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CAR T Cell Therapy Progress and Challenges for Solid Tumors

Lawrence A. Stern, Vanessa D. Jonsson and Saul J. Priceman

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L. A. Stern · V. D. Jonsson · S. J. Priceman (🖂)

Department of Immuno-Oncology, Beckman Research Institute, City of Hope, Duarte, CA, USA

e-mail: spriceman@coh.org

Department of Hematology and Hematopoietic Cell Transplantation,

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11.1 Introduction to Immunotherapy

The past two decades have marked the beginning of an unprecedented success story for cancer therapy through redirecting antitumor immunity [1]. While the mechanisms that control the initial and ongoing immune responses against tumors remain a strong research focus, the clinical development of technologies that engage the immune system to target and kill cancer cells has become a translational research priority. Early attempts documented in the late 1800s aimed at sparking immunity with cancer vaccines were difficult to interpret but demonstrated an opportunity that more than 100 years later has blossomed into the current field of cancer immunotherapy. Perhaps the most recent and greatest illustration of this is the widespread appreciation that tumors actively shut down antitumor immunity, which has led to the emergence of checkpoint pathway inhibitors that re-invigorate the body's own immune system to target cancer [2, 3]. This class of drugs, with first FDA approvals in 2011, has demonstrated impressive durable clinical responses in several cancer types, including melanoma, lung cancer, Hodgkin's lymphoma, and renal cell carcinoma, with the ongoing investigation in others. The biology and ultimate therapeutic successes of these drugs led to the 2018 Nobel Prize in Physiology or Medicine, awarded to Dr. James Allison and Dr. Tasuku Honjo for their contributions to cancer therapy [4]. In parallel to the emerging science that aided in unleashing the body's own antitumor immunity with checkpoint pathway inhibitors, researchers were also identifying ways to re-engineer antitumor immunity through adoptive cellular immunotherapy approaches. Chimeric antigen receptor (CAR)-based T cell therapy has achieved an early head start in the field, with two recent FDA approvals in 2017 for the treatment of B-cell malignancies [5]. There is an explosion of preclinical and clinical efforts to expand the therapeutic indications for CAR T cell therapies, with a specific focus on improving their clinical utility, particularly for the treatment of solid tumors. In this chapter, we will highlight the recent progress, challenges, and future perspectives surrounding the development of CAR T cell therapies for solid tumors.

11.2 CAR T Cell Therapy

The development of effective CAR T cell therapies for any cancer type lies in several key variables [6]—(1) design of CAR constructs, (2) manufacturing processes that lead to the final therapeutic product, and (3) clinical study design to comprehensively assess safety and efficacy of CAR T cell therapies and combinatorial immunotherapy strategies. We will summarize key findings in these areas that have contributed to the successes in CAR T cells in treating hematological malignancies to date, as well as address many of the challenges facing CAR T cell therapy for treating solid tumors.

11.2.1 CAR Design

CARs are modular synthetic receptors that redirect antigen specificity of T cells to cell surface targets expressed by tumor cells, thereby eliciting a potent T cell functional output primarily through cytolytic activity and production of inflammatory cytokines. CARs consist of four major components: the antigen-binding domain, the extracellular spacer domain, the transmembrane domain, and the intracellular signaling region consisting of co-stimulatory and CD3 ζ cytolytic domains (Fig. 11.1). Observations from engineering these components were recently reviewed extensively [7]. Here, we will summarize major findings that contribute to the current convention for designing new CARs.

The antigen-binding domain confers target specificity to the CAR. These domains are often derived from the variable regions of monoclonal antibodies termed single-chain variable fragments (scFv), although other targeting moieties have been described, including but not limited to: natural or engineered receptor ligands [8, 9], receptor extracellular domains [10, 11], and engineered non-immunoglobulin binding proteins [12, 13]. The majority of solid tumor targets evaluated to date are also expressed on normal tissue at various levels, which raises toxicity concerns with "on-target off-tumor" targeting of normal tissue [14]. Therefore, the optimization of CAR selectivity and potency has been heavily studied by modulating properties of the antigen-binding domain. For instance, fine-tuning scFv affinity has been impactful in setting antigen expression thresholds



Fig. 11.1 Illustrations of a T cell receptor (TCR) and a chimeric antigen receptor (CAR). **a** TCR complex on the surface of a T cell, composed of six subunits including TCR alpha (α) and beta (β), a homodimer of CD3 zeta (ζ), and dimers of CD3 epsilon (ϵ) with either CD3 gamma (γ) or CD3 delta (δ). **b** CAR construct on the surface of a T cell, composed of an antigen-binding domain (e.g., a single-chain variable fragment, or scFv), an extracellular spacer domain (e.g., an IgG4 Fc molecule), a transmembrane domain, an intracellular co-stimulatory domain, and an intracellular CD3 ζ cytolytic domain

required for CAR activation [15-18]. One potential avenue for decreasing toxicity concerns is reducing the affinity of the antigen-binding domain. This may increase the requirement for higher antigen density on tumor cells for optimal activation of CAR T cells, and therefore, bypass targeting of antigen-low healthy tissue. This rationale was explored by Liu and colleagues, who generated HER2-specific CARs targeting solid tumors with a 4-log range of binding affinities [15]. This study observed that the threshold for antigen density that results in CAR activation correlates with antigen-binding domain affinity. Antigen-binding domains with low nanomolar and sub-nanomolar affinity mediated T cell activation against all HER2⁺ cell lines tested, whereas antigen-binding domains with micromolar affinity were much more selective for tumors with higher levels of HER2 expression. This observation was confirmed using EGFR-specific CAR T cells, as both HER2 and EGFR are expressed at lower levels on several critical normal tissues. Other studies have made similar observations correlating decreased CAR binding affinity and improved selectivity for high target antigen expression [16, 17]. However, in each case, decreased affinity also correlated with lower CAR T cell-mediated cytokine secretion, even with high target antigen expression. Thus, the interplay of binding affinity, selectivity for disease-specific target antigen density, and functional activation of CAR T cells must be carefully considered when designing new CARs.

The extracellular spacer domain provides an extension from the T cell membrane and flexibility to allow the antigen-binding domain to optimally access the targeted epitope. The selected spacer can impact CAR expression, flexibility, epitope accessibility, and strength of activation outputs [19, 20], which ultimately affects CAR functionality. Most often, extracellular spacer domains are derived from natural molecules. Common examples include CD8a hinge, CD28 hinge, and IgG hinge and Fc regions. The proper spacer length for a particular binding domainantigen pair is often empirically determined and likely depends on the target epitope location and relative level of steric hindrances present on the target cell. Notable examples of CARs requiring short spacers (CD19, CEA) [21] and long spacers (MUC1, membrane-proximal epitopes of ROR1) [19, 22] for optimal activity exist in literature. In some contexts, extracellular spacer domains can also mediate undesired effects, including antigen-independent tonic signaling [23] and interaction of IgG-derived spacers with FcyR-expressing cells [23, 24]. Importantly, these effects can be abrogated by either selecting different spacer domains or by further engineering of the spacer based on structural or functional considerations.

The transmembrane domain serves to anchor the CAR to the T cell membrane. Like the extracellular spacer domains, transmembrane domains are derived from natural proteins, with the most common versions including CD4, CD8, and CD28. The impact of the transmembrane domain on CAR activity is not well-studied as this domain is often changed as required by either the extracellular spacer domain or the intracellular signaling domains. Experiences with CARs to date show that the transmembrane domain can be active in signaling [25] and dimerization with endogenous signaling molecules [26], and can also influence CAR expression level [11].

Perhaps the greatest attention in optimization strategies has surrounded the intracellular co-stimulatory signaling domain. The first version of engineered CARs in the late 1990s (termed immunoglobulin-T-cell chimeric receptor molecules) [27] were so-called "first-generation" CARs, which included an antigen-binding domain, an extracellular spacer domain and transmembrane domain, and an intracellular CD3 ζ or FcR γ signaling domain. In vitro, these CAR T cells showed potent antitumor activity, yet demonstrated limited persistence and durability of therapy [28]. As these first-generation CAR T cells moved to clinical investigation, a lack of efficacy became clear in a variety of diseases. In hematologic malignancy, a phase 1 trial targeting CD20 in indolent non-Hodgkin lymphoma and mantle cell lymphoma reported safety and feasibility with modest efficacy [29]. In the context of solid tumors, a GD2-targeted first-generation CAR to treat neuroblastoma reports one patient achieving a complete response and one patient having disease cleared from the bone marrow [30]. Clinical studies targeting FR α in ovarian cancer [31], TAG72 in colorectal cancer [32], and CAIX in renal cell carcinoma [33] showed no objective clinical responses, and many of these studies remarked a lack of T cell persistence.

To address the lack of durable CAR T cell therapy, early in vivo models of malignancies illuminated the importance of co-stimulation with B-cell CD19-targeted CAR T cells [34]. In this study, durable antitumor response was observed when treating Raji Burkitt lymphoma (expressing co-stimulatory molecules CD80 and CD86) but not when treating NALM-6 pre-B-cell ALL (lacking co-stimulatory molecule expression). In vivo efficacy of CD19-CAR T cells in the second model was rescued with an engineered expression of CD80 in NALM-6 cells. Importantly, in the Raji model, CD19-CAR T cells were detected in the bone marrow of treated mice 21 days post-infusion, further accentuating the importance of co-stimulation in CAR T cell persistence. Similar phenomena were contemporaneously observed using solid tumor-directed CAR T cells targeting PSMA [28]. With this new understanding, "second-generation" CARs, which contain one co-stimulatory domain in series with the CD3 ζ intracellular domain were developed [35, 36]. These CARs were able to mediate CAR T cell expansion after repeated antigen exposure while maintaining antigen-specific cytotoxic activity. The most common co-stimulatory domains added to second-generation CARs were derived from CD28 [35] and 4-1BB [36], but other domains including ICOS [37], OX40 [37], and CD27 [38] have also been explored. Clinical translation of these second-generation CAR T cells has thus far resulted in strong therapeutic responses in hematologic malignancies including chronic lymphocytic leukemia [39], B-cell acute lymphoblastic leukemia [40], diffuse large B-cell lymphoma [41], and multiple myeloma [42]. Second-generation CAR T cells have now entered clinical investigation for solid tumors, including glioblastoma [43–45], advanced sarcoma [46], liver metastases [47], as well as mesothelioma, ovarian cancer, and pancreatic cancer [48]. A summary of clinical trials evaluating CAR T cells for solid tumors has been recently detailed elsewhere [49].

Despite the success of second-generation CAR T cells, the hypothesis remained that co-stimulation via only one domain would lead to incomplete T cell activation.

Thus, "third-generation" CARs, which incorporated two co-stimulatory domains in series with the CD3 ζ , have been evaluated [50]. The most common combinations of co-stimulatory domains are CD28-OX40 and CD28-4-1BB [51]. Preclinical studies with third-generation CARs show mixed results. CARs incorporating CD28 and 4-1BB signaling demonstrated stronger cytokine production and improved in vivo antitumor response in lymphoma [52] and pulmonary metastasis [52] models relative to second-generation CARs. However, they failed to outperform a secondgeneration counterpart in a pancreatic cancer model [53], and resulted in decreased in vitro cytokine production and no in vivo treatment benefit relative to second-generation CARs in a leukemia model [54]. Incorporation of CD28 and OX40 signaling domains resulted in improved therapy of colon adenocarcinoma in vivo [55] and shows improved activation and cytokine production in vitro [50]. Clinical application of third-generation CAR T cells to date has been limited, but has not shown marked improvement over second-generation CAR T cells [29, 56]. However, additional investigations are warranted to define the optimal intracellular co-stimulatory domain required for CARs based on disease indication and tumor antigen target.

11.2.2 CAR T Cell Manufacturing

The processes used for manufacturing CAR T cells can have a profound impact on the efficacy of a clinical product. Across many clinical trials, there are significant variations to the T cell subsets chosen for CAR engineering and cell expansion protocols used. Unfortunately, due to the relatively small number of CAR T cell clinical trials completed to date, there have been few direct comparisons of manufacturing methods for a single CAR product. A recent review details many of these parameters [57].

11.2.3 T Cell Subsets for CAR Engineering

The majority of CAR T cell clinical trials do not select particular T cell subsets, choosing rather to isolate and engineer peripheral blood mononuclear cells (PBMCs), using stimulation methods and cytokine regimens to selectively expand T cells [57]. Early clinical trials with first-generation CARs in neurological malignancies isolated and expanded CD8⁺ T cell clones for manufacturing [58, 59], but this procedure led to products with low persistence in patients, likely due to exhaustion from clonal ex vivo T cell expansion. More recent innovation involves the engineering of stem-like T cell subsets [60]. Preclinical data showing the ability of central memory T cells to persist after adoptive transfer due to their stemness [61, 62] led to the use of this subset in a phase 1 clinical trial in non-Hodgkin lymphoma demonstrating safety [63] and in a unique case study of complete response in a glioblastoma patient [43]. Further, preclinical investigation of central memory T cells showed that using defined 1:1 mixtures CD4⁺ and CD8⁺ yielded more

consistent potency relative to unenriched central memory T cell products [64], leading to a phase 1 clinical trial in adult B-cell acute lymphoblastic leukemia with 93% remission rate [65]. Additional phase 1 clinical trials have been applied to define CD4⁺/CD8⁺ mixtures with similar success in other B-cell diseases [66, 67]. In phase 1 clinical trial for B-cell non-Hodgkin lymphoma, CD19-CAR T cells derived from central memory T cells and naïve/memory T cells were directly compared [68]. Both arms of treatment showed efficacy in patients, but naïve/memory T cells were viewed as the superior platform because of their greater vield from apheresis as it required fewer enrichment steps, shorter ex vivo expansion time, and superior in vivo expansion. Application of naïve/memory CAR T cells in a recent phase 1 clinical trial for adult relapsed/refractory B-cell acute lymphoblastic leukemia yielded a 100% complete response rate in 13 patients treated [69]. In a retrospective study of PBMC-derived CTL019 CAR T cell products manufactured for the treatment of chronic lymphocytic leukemia, memory phenotype was correlated with complete-response in patients [70]. The impressive in vivo persistence and efficacy of CAR T cells with memory phenotype in these hematological trials motivate the application of these subsets in solid tumor indications. Further, CD8⁺ tumor-infiltrating lymphocytes from patient breast and melanoma tumors dominantly display memory phenotype and retain polyfunctionality despite the expression of checkpoint molecules [71]. Preclinical studies have also underscored the importance of memory phenotype in both syngeneic [72] and humanized [73] solid tumor models.

The majority of CAR T cell products are engineered from a patient's own PBMCs or autologous products. This can lead to several issues in manufacturing, including high cost, manufacturing failures due to dysfunctional cells in the presence of disease and subsequent pre-treatment, disease progression during manufacturing, and contamination of circulating tumor cells in the apheresis product [74]. Because of these challenges, avenues for developing "off-the-shelf" or allogeneic cell-based immunotherapies, which can be obtained from healthy donors and banked, are actively being explored. Recent advances in genome modification enable engineering of healthy donor T cells or inducible pluripotent stem cells to remove endogenous HLA and TCR [75, 76]. Clinical trials are currently underway using off-the-shelf CD19-CAR T cells (NCT03939026), CD123-CAR T cells (NCT03190278), and BCMA-CAR T cells (NCT03752541). Likely, the solid tumor CAR T cell field may follow suit with evaluating allogeneic CAR T cell therapies, as another potential benefit of this approach is the removal of heterogeneity and potential immunosuppressive immune cell populations in the blood of advanced cancer patients.

11.2.4 Ex Vivo T Cell Expansion

Several methods for ex vivo T cell activation and expansion have been explored. Generally, isolated T cells are stimulated through the T-cell receptor, and co-stimulation through agonistic antibodies, cytokines, and/or feeder cells sustains the expansion [57]. Early protocols regularly used a monoclonal anti-CD3 antibody (OKT3) for TCR stimulation and IL-2 for T cell expansion [77]. This method was later shown to promote a more effector memory phenotype in expanded T cells, whereas the application of anti-CD28/anti-CD3 antibody-coated magnetic beads for stimulation promoted a more central memory phenotype [78]. Other studies have shown that the application of high concentrations of IL-2 in T cell culture leads to a more exhausted T cell product with poor effector function [73]. Investigations of the appropriate cytokine cocktails to sustain ex vivo expansion while maintaining memory phenotype revealed that culture with IL-7 and IL-15 cytokines increased the frequency of stem cell memory CD8⁺ T cells, which displayed greater antitumor activity via increased resistance to activation-induced cell death when compared to IL-2 expanded T cells [79]. A recent study showed that T cell expansion with IL-15 alone produced similar retention of stem cell memory phenotype, decreased mTORC1 activity, reduced expression of glycolytic enzymes, and improved mitochondrial fitness relative to T cells cultured with IL-2 [80].

11.2.5 Preconditioning and Chemotherapy Combinations to Enhance CAR T Cell Therapy

Through clinical experience with adoptive cell therapy, non-myeloablative lymphodepleting preconditioning is known to enhance outcomes. Preclinical studies have shown that the removal of host immune cells prior to adoptive cell transfer increases the in vivo availability of yc cytokines important to T cell functionality [81]. One lymphodepleting agent, cyclophosphamide, is known to enhance immune function further due to the depletion of regulatory T cells, which are hypersensitive to its effects [82]. Preclinical study has also shown that cyclophosphamide treatment can deplete myeloid-derived suppressor cells and, in combination with IL-12, increase the presence of inflammatory monocytes and neutrophils in colon cancer models [83]. A clinical comparison of preconditioning with cyclophosphamide with or without fludarabine revealed the combination approach yielded superior treatment of non-Hodgkin lymphoma, likely due to increased persistence of the engineered T cells due in part to a decreased immune response against the transgene [84]. Improved CAR T cell engraftment after preconditioning with cyclophosphamide and fludarabine was also observed in a clinical trial using first-generation CEACAM5-specific CAR T cells [85]. In addition to improving CAR T cell persistence, chemotherapies can have other synergistic effects with CAR T cell therapy. Lenalidomide, an immunomodulatory drug that has anti-multiple myeloma effects and co-stimulatory effects on T cells, enhanced CS1-targeted CAR T cell treatment in preclinical models of multiple myeloma via enhancement of the immune synapse [86]. Decitabine, a DNA methyltransferase inhibitor, enhanced CD19 expression, and thus susceptibility to CD19-targeted CAR T cell therapy, in both in vitro lymphoma models and in two treated patients [87]. The combination of temozolomide with EGFRvIII-targeted CAR T cells improved treatment of glioblastoma xenografts in mice and has been explored with an escalated dose in preclinical models as the sole lymphodepleting agent prior to CAR T cell therapy [88]. Interestingly, this study showed that the application of dose-intensified temozolomide significantly increased CAR T cell infiltration into tumors without significantly decreasing the presence of regulatory T cells. In sum, the utility of preconditioning and chemotherapy has been validated in combination with CAR T cell therapies for hematological malignancies and has become an attractive area of clinical research for the development of solid tumor CAR T cell therapies.

11.2.6 CAR T Cell Route of Administration

Although targeting hematological malignancies has nearly strictly required intravenous administration of CAR T cells, solid tumors introduce a unique opportunity to localize CAR T cell delivery to target tumors in selected disease sites. Two major reasons to take advantage of different routes of CAR T cell administration compared with systemic delivery are (1) to avoid the requirement of trafficking of CAR T cells to sites of disease, and (2) to direct the on-target activity of CAR T cells in tumors, thereby minimizing their opportunity to target normal tissues. Trafficking of CAR T cells in solid tumors may be hampered by their inability to penetrate tumor stroma and other physical barriers, as well as by the harsh immunosuppressive microenvironment that may impede their mobility into the tumor [89] (more details on the immunosuppressive tumor microenvironment may be found later in the chapter). Additionally, since many solid tumor antigens targeted by CAR T cells are expressed at varying levels in select normal tissue, local or regional CAR T cell delivery may mitigate the potential for on-target off-tumor toxicities [90] (more details on the selection of solid tumor antigens may be found later in the chapter).

Several examples of local or regional delivery of CAR T cells have been evaluated preclinically and in phase 1 trials. Intraperitoneal injection significantly outperformed the systemic injection of CAR T cells in preclinical models of ovarian cancer [91, 92] and peritoneal carcinomatosis [93]. On the strength of preclinical success, a phase 1 clinical trial is currently ongoing comparing intravenous and intraperitoneal infusion of MUC16-targeted CAR T cells in ovarian cancer (NCT02498912). Intravenous administration of mesothelin-targeted CAR T cells that use a murine-derived scFv for antigen recognition has yielded antibody responses against the murine component and anaphylaxis [94, 95]. To improve the efficacy of this therapy and potentially shield the CAR T cells from endogenous immunity, intrapleural delivery of CAR T cells was explored. This route of administration showed superior treatment of a preclinical orthotopic model of malignant pleural mesothelioma in both lung and extrathoracic sites compared to intravenous administration [96], leading to an ongoing phase 1 clinical trial (NCT02414269). Intrahepatic arterial delivery of CAR T cells for liver metastases has been explored preclinically, revealing the challenges of liver myeloid-derived suppressor cells to immunotherapy [97] and in phase 1 clinical trial, demonstrating safety in four patients [47]. Intraventricular administration of CAR T cells targeting HER2 in breast cancer brain metastases [98] and IL13R α 2 in glioblastoma [99] showed superior therapy relative to intravenous injection in orthotopic xenograft models. Importantly, this route of administration offers advantages over intravenous and intracranial delivery in the treatment of multifocal disease. In one patient, intraventricular infusion of IL13R α 2-targeted CAR T cells resulted in a complete response of glioblastoma [43]. Phase 1 clinical trials for intraventricular injection of CAR T cells in glioblastoma (NCT02208362, NCT03389230) and recurrent brain or leptomeningeal metastases (NCT03696030) are ongoing.

11.3 Barriers to Solid Tumor CAR T Cell Therapies

This chapter has highlighted multiple aspects critical to developing effective CAR T cell strategies for the treatment of solid tumors. The three most challenging areas that require attention in the development of next-generation CARs for solid tumors are (1) selective targeting of tumor antigens, (2) tumor antigen heterogeneity, and (3) the immunosuppressive tumor microenvironment. These challenges are active areas of translational research, and will likely require empirical testing for each tumor type, molecular signature, and disease stage of therapeutic intervention.

11.3.1 Solid Tumor Target Antigen Selection

There are nearly 300 CAR T cell clinical trials currently listed on NIH's U.S. National Library of Medicine (ClinicalTrials.gov), with over 50 trials in solid tumors. The solid tumor antigens most frequently targeted by CAR T cell therapy include CEA, EGFR, EGFRvIII, GD2, HER2, IL13R α 2, PSCA, and PSMA [14] and more are summarized in Table 11.1. While all of these antigens are either over-expressed and/or amplified in tumors compared with normal tissue, their protein expression is not uniquely restricted to tumor cells, with the exception of EGFRvIII, a common oncogenic rearrangement in glioblastoma marked by deletion of exons 2–7 of EGFR. Therefore, unlike CD19, a B-cell restricted antigen that is expressed in many B-cell malignancies, solid tumor antigen targets pose significant toxicity concerns that may limit their utility in CAR T cell therapy.

Examples of such toxicities have been observed in clinical trials. A phase 1 trial at the NIH treated three patients using a murine TCR-expressing autologous T cell therapy targeting CEA, and although bioactivity was observed in all three patients with an objective regression in one patient, all patients developed severe transient inflammatory colitis [121]. Similar on-target toxicities were observed in a clinical trial evaluating CAIX-specific CAR T cells in patients with renal cell carcinoma, demonstrating targeting of normal bile duct epithelial cells known to express low levels of CAIX [33]. Perhaps most famously, a serious adverse event was observed in a metastatic colon cancer patient treated with a third-generation HER2-CAR T cell product at the NCI, which resulted in acute respiratory distress syndrome and

Target	Aliases	Cancers targeted with CAR T cell therapy
B7-H3	CD276	Pancreatic ductal adenocarcinoma, ovarian cancer, neuroblastoma [100]; osteosarcoma, Ewing sarcoma, medulloblastoma [101]; glioblastoma [102]
CAIX	Carbonic anhydrase IX	Renal cell carcinoma [33]
CD44v6	CD44 variant 6	Sarcoma [103]; colon cancer [104]
CEA	Carcinoembryonic antigen	Liver metastases [47]
EGFR	Epidermal growth factor receptor; HER1; ERBB1	Non-small cell lung cancer [105]
EGFRvIII	Epidermal growth factor receptor variant III	Glioblastoma [44]
ЕрСАМ	Epithelial cell adhesion molecule	Prostate cancer [106]
FRα	Folate receptor alpha	Ovarian cancer [107]; colon cancer, pancreatic cancer [108]
GD2	Disialoganglioside 2	Neuroblastoma [109]; diffuse midline glioma [110]; melanoma [111]
GPC3	Glypican-3	Hepatocellular carcinoma [112]; lung squamous cell carcinoma [113]
HER2	Human epidermal growth factor receptor 2; ERBB2	Biliary tract cancer and pancreatic cancer [114]; sarcoma [46]; colon cancer [56]; medulloblastoma [115]; breast cancer [116]; brain metastases [98]
IL13Ra2	Interleukin 13 receptor alpha 2	Glioblastoma [43]
MSLN	Mesothelin	Malignant pleural mesothelioma, ovarian carcinoma, pancreatic ductal adenocarcinoma [48]
MUC1*	MUC1 cleavage product	Breast cancer [117]
MUC16	Mucin 16	Ovarian cancer [91]
PSCA	Prostate stem cell antigen	Prostate cancer [118]
PSMA	Prostate-specific membrane antigen	Prostate cancer [119]
TAG72	Tumor-associated glycoprotein 72	Ovarian cancer [92]; colorectal cancer [32]
Tn-MUC1	Tn-glycoform of MUC1	Pancreatic cancer [120]

Table 11.1 Solid tumor targets for CAR T cell therapy

death of the patient five days after treatment [56]. Two recent studies, however, have reported safety and bioactivity in two clinical trials evaluating second-generation HER2-CAR T cells in patients with advanced sarcoma and glioblastoma [45, 46]. One potential avenue for overcoming on-target off-tumor toxicity is the implementation of a suicide gene strategy, which would allow selective depletion of engineered cells via treatment with a secondary inducing agent at the onset of adverse events [122]. Apoptosis can be mediated by the

expression of engineered endogenous apoptotic molecules that can be dimerized via small molecule drugs [123]. Examples of this strategy include inducible FAS [124] or inducible Caspase 9 [125, 126]. Co-expression of transmembrane-anchored proteins or peptides can mark engineered cells for destruction through monoclonal antibody therapy. Expression of full-length CD20 [127] or CD20 mimotope independently [128] or as part of the CAR construct [129, 130] enables the depletion of CAR T cells by Rituximab treatment. A truncated, non-signaling version of EGFR has been shown to facilitate CAR T cell depletion with Cetuximab therapy [131]. Importantly, while suicide gene strategies are attractive for ensuring safety, their implementation abruptly terminates therapy of potentially rapidly progressing disease. This motivates the development of other strategies to ensure safety in treatment, leaving suicide gene activation as a last resort for high-grade adverse events. One such approach was recently reported using Dasatinib, an FDA-approved tyrosine kinase inhibitor for the treatment of t(9;22) chronic myelogenous leukemia and Philadelphia chromosome + acute lymphoblastic leukemia, which suppresses T-cell activation via inhibition of proximal TCR signaling kinases, such as Src, Fyn, and Lck [132, 133]. This pharmacological approach to transiently inhibiting CAR T cell function may allow for the rescue of CAR T cell therapy once toxicities subside.

To overcome targeting tumor antigens that are also found in normal tissues, such as CEA and HER2, targeting tumor-restricted post-translational modifications may provide a unique opportunity for the development of CAR T cell therapy for solid tumors. One of the well-characterized post-translational processes that are differentially regulated in tumor cells is protein glycosylation. The most prevalent of these aberrantly glycosylated antigens are truncated O-glycans, including Tn (GalNAca1-O-Ser/Thr) and sialyl-Tn (STn) (NeuAca2-6-GalNAca1-O-Ser/Thr), which are found over-expressed in many solid tumor types [134]. The four major examples that have been evaluated as CAR T cell targets are MUC1, MUC16, B7-H3, and TAG72. A report of a first-generation CAR T cell therapy for patients with colorectal cancer targeting the tumor-associated glycoprotein TAG72 [32] demonstrated safety and bioactivity, but no tumor responses were observed. Two potential explanations for the lack of therapy in this trial was the use of first-generation CARs and the observed anti-CAR immune responses. Newer versions of TAG72-CAR T cells are being investigated, including second-generation CAR T cells [92] and modifications to the scFv to avoid anti-idiotype immunogenicity [135], and will inform the field on the utility of targeting TAG72⁺ solid tumors with CAR T cells.

Several ongoing phase 1 clinical trials are evaluating and targeting MUC1 with CAR T cells, which is highly over-expressed and aberrantly glycosylated in many solid tumor types [22]. Given the expression of full-length MUC1 in normal tissue, however, novel engineering strategies are warranted to avoid on-target toxicities that have been observed in prior studies mentioned above. Two novel tumor-specific versions of MUC1-targeted CAR T cells are now being evaluated in early clinical trials. The first is a CAR targeting the tumor-associated Tn-glycoform of MUC1 (Tn-MUC1) [120, 136], which was shown to be highly expressed in

tumor tissue, but absent in normal tissue, as compared with full-length MUC1. A similar approach was recently evaluated in mice with CAR T cells targeting the novel cleavage product, MUC1*, shown to be expressed on the cell surface of tumor cells but not in normal tissue [137]. A phase 1 trial has just begun testing MUC1*-CAR T cells for patients with breast cancer (NCT04020575). Additionally, MUC16 has been explored as a target for multiple solid tumor types, and an ongoing phase 1 trial is exploring MUC16ecto-CAR T cells for the treatment of solid tumors [138, 139] (NCT02498912). More recently, the glycoprotein B7-H3 was found to be over-expressed and aberrantly glycosylated in multiple solid tumor types. Preclinical studies have demonstrated the safety and efficacy of CAR T cells targeting B7-H3 [100–102], and a phase 1 clinical trial has just begun to evaluate the safety and efficacy of B7-H3-targeted CAR T cells in patients with recurrent glioblastoma (NCT04077866).

11.3.2 Improving Tumor Antigen Selectivity of CARs

Novel strategies have emerged in CAR design to further control the specificity and activity of CAR T cells for improved safety and antitumor efficacy. Perhaps the earliest example of this for solid tumors was investigated by Kloss and colleagues, using a combinatorial CAR targeting PSCA and PSMA in prostate cancer models. In this system, the co-stimulatory domain and the CD3ζ cytolytic domain were uncoupled and required two antigens to be simultaneously expressed on tumors for optimal CAR T cell activation [140]. Uniquely, the greatest antitumor activity in preclinical models was achieved using a first-generation PSCA-CAR, which was further affinity-tuned for optimal tumor targeting, along with a PSMA-CAR containing a 4-1BB co-stimulatory domain that lacked cytolytic activity (no CD3 ζ domain). More recent versions of controlled CARs include drug-inducible platforms. One of the most promising examples of this uses an inducible MyD88/CD40 (iMC), which can be triggered in vivo with the synthetic dimerizing ligand, rimiducid, for potent co-stimulation of CAR T cells [141]. This strategy has been employed effectively in preclinical studies targeting HER2, demonstrating superiority compared with second-generation HER2-CAR T cells with CD28 co-stimulation [142]. The major improvement in this strategy may involve the ability to modulate signaling of the CAR, controlling both safety and efficacy. A phase 1 trial with this approach has been initiated in targeting PSCA⁺ pancreatic cancers, with interim results demonstrating safety and bioactivity in patients [143].

11.3.3 Tumor Antigen Heterogeneity and Escape

One of the major limitations to current CAR T cell therapies is single antigen targeting. Tumor resistance to single therapeutic agents is well-established as the majority of tumors are heterogenous, and prolonged targeting of a single drug-sensitive pathway can ultimately lead to drug-resistant tumor recurrences.

Acquired or intrinsic resistance patterns following CAR T cell therapy have also been observed. CD19-CAR T cell therapy has demonstrated durable clinical remissions in 70-90% of patients with B-cell malignancies including acute lymphoblastic leukemia (ALL), however, emerging follow-up data from clinical trials show a common mechanism of resistance including loss and/or downregulation of CD19 antigen in up to 70% of patients who recur following treatment [144, 145]. Early clinical findings using CAR T cells for solid tumors have observed similar antigen escape resistance mechanisms. For instance, a phase 1 trial evaluating intravenous delivery of EGFRvIII-specific CAR T cells in patients with recurrent glioblastoma, known for its antigen heterogeneity, showed antigen loss resulting in tumor resistance [44]. A case report targeting IL13R α 2 in glioblastoma with CAR T cells demonstrated decreased IL13R α 2 expression in tumor recurrences [43], suggesting that antigen escape also may have contributed to tumor relapse. Multiple mechanisms may exist that underlay antigen escape following CAR T cell therapy. Hamieh and colleagues recently demonstrated decreased tumor target density by extracting surface expressed antigen from tumor cells by CAR T cells through a process known as trogocytosis [146]. These studies strongly suggest that treatment optimization through CAR design or the rational design of combination and/or sequential CAR T cell strategies targeting distinct tumor antigens will be necessary for effective disease control.

11.3.4 Multi-targeted CAR T Cells

To reduce the relapse rate in CAR T cells for the treatment of hematological malignancies, studies have emerged using dual-targeted CAR T cells. Such approaches have utilized either dual CAR constructs, or two scFvs ("OR"-gate) within a single CAR construct (known as tandem CARs) to simultaneously target different tumor antigens. Both strategies have been employed targeting CD19 and CD22 in relapsed/refractory ALL, with promising early clinical data suggesting that dual-targeting may prolong durable remission rates [147]. Additional studies are ongoing with simultaneous targeting of CD19 and CD20 with "OR"-gate CARs [148, 149], as well as CD19 and CD123 co-targeting [150] and others [151]. These approaches are also being evaluated for CAR T cells targeting multiple myeloma [152].

In solid tumors, HER2 and MUC1 tandem CARs have been evaluated in preclinical models of breast cancer with improved activity over single antigen targeting CARs [153]. Similarly, dual-targeting of HER2 and IL13R α 2 in glioblastoma has been studied in xenograft models [154]. In this study, tandem CARs were evaluated in both human xenograft and syngeneic immunocompetent mouse models of glioblastoma, and compared to T cells expressing both CARs, or pooling single-specific CAR T cells. Interestingly, tandem targeting of HER2 and IL13R α 2 resulted in superior antitumor activity, and reduced antigen escape compared with the two other dual-targeting approaches. While this finding may be specific for different antigens being targeted in solid tumors, it highlights the need to empirically define dual-targeting approaches that improve antitumor responses and potentially mitigate antigen escape mechanisms of tumor resistance.

Additional innovative approaches have been developed to target multiple antigens in attempts to overcome antigen escape in solid tumors. Recently in glioblastoma, a novel EGFRvIII-specific CAR was designed to secrete a bispecific T cell-engager (BiTE) targeting EGFR. In this study, the co-targeting of EGFRvIII and EGFR using this strategy successfully controlled heterogeneous model tumors compared with either strategy alone [155]. Another novel approach to target two tumor antigens was recently investigated in preclinical studies using oncolytic viruses to infiltrate tumors and secrete EGFR-BiTEs, in combination with CAR T cells targeting FR α , which improved antitumor activity over monotherapy [108].

Multi-targeting introduces additional toxicity concerns as each new target potentially compounds healthy tissue targeting. Newer synthetic CAR switches are being developed to circumvent this likelihood of exacerbating toxicities to normal tissue. Perhaps the most intriguing approach in recent years has been demonstrated using modular synthetic Notch receptors (synNotch) for "AND"-gate CAR T cell regulation, requiring tumor cells to express two antigens for controlling CAR T cell activation, sparing normal tissues that express either of the antigen alone [156]. This approach was further validated with co-targeting of ROR1⁺ tumors expressing EpCAM or B7-H3 for reduced toxicity to normal tissue [157]. One additional strategy for improved target selectivity of tumors is the use of "NOT"-gate inhibitory CAR T cells (iCARs), which use checkpoint pathway inhibition of one target while simultaneously activating CAR T cells with another target [158]. These approaches potentially provide further improvements over the previous combinatorial targeting approach mentioned above [140], and are anticipated to enter clinical testing soon for patients with solid tumors.

While requiring dual antigen expression on tumor cells for optimal activation of CAR T cells is an exciting advancement over single antigen-specific CARs, another versatile approach to engineer target specificity, called switchable or universal CARs, has recently been developed. These programmable CARs come in several forms, but each has in common the ability to redirect the specificity of CAR T cells to different antigens based on a druggable reagent. The first example of this strategy was demonstrated using CARs with an antigen-binding domain specific for the common fluorophore FITC, which controlled the activation of CAR T cells to antibodies tagged with FITC, redirecting specificity to EGFR, HER2, or CD20 [159]. Further validation of FITC-specific CAR T cells has been documented [160], as well as for biotinylated antibodies using Streptavidin-specific CAR T cells [161] and peptide neo-epitopes from the yeast transcription factor GCN4 [162, 163]. A more recent iteration of this strategy has been demonstrated using a switch, universal, and programmable (SUPRA) CAR, which employs a leucine zipper as the targeting domain on the CAR, along with an antibody tagged with the cognate leucine zipper [164]. Compared to conventional single or dual-targeted CAR T cells, these modular approaches offer improved safety with robust efficacy of CAR T cell activation, allowing for "smart" targeting of solid tumors.

11.3.5 Improving CAR T Cell Therapies in Immunosuppressive Solid Tumors

Another major challenge for effectively targeting solid tumors with CAR T cell therapies is the immunosuppressive tumor microenvironment. Distinct from most of the hematological malignancies that lack local immunosuppressive pathways that hamper antitumor immunity and limit adoptive T cell therapies, solid tumors can be heavily infiltrated by multiple cell types that support tumor growth, vasculature, metastasis, and may dictate therapeutic responses [165]. The most prominently studied cell types that drive immunosuppression in tumors are regulatory T cells (Tress), M2 tumor-associated macrophages (TAMs), and myeloid-derived suppressor cells (MDSCs) [166]. These immune cell infiltrates, in addition to the tumor cells themselves, drive local cytokine, chemokine, and growth factor production in solid tumors, including IL-4, IL-10, VEGF, and TGFB, that can facilitate tumor growth and progression. Likewise, immune checkpoint pathways, including PD-1 and CTLA-4, can be highly active in tumors to dampen antitumor immunity. Considerable evidence suggests that the tumor microenvironment also controls response and resistance to immunotherapies [167], and can limit the effectiveness of CAR T cell therapy [168].

A number of recent studies have aimed to boost CAR T cell functionality by blocking immune checkpoint pathways. Multiple studies have demonstrated that following CAR T cell therapy, PD-1/PD-L1 and other checkpoint pathways are induced, thereby limiting durable therapy [169]. The simplest of these methods has been demonstrated by combining CAR T cells with immune checkpoint blockade [170–172]. Phase 1 clinical trials are underway evaluating this combination approach to improve response rates in hematological malignancies and solid tumors [173, 174] (NCT03545815). Novel strategies to intrinsically circumvent PD-1/PD-L1 signaling pathways to prolong CAR T cell functionality have been explored. For example, a chimeric PD1-CD28 receptor allowed for redirecting PD-1-signaling in T cells towards co-stimulation [175, 176]. Cherkasskey and colleagues evaluated multiple methods of intrinsic blockade of PD-1 in CAR T cells, including shRNA knockdown of PD-1 or a PD-1 dominant negative receptor, showing improved antitumor responses in multiple preclinical models by blunting PD-1 signaling in adoptively transferred T cells [177]. More recently, the secretion of PD1 blocking antibodies by CAR T cells was shown to similarly improve therapy [178, 179]. CRISPR/Cas9-mediated disruption of PD-1 in CAR T cells has also been explored, and clinical trials are now underway evaluating this approach in patients [180-182]. In the context of the most well-studied PD-1 and CTLA-4 inhibitors, it has been demonstrated that potential mechanisms of tumor resistance include compensatory upregulation of alternative immune checkpoint pathways. Therefore, it will be imperative to evaluate and overcome these resistance mechanisms in the context of combinatorial CAR T cell - immune checkpoint blockade.

Expression of TGF β , a multi-functional cytokine that is dysregulated in many cancers, has been associated with an immune phenotype characterized by a lack of tumor T cell infiltration [183]. Hence, a recent pursuit has been dedicated to

blocking TGF β signaling in CAR T cells and in the immunosuppressive tumor microenvironment to promote adoptive and adaptive T cell antitumor immunity. Preclinical studies suggest that CAR T cells containing a CD28 co-stimulatory domain may resist TGF_β-mediated inhibitory signals predominantly through IL-2 signaling [184]. Despite recent evidence pointing to superior T cell persistence and antitumor activity, 4-1BB-containing CAR T cells may lack the ability to resist TGF β -mediated immunosuppression. Therefore, CAR T cells engineered to be refractory to immunosuppressive factors present in the tumor microenvironment, including TGFB, have been developed [185]. Based on these strong preclinical findings, a phase 1 clinical trial has been initiated to evaluate PSMA-targeted CAR T cells with a dominant negative TGF β receptor in patients with metastatic castration-resistant prostate cancer (NCT03089203). Other approaches include redirecting TGF β signaling in T cells towards 4-1BB co-stimulation [186] or IL-12 signaling [187] using chimeric receptors. Uniquely, CAR T cells targeting soluble TGF β have also been engineered [20], which can be used in a dual-targeted CAR T cell approach to simultaneously target tumor cells and inhibit TGFB signaling [188].

In addition to engineering CAR T cells to block inhibitory signals in the immunosuppressive tumor microenvironment, the expression of pro-inflammatory cytokines with the ability to shape the tumor microenvironment for improved T cell trafficking, survival, persistence, and antitumor functionality has been explored. The earliest example of this strategy was shown using CD19-CAR T cells engineered to secrete IL-12. In addition to increased IFNy production, CAR T cell persistence, and overall therapeutic activity, this therapy also eliminated tumors in absence of lymphodepleting preconditioning [189]. IL-12 secreting the MUC16-directed CAR T cells also produced elevated levels of IFNy, increased survival and persistence of CAR T cells, and improved overall therapy in xenograft models of ovarian cancer [91]. Follow-up studies in immunocompetent mice showed that IL-12-secreting MUC16-CAR T cells also shaped the immunosuppressive microenvironment in ovarian cancers by depleting tumor-associated macrophages and overcoming PD-L1-mediated T cell inhibition [190]. These preclinical studies have resulted in a clinical trial testing this approach in MUC16⁺ solid tumors (NCT02498912). CD19-CAR T cells have also been engineered to express IL-15 tethered to the surface of T cells (mbIL-15). mbIL-15 CAR T cells showed improved stem/memory phenotype with increased T cell persistence and durable antitumor activity [191]. Alternative platforms for intrinsic IL-15 production by CAR T cells have been investigated, including CAR T cells engineered to secrete soluble IL-15 [192], and a novel nanoparticle drug delivery platform carrying an IL-15 super-agonist complex [193]. Other approaches have introduced novel ways to redirect immunosuppressive cytokines toward pro-inflammatory pathways, including CAR T cells with chimeras in which the IL-4 receptor ectodomain is fused to the IL-7 receptor endodomain. This platform was utilized in xenograft models of pancreatic cancer using PSCA-directed CAR T cells [194]. A similar strategy was utilized to redirect IL-4 signaling towards another pro-inflammatory cytokine, IL-21 [195].

The immunosuppressive tumor microenvironment, in addition to suppressing the function of CAR T cells once they arrive at the tumor site, likely also intrinsically blocks trafficking of CAR T cells. Therefore, in addition to increasing doses of infused CAR T cells to achieve a required threshold of recruitment at the tumor site, combination approaches to amplify endogenous immunity to aid in CAR T cell responses have been explored. Oncolytic viruses (OV) can be selectively programmed to target, infect, and kill cancer cells, and genetically modified to express therapeutic genes selectively in the tumor microenvironment [196, 197]. Through cancer cell infection and lysis, OV has been used for tumor debulking, reversing tumor immunosuppression, and initiating systemic antitumor immune responses. Watanabe and colleagues showed that the combination of mesothelin-targeted CAR T cell therapy with an oncolytic adenovirus driving tumor expression of TNFα and IL-2 induced significant tumor regression in a syngeneic mouse model of pancreatic cancer. This antitumor response was accompanied by an increase in CAR T cell and endogenous T cell infiltration, pro-inflammatory M1 macrophage polarization, and dendritic cell maturation [198]. Additional studies have utilized OV to express multiple transgenes in cancer cells simultaneously, consisting of immune checkpoint inhibitors and pro-inflammatory cytokines, that, when combined with CAR T cells, showed enhanced T cell effector function [199]. These findings indicate that combining cytokine-armed oncolytic adenoviruses to enhance the efficacy of CAR T cell therapy is a promising approach to overcome the immunosuppressive tumor microenvironment and to also amplify endogenous antitumor immunity.

11.3.6 Pre-existing T Cell Immunity and CAR T Cell-Induced Endogenous Immunity

Current understanding suggests that the effectiveness of immunotherapy depends on the presence of pre-existing immunity and the ability to effectively modulate the baseline immune response. Clinical studies are beginning to define predictive tumor and immunological factors governing the anticancer response—one such measure is the immune classification of cancer.

The immune classification of cancer is an evolving measure that characterizes tumors with respect to their immune infiltration in two broad classifications: immunologically "hot" and immunologically "cold" tumors (Fig. 11.2). Immuno-logically hot, or immune-inflamed tumors, are characterized predominantly with a high infiltrate of T cells, low infiltration of immune-suppressive cells including regulatory T cells (T_{reg}) and myeloid-derived suppressor cells (MDSC) and include additional features like PD-L1 expression on tumor cells and tumor-associated immune cells, potential genomic instability and the presence of a pre-existing antitumor immune response. Immunologically cold, immune-excluded, or immune-deserted tumors typically have poor antitumor T cell infiltration, high immune-suppressive cell infiltration, low PD-L1 expression, with high proliferation of cancer cells and low mutational burden [167]. Studies have recently proposed a



Fig. 11.2 The immune landscape of solid tumors. **a** A representative immunologically "hot" tumor containing a high frequency of antitumor CD8 T cells, and a relatively low frequency of immunosuppressive regulatory T cells (T_{reg}) and myeloid cell subsets including tumor-associated macrophages (TAM), mononuclear myeloid-derived suppressor cells (MO-MDSC), neutrophils, and polymorphonuclear myeloid-derived suppressor cells (PMN-MDSC), along with tumor vasculature and stromal cells.**b** A representative immunologically "cold" tumor containing a higher frequency of immunosuppressive cell subsets and a relatively low frequency of antitumor CD8 T cells

combinatorial set of parameters to augment this classification: the T cell phenotype (follicular helper T (Tfh), T helper 1 (Th1), memory and exhausted T cells) at the tumor, dependent on location (invasive margin, tumor core, and tertiary lymphoid structures), density (immune density and quantity), and functional immune orientation (chemokines, cytokines, cytotoxic factors, adhesion, attraction) [200]. These factors combine to represent cancer immune interactions for an individual patient [201], and together, they can help define immunomodulation strategies to optimize personalized treatment choices [202].

It has yet to be determined whether antitumor responses with CAR T cell therapy is impacted by pre-existing T cell immunity. In the context of immune checkpoint blockade, response to therapy may rely on the reactivation of pre-existing T cells, the recruitment of new T cells to the tumor, or a combination of both [203, 204]. T cell exhaustion represents a distinct state of T cell differentiation and can be driven by cell signaling, prolonged TCR engagement, co-stimulatory/inhibitory signals, soluble factors (e.g. excessive suppressive cytokines), and microenvironment features (e.g. chemokine receptor expression, adhesion molecules). Exhausted T cells acquire an epigenetic profile that is distinct from T effector cells, and despite the ability to revert to an effector using PD-1 blockade, these cells may never acquire a memory phenotype [204]. This limits the durability of immunotherapy, and an understanding of how to permanently reverse T cell exhaustion is currently incomplete. These phenotypes may heavily impact CAR T cells once they arrive at tumors, and may overcome in part by addressing immunosuppression, as covered in the section above.

The presence and density of tumor-infiltrating lymphocytes (TILs) are often interpreted as an indication of pre-existing T cell immune recognition, though recent studies have highlighted that reactivity of TILs with respect to cognate tumor antigens is rare and variable [205]. A recent study that analyzed phenotype and TCR repertoire in site matched tumors, from basal or squamous cell carcinoma patients, pre- and post-therapy showed that response to PD-1 blockade associated with the expansion of a distinct repertoire of T cell clones from pre-therapy TILs [206]. Together, these studies suggest that increasing the frequency and breadth of the tumor-specific TCR repertoire may be critical to boost the response towards immunotherapy, thereby increasing infiltration of tumor reactive T cells, and amplifying secondary immune responses. These studies also indicate that priming the tumor microenvironment prior to, and during, CAR T cell therapy may greatly impact the overall antitumor responses and provide for more durable clinical outcomes in patients.

One suggested mechanism by which adoptive T cell therapy is able to promote durable antitumor responses is through the stimulation of epitope spreading—a dynamic process that underlies the expansion of an immune response to secondary epitopes that are not targeted by therapy. In particular, epitope spreading may be initiated by the presence of a tumor-specific endogenous immune response responsible for the release of immunosuppressive mechanisms and promotion of T cell chemo-attracting cytokines at the tumor site. In the context of CAR T cell therapy, this resulting immune recruitment may confer the ability to produce a secondary immune response to cancer cells that do not express the CAR target antigen.

The potential for CAR T cells to induce epitope spreading has not been extensively studied with the exception of a few preclinical studies. In a murine CAR model targeting EGFR⁺ glioblastoma, mice that were cured of EGFR⁺ tumors later rejected EGFR-tumors when re-challenged, suggesting the generation of endogenous immunity against additional tumor antigens [207]. Pituch and colleagues showed significant changes in the tumor microenvironment and endogenous immune infiltration after IL13R α 2-CAR T cell therapy in an immunocompetent mouse model of malignant glioblastoma [208]. These changes included a decrease of immunosuppressive MDSCs and an increase in both endogenous CD4⁺ and CD8⁺ T cells, as well as CD8 α^+ dendritic cells. The presence of these factors along with a lack of tumor development upon re-challenge with an IL13R α 2 negative tumor, suggests these mice could acquire antitumor immunity in response to CAR T cell therapy. Modifications to the cytokine/chemokine expression of CAR T cells, namely inclusion of IL-7 and CCL19, resulted in superior antitumor activity coupled

with increased endogenous immune infiltration and protection against CAR-targeted antigen-negative tumor growth [209]. These preclinical studies have underscored that CAR T cell therapy may not only modulate the immune landscape by creating a pro-inflammatory tumor microenvironment, but also recruit endogenous antitumor immunity in response to CAR T cell therapy.

Recent clinical studies have suggested that CAR T cells show evidence for inducing a secondary immune response. A first-in-human study of intravenous delivery of EGFRvIII-CAR T cells reported that the CAR T cells trafficked to the brain tumor proliferated, and exerted some bioactivity in patients with recurrent glioblastoma [44]. Although the T cell receptor clonotypes present in the CAR T product were a large fraction of the T cell repertoire infiltrating the tumor after CAR T infusion, a significant portion were not, suggesting that CAR T cell infusions could potentially increase endogenous TCR repertoire diversity to the tumor, with the potential to induce a secondary immune response targeting secondary epitopes on EGFRvIII- tumor cells [44]. CAR T cell-mediated epitope spreading was suggested in a patient with recurrent multifocal glioblastoma that received IL13R α 2-CAR T cells [43]. Following 10 intraventricular infusions, regression of all intracranial and spinal tumors with a continued clinical response in the patient for 7.5 months was observed. Evidence of endogenous T cell recruitment and stimulation in the CSF after every CAR T cell infusion was associated with increases in T cell chemo-attractants CXCL9/CXCL10, as well as IFNy.

Together, these studies suggest that CAR T cells have the capacity to amplify an inflammatory immune response and recruit endogenous T cells to tumor sites. Increasing the frequency and breadth of the tumor-specific TCR repertoire at tumor sites may be critical to boost CAR T cell therapy by inciting a secondary immune response. This phenomenon will likely be an important component of durable clinical outcomes in patients with single or multi-targeted CAR T cell therapy to ultimately overcome resistance driven by heterogeneity and in solid tumors.

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