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# Chimeric Antigen Receptor Modified T Cells – Clinical Translation in Stem Cell Transplantation and Beyond

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#### Introduction

The adoptive transfer of re-directed T cells that are engineered by gene transfer to express receptors that target tumor associated molecules and signal T cell effector functions is emerging as an effective modality for cancer therapy, and holds promise for treating a broad range of malignancies, including those that are commonly managed with autologous or allogeneic hematopoietic stem cell transplantation. This review will focus on current issues for engineering T cells with chimeric antigen receptors (CARs) that are specific for molecules on hematologic malignancies, and discuss the prospects and challenges for using such T cells either alone or as an adjunct to hematopoietic stem cell transplantation to reduce the unacceptably high rates of relapse associated with this procedure <sup>1</sup>.

# Chimeric Antigen Receptors - Target Molecules and Receptor Design

Genetic modification of T cells to confer tumor specificity surmounts many of the technical difficulties in isolating and expanding rare tumor-reactive T cells from patients, and can circumvent mechanisms that interfere with the antitumor activity of endogenous T cells by providing control over the specificity, avidity, function, and cellular composition of the antitumor immune response. A notable advance in the field of T cell engineering was the development of chimeric antigen receptors that link recognition domains of antibodies to molecules involved in signaling T cell effector function  $^2$ . CARs usually consist of a single chain variable fragment (scFV) derived from the  $V_H$  and  $V_L$  sequences of a monoclonal antibody specific for a tumor cell surface molecule, fused to sequences that encode a transmembrane domain, the CD3 $\zeta$  signaling domain and one or more costimulatory signaling modules, such as CD28, 4-1BB, OX40, or CD27 (Figure 1)  $^{3,4}$ . Costimulation can also be provided in trans by encoding separately a ligand for a costimulatory receptor in the vector  $^5$ . Although an scFV is the most common tumor recognition domain for CARs, other molecules such as cytokines or peptides that bind to tumor cells have been incorporated as tumor targeting domains in the CAR design  $^6$ .

CARs have been developed that encode scFvs specific for molecules expressed on several malignancies routinely treated with hematopoietic stem cell transplantation. These include the CD19, CD20 and ROR1 molecules expressed on B cell lymphomas and leukemias; CD33 and Lewis Y expressed on acute myeloid leukemia; and NKG2D ligands and Lewis Y expressed on multiple myeloma <sup>7–12</sup>. One of the potential problems with the current targets

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is they do not completely distinguish normal from malignant cells, and on-target toxicity to normal cells expressing the molecule is an expected complication of therapy. This may be tolerated temporarily for lineage restricted molecules such as CD19 and CD20 that are also expressed on normal mature B cells and where intravenous immunoglobulin can be used to replace deficient antibody function. However, on target toxicity to normal tissues remains an issue for which novel strategies to mitigate toxicity such as by regulating CAR expression or T cell survival are needed. Exome sequencing, analysis of gene expression profiles, and elucidation of glycosylation of cell surface molecules on tumors versus normal cells may identify novel surface molecules or isoforms that can be selectively targeted on the malignancy without harming normal cells.

CARs, unlike conventional T cell receptors, are not MHC restricted and have the advantage that a single construct can be used to treat all patients that express the target molecule. However, CARs are synthetic and much remains to be understood concerning how to optimally design the receptor construct to ensure that T cells respond appropriately in vivo to tumor cells. The earliest CARs, termed "first generation", linked the scFV to the CD3\(\zeta\) chain and did not incorporate costimulation into the signaling module <sup>13</sup>. These CARs entered clinical trials but significant antitumor activity was not observed <sup>14,15</sup>. The lack of in vivo activity was attributed to absent costimulation in the CAR, although it must be acknowledged that methods for activating T cells, introducing the receptor constructs, and expanding cells for therapy have improved markedly. Recent studies have used efficient retroviral or lentiviral transduction to introduce CARs that contain one (second generation) or two (third generation) costimulatory domains into T cells and employed culture methods that rapidly expand T cells for adoptive transfer <sup>16</sup>. T cells that express second and third generation CARs, or provide trans costimulation have more potent antitumor activity in vitro and in animal models 5,13,17. It must be conceded that the optimal design for clinical applications is still uncertain and may differ depending on the target molecule and malignancy being treated. These issues may eventually only be resolved by clinical trials that compare distinct constructs, both for efficacy and toxicity.

There are aspects of CAR design in addition to costimulation that may need to be considered including the density of target molecules on tumor cells required for effective T cell signaling, their proximity and accessibility on the tumor cell membrane, and the affinity of the scFv. In CLL for example, the number of cell surface CD19 molecules is several fold greater than that of ROR1, and how CAR occupancy might influence T cell signaling has not been carefully studied. The affinity of the scFv and steric constraints imposed by the location of the target epitope may require alterations in the extracellular spacer length to promote synapse formation between effector T cell and the tumor cell <sup>18</sup>. Structural modeling of CARs interacting with their target molecules and measurement of on and offrates of the CAR and tumor cell ligand may provide further insights into how to optimally design CARs specific for individual targets.

## **Clinical Applications of CAR-Modified T Cells**

CAR engineered T cells have been tested for antitumor activity in pilot clinical trials in which autologous T cells have been removed from the patient, modified to express a CAR, and re-infused into the patient after expansion in vitro. Results of several trials of CARs specific for CD19, CD20, GD2, and other targets have been reported <sup>19–24</sup>. Extremely promising results have been obtained targeting the B-cell lineage restricted CD19 molecule that is expressed on B-cell leukemias and lymphomas with CD19-specific CAR T cells <sup>19–22,25</sup>. Kalos, Porter and June et al. reported durable remissions in patients with B cell chronic lymphocytic leukemia (CLL) after the infusion of autologous T cells transduced to express a CD19-specific CAR that contained a 4-1BB costimulatory domain. In these

studies, the infusion of low doses of T cells led to in vivo expansion and persistence of CAR T cells, subsequent tumor lysis, and a sustained deficiency of normal CD19<sup>+</sup> B cells <sup>19</sup>. Significant antitumor activity, depletion of normal B cells, and side effects related to tumor lysis and cytokine release have also been reported in patients with CLL and lymphoma by groups at the National Institutes of Health, Memorial Sloan Kettering Cancer Center and Baylor College of Medicine in trials in which autologous T cells were modified to express CD19 CARs that contain a CD28 costimulatory domain <sup>20,22,26</sup>. That durable responses can be achieved in patients with advanced B cell malignancies illustrates the potency of CAR-T cells, and suggests that future work to define optimal CD19 CAR constructs, and to integrate T cell therapy earlier after diagnosis or after autologous transplant where marked tumor cytoreduction can be achieved by intensive conditioning may improve outcome and lessen the toxicity resulting from CAR mediated tumor lysis.

The success of targeting CD19 on tumor cells predictably leads to the elimination of normal mature and immature B cells that also express CD19. Moreover, because CD19 is expressed when hematopoietic stem cells commit to the B cell lineage, no new B cells would be expected to emerge and depletion of B cells is likely to persist as long as the transferred T cells survive, express the CAR, and remain functional. Strategies to regulate survival of CAR T cells or turn off CAR expression are being developed as discussed below, and could allow the regeneration of normal B cells from hematopoietic progenitors. CD20 and ROR1 are alternative targets on B cell malignancies, however CD20 is also expressed on normal B cells and would be expected to result in B cell depletion. ROR1, which is uniformly expressed on CLL, mantle cell lymphoma and a subset of ALL, is not expressed on mature B cells in the periphery and could spare this compartment <sup>8</sup>. ROR1 is transiently expressed on early B cell precursors in the bone marrow and on some normal tissues, thus toxicity remains a concern with targeting ROR1. Animal studies are in progress to determine if ROR1 is a safe target for CAR therapy.

CAR-modified T cells may also have applications in solid tumors, including those in children that can be managed with high dose therapy and autologous stem cell transplantation. Candidate surface molecules on solid tumors, including mesothelin, folate receptor, L1CAM, and GD2 have been identified and are being developed for clinical applications. Significant antitumor activity without toxicity has been reported in patients with neuroblastoma treated with T cells modified with a first generation GD2 CAR <sup>27</sup>. The persistence of the transferred cells was relatively short in this study, perhaps related to the lack of costimulation in the CAR. GD2 is expressed on normal peripheral nerves and ontarget toxicity from a sustained T cell response will need to be monitored if more potent CARs are employed <sup>24</sup>.

# **Composition of CAR-T Cell Products**

The initial clinical applications of CAR-T cells have primarily derived cell products for administration by inserting the CAR gene into unselected polyclonal T cells obtained from the patient by leukapheresis. It is now well established that peripheral T cell pool contains different proportions of naïve and memory CD4<sup>+</sup> and CD8<sup>+</sup> cell subsets, and that these subsets diverge in function, transcriptional profiles and epigenetic programming, with potential consequences for their efficacy in adoptive therapy  $^{28}$ . Direct comparisons have revealed profound differences in the ability of T cells from different subsets to persist long term in vivo and revert to the memory pool. For example, in animal models the adoptive transfer of CD8<sup>+</sup> CD62L<sup>+</sup> central memory T cells ( $T_{CM}$ ) or a rare subset of CD62L<sup>+</sup> memory T cells (termed memory stem cells;  $T_{SCM}$ ) that has cell surface markers shared by both naïve ( $T_N$ ) and  $T_{CM}$  cells, exhibit superior survival in vivo and/or mediate superior antitumor activity compared with more differentiated CD62L<sup>-</sup> effector memory ( $T_{EM}$ )

cells <sup>29–31</sup>. CD8<sup>+</sup> T<sub>N</sub> cells are undifferentiated, have long telomeres and substantial proliferative capacity, and as a consequence might be directed by cytokines or small molecules during in vitro expansion and transduction to have properties that confer superior antitumor activity <sup>32</sup>. The CD4<sup>+</sup> T cell compartment is also heterogeneous, with defined naïve, memory and FoxP3<sup>+</sup> regulatory subsets <sup>33</sup>. CD4<sup>+</sup> T cells alone have been shown to exhibit antitumor activity after adoptive transfer in animal models and humans, at least in part by providing help for CD8<sup>+</sup> T cells <sup>34</sup>, and the potential to formulate CAR-modified T cell products that contain defined proportions of CD8<sup>+</sup> and CD4<sup>+</sup> T cells from distinct subsets is beginning to be explored. Because the phenotypic distribution of T cells in the blood is influenced by age, pathogen exposure and prior chemotherapy, the use of unselected populations for gene transduction results in a lack of uniformity in the composition of T cell products administered to individual patients, and could affect potency and toxicity. The disadvantage of having to select distinct subsets for therapy is the additional cost and time associated with deriving the cell product for therapy. Transduction of unselected T cells should provide some proportion of T cells with the desired traits, and for malignancies that are sensitive to immunotherapy, may be sufficient for efficacy. However, differences in the composition of cell products will make it difficult to define dose/response relationships and to ensure immediate potency against less responsive tumors.

## **Incorporating CAR-T Cells Into Stem Cell Transplant Regimens**

The antitumor activity of autologous CD19 CAR-T cells in patients with advanced B cell malignancies, illustrates the potential to incorporate the administration of T cells modified with a CAR to reduce relapse after autologous stem cell transplant, and to utilize donor T cells modified with a CAR to augment the graft versus leukemia (GVL) effect of allogeneic HCT. The importance of lymphodepleting chemoradiotherapy for improving the persistence and antitumor efficacy of transferred T cells was established in studies in melanoma <sup>35</sup>, and suggests that intensive conditioning administered before autologous HCT could provide an ideal environment for administering CAR-T cells. Studies targeting CD19 in lymphoma patients undergoing autologous HCT are in progress and the results will be of interest. If effective, this could renew efforts to use intensive conditioning and autologous HCT combined with T cell immunotherapy for selected solid tumors that express unique cell surface molecules for which CARs have been designed.

Leukemia relapse remains a major cause of failure after allogeneic HCT and the long sought goal of augmenting the graft versus leukemia (GVL) effect without aggravating graft versus host disease remains elusive. This could potentially be accomplished using CAR-T cells, however in this setting the composition of CAR-T cell products will be critical to avoid infusing alloreactive T cells that can cause GVHD. One strategy involves selecting donor CD8+ cytomegalovirus (CMV) or Epstein Barr virus (EBV)-specific T cells that have been administered without GVHD in prior studies <sup>36</sup>, and engineering these cells to express the tumor-reactive CAR <sup>37</sup>. In principle, virus-specific CD4+ T cells could be similarly modified if it essential for therapeutic efficacy to have both subsets in the cell product. Clinical trials of CD19 CAR-modified T cells following allogeneic HCT are now accruing patients, and should assist in defining both the potential for efficacy and limitations of this approach in patients with B cell malignancies. Ideally, CARs specific for molecules on myeloid leukemias will soon be validated and advance to clinical testing to provide therapeutic options for this subset of patients.

# Modification to enhance safety and efficacy of CAR-T cells

Because most of the current CARs do not completely discriminate normal and malignant cells, strategies to improve safety are a significant area of research. A conditional suicide

gene that encodes human caspase 9 fused to a modified human FK-binding protein confers sensitivity to a synthetic small molecule that induces apoptosis through activation of caspase 9. This approach effectively ablated alloreactive T cells in a clinical trial after human hematopoietic cell transplant, and may prove useful to abrogate on-target or off-target toxicities of CAR engineered T cells <sup>38</sup>. Expressing CARs through RNA transfection of T cells ensures a gradual loss of surface CAR expression as T cells divide and may be a useful strategy to define potential toxicity of novel constructs <sup>39</sup>. Another approach is to regulate expression of the tumor-targeting CAR by placing it under the control of regulatory elements that can be turned on (or off) by delivering a small molecule <sup>40</sup>.

Strategies to improve efficacy are also likely to be necessary as clinical trials are extended to more aggressive tumors. Genetic strategies to overcome local tumor immunosuppression mediated by stromal cells, cytokines, and negative signaling pathways in T cells have been devised. One approach expresses a single chain IL-12 molecule in tumor-specific T cells to re-program tumor associated myeloid cells and eliminate the need for lymphodepleting chemotherapy prior to CAR-T cell therapy <sup>41</sup>. Gene editing technology could allow permanent disruption of negative signaling pathways such as PD-1, although combinatorial approaches utilizing blocking antibodies that themselves have some efficacy may be a more convenient approach <sup>42</sup>. The development of strategies to enhance efficacy will best be accomplished by careful analysis of treated patients to define the basis for success and failure.

#### Conclusions

Immunotherapy for cancer has undergone a renaissance in recent years, based both on technologic developments such as the ease in introducing genes into T cells and on an improved understanding of the obstacles to eradicating cancer through immune mechanisms. Although much remains to be discovered, there is considerable optimism that this modality can have a meaningful impact for patients, both alone and as an adjunct to hematopoietic stem cell transplant.

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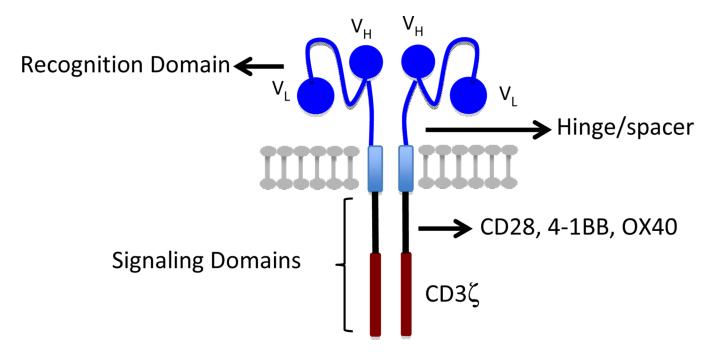
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**Figure 1.** Basic elements of chimeric antigen receptor design