansion of this T-cells subset which is followed by an increase of effector molecules such as granzyme A, interferon  $\chi$  (IFN- $\chi$ ), perforin, and different cytokines, driving CD8<sup>+</sup> memory T-cells toward a differentiated effector phenotype <sup>[35][36]</sup>. In addition, the CD137 receptor seems to have a strong and prevalent role in increasing T-cell survival by preventing activation-induced T-cell death [37][38] and this appears to be in line with the physiological timing of the CD137 signal that is subsequent to the TCR and CD28 mediated signals. The BIM downregulation and the induction of Bcl-XL and Bfl-1 were pointed out as responsible for the inhibition of the activation-induced cell death, after the CD137 engagement [39]. Further studies also showed that the CD137 engagement is able to stimulate the mitochondrial metabolism in order to increase T-cell respiratory capacities [40][41] and to induce DNA demethylation in CD8<sup>+</sup> T-cells main genes and chromatin reprogramming [42]. As above mentioned, different studies highlighted a preferential role of CD137 in CD8<sup>+</sup> T cells rather than in CD4<sup>+</sup> T cells, even if it can be induced on both the T-cell subsets, including CD4<sup>+</sup> regulatory T-cells (Tregs) [43]. However, the effective function of CD137 signaling on CD4<sup>+</sup> T-cells is still unclear and may thus not be as physiologically relevant as for CD8<sup>+</sup> T cells <sup>[44]</sup>. As confirmation, mice deficient for CD137 show an impaired antiviral response mediated by CD8<sup>+</sup> T cells [45][46][47]. Similarly, mice deficient for CD137L that were adoptively transferred with OT-1 derived CD8<sup>+</sup> T-cells, showed a marked reduction of these OVA-specific T-cells in both the late primary response and the secondary expansion to OVA/LPS <sup>[19]</sup>. Moreover, when anti-CD137 monoclonal antibodies or CD137L injections were tested in cancer therapy, a significant benefit in terms of antitumor response was noticed [21][24][38][48][49][50][51][52][53][54][55][56][57]. When CD137 was targeted by in vivo treatments with an agonistic anti-CD137 monoclonal antibody, it resulted in an increased effect on CD8<sup>+</sup> T-cells expansion and activation with just a modest effect on CD4<sup>+</sup> T-cells, thus confirming that in vivo its signal affects predominantly this subset of T-cells [50][58].

Finally, as evidence about the importance of the CD137 receptor in marking those T-cells that were activated to eliminate a non-self-antigen, it was reported that the CD137 resulted specifically upregulated after an alloantigen stimulation and, upon CD137 depletion, it was possible to remove alloreactive T-cells during hematopoietic transplantation <sup>[59]</sup>.

An important step forward in the field of tumor immunology, was made when CD137<sup>+</sup> T-cells were clearly identified as those cells that were terminally differentiated and antigen-specific effector cells, regardless of the antigen specificity <sup>[60]</sup>. This allowed the isolation of those cells that were considered the real effector cells activated against tumor antigens <sup>[61]</sup>.

## 3. CD137<sup>+</sup> T-Cells: The Natural Tumor-Specific Population

The discovery that CD137 is expressed by most of activated and antigen-specific (both against viral and tumor antigens) CD8<sup>+</sup> T-cells, allowed the isolation of tumor-specific effector T-cells from blood, without knowing the immunogenic epitopes or the MHC-restriction complex. These cells, even if present at low frequencies, were able to kill antigen-expressing cancer cells upon expansion, although this required an ex vivo restimulation with the defined tumor antigen [60][61].

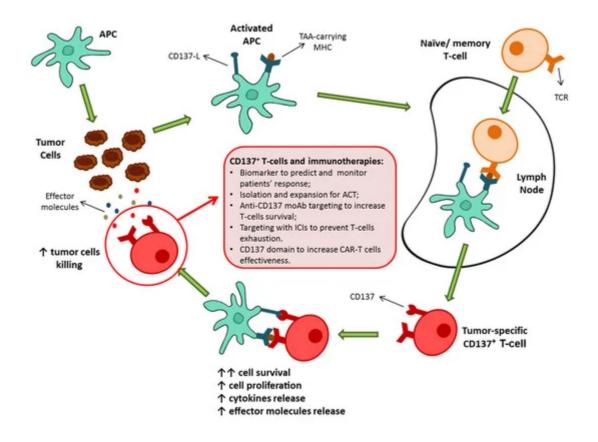
This evidence raised a strong interest in investigating this cell repertoire also inside the tumor. In fact, the tumor microenvironment (TME) is enriched for T-cells specific for defined antigens with cytolytic ability against cancer cells <sup>[62]</sup>. In addition, even if defined antigens are known for different tumor models, exomic sequencing data in different solid tumors proved that cancer cells express a various and heterogeneous set of mutated neo-antigens that are characteristic for every single patient and thus can be recognized by TILs that are able to exert an antitumor response <sup>[63]</sup>. As confirmation, T-cell receptors (TCRs) isolated from CD137<sup>+</sup> TILs, showed a reactivity against various mutations of tumor-derived antigens <sup>[64]</sup>. Given this evidence, the possibility of identifying a tumor-specific T effector population inside the TME without the knowledge of the antigen epitopes seemed very promising.

Initial evidence proved that CD137 is strongly expressed by TILs if compared to spleen- or lymph nodes-derived T-cells and its expression is induced by hypoxia through hypoxia-inducible factor  $1\alpha$  <sup>[65]</sup>.

Recently, Ye et al. decided to investigate the CD137<sup>+</sup> T-cells population in ovarian cancer patients, comparing three different locations in which this subset of cells could be found: TME, ascites, and peripheral blood <sup>[66]</sup>. They demonstrated that CD137<sup>+</sup> T-cells are present in small percentages in the peripheral blood and, to a larger extent, in ascites and even more inside the tumor, showing a progressive hierarchy with the T-cells in a closer proximity to cancer cells expressing the higher percentages of CD137 and then decreasing gradually toward the periphery. Overnight incubation with autologous cancer cells largely increased the percentage of CD137<sup>+</sup> T-cells and their ability of producing a consistent amount of IFN-γ. Additionally, CD137 expression was further increased when T-cell lines with a known antigen specificity were used. Most importantly, when human TILs and tumor cells where transferred into immunodeficient mice, only CD137<sup>+</sup> T-cells (but not CD137<sup>-</sup> T-cells) were able to inhibit tumor

growth <sup>[66]</sup>. Thus, they demonstrated that CD137<sup>+</sup> T-cells are those cells that naturally show the real antitumor reactivity, confirming also that they represent a subset of newly recruited antitumor T-effector cells, being CD137 expression a rapid and transient event upon specific activation. Overall, this study proposed a novel method to isolate and expand tumor reactive TILs that can be used for adoptive T-cell transfer approaches; the vast heterogeneity of TCRs is indeed conserved with this strategy thus helping to prevent the escape of those tumor cells that do not express a determined antigen or those that express mutated antigens.

These findings suggested the potential role of CD137<sup>+</sup> T-cells as key contributors of the antitumor immune responses and thus as potential determiners of the success of immunotherapies as well as novel protagonists of immune-based approaches (<u>Figure 1</u>).



**Figure 1.** Schematic representation of CD137<sup>+</sup> T-cell activation as a result of priming by TAA-carrying activated APC. The consequence of the CD137/CD137-L engagement is a marked increase in cell survival, followed by an increased proliferation, cytokine production, and effector molecules release. Then, possible roles of CD137<sup>+</sup> T-cells population in the present and future of immunotherapy. APC, antigen presenting cell; TAA, tumor-associated antigens; MHC, major histocompatibility complex; ACT, adoptive cell therapy; moAb, monoclonal antibodies; ICIs, immune checkpoint inhibitors; CAR, chimeric antigen receptor.4. CD137<sup>+</sup> T-cells can predict cancer patients' response to immune-based therapies.

Despite these clear results showing the importance of the CD137<sup>+</sup> T-cell population in eliciting an antitumor response, evidence about the role of these T-cells in oncologic patients have only recently emerged (<u>Table 1</u>).

**Table 1.** Summary of the results showing the power of CD137<sup>+</sup> T-cells population as a biomarker able to predict and monitor patients' response to different immune-based therapies in various tumor models.

Cancer Type	Treatment	Results	References
Metastatic NSCLC	Anti-PD-1 ICIs	Higher percentages of CD137 <sup>+</sup> T-cells in PBMC predicted a prolonged patients' OS and PFS.	[ <u>67,68</u> ] Ugolini et al., 2020
Metastatic RCCC	Anti-VEGF-R TKIs and anti-PD-1 ICIS	Percentage of CD137 <sup>+</sup> T-cells in PBMC decreased during patients' progression.	[ <u>69</u> ] Zizzari et al., 2018
Metastatic RCCC	TKIs	Higher percentages of CD137 <sup>+</sup> T-cells in PBMC were associated with responder patients.	[ <u>70</u> ] Zizzari et al., 2020
Metastatic Melanoma	Anti-PD-1 ICIS	CD137 mRNA levels at the tumor site were positively associated with a prolonged OS, PFS, and a better response to the therapy.	[ <u>71</u> ] Fröhlich et al., 2020

In 2020, for the first time we provided evidence about the importance of CD137<sup>+</sup> T-cells in determining the outcome of metastatic non-small cells lung cancer (NSCLC) patients undergoing immunotherapies <sup>[67][68]</sup>. Patients that were positive for the autoantibody IgM-Rheumatoid Factor (IgM-RF) showed indeed a reduced frequency of CD137<sup>+</sup> T-cells in peripheral blood and an increased tendency to develop an early progression, in addition to a markedly reduced progression-free survival (PFS) and overall survival (OS) after the anti-PD-1 treatment <sup>[68]</sup>. In addition, to confirm the importance of this population as an independent prognostic factor, it was reported how a higher percentage of CD137<sup>+</sup> T-cells in peripheral blood mononuclear cells (PBMC) at baseline, was alone associated with a prolonged OS as well as PFS of patients in treatment with an anti-PD-1 ICI <sup>[68]</sup>.

In addition, in 2018 it was proven that, in metastatic renal clear cell carcinoma (mRCCC) patients undergoing the anti-PD-1 treatment, the percentage of CD137<sup>+</sup> T-cells decreased during tumor progression <sup>[69]</sup>. Moreover, patients pretreated with Tyrosin-kinase inhibitor Pazopanib, showed a robust increase in DC activation profile and a subsequent increase of the frequency of CD137<sup>+</sup> T-cells when compared to Sunitinib <sup>[69]</sup>. Still in mRCCC, Zizzari et al. demonstrated that CD137<sup>+</sup> T-cells were positively associated with patients response to TKI <sup>[70]</sup>. In fact, responder patients showed a markedly higher percentage of this T-cell subset when compared to non-responders. These results highlight the importance of this T-cell subset in oncologic patients response to therapies that require, even if in an indirect way, the immune system's ability of killing tumor cells. In this scenario, the percentage of this population in peripheral blood (and most likely also in other districts as draining lymph nodes and TME) could serve as a possible biomarker able to identify those patients that would benefit the most from a determinate treatment that relies on T-cells as final effectors.

Finally, in 2020, indirect evidence of the CD137<sup>+</sup> T-cells power in determining a prolonged survival for cancer patients came from a study on melanoma patients where it was shown that TNFRSF9 low methylation levels and the subsequent increased mRNA expression at the tumor site, that was prevalently identified inside T-cells, correlated with a better OS of patients as well as a better PFS and response to the anti-PD-1 treatment <sup>[71]</sup>.

TNFRSF9 mRNA expression positively correlated also with the frequency of effector and memory tumor infiltrating lymphocytes, while it was inversely correlated with the frequency of naïve tumor infiltrating lymphocytes <sup>[71]</sup>. As a confirmation of its power as biomarker for the identification of activated effector T-cells, TNFRSF9 mRNA expression levels positively correlated with an increased IFN-y signature <sup>[71]</sup>.

These results indicate the potential role of this population as the driver of a successful immunotherapy, thus suggesting the possibility of investigating its presence in patients before undergoing immune-based treatments. In fact, a reduction in its frequency could account for the impossibility of getting a complete or even partial response at least in part of the oncologic patients. In this scenario, strategies aimed at increasing their numbers could be considered at an initial stage, in order to make the patient more prone to efficiently receive an immunotherapeutic treatment.

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