

Review





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Immunosuppression in tumor immune microenvironment and its optimization from CAR-T cell therapy

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Abstract

Chimeric antigen receptor (CAR)-T cell therapy represents a landmark advance in personalized cancer treatment. CAR-T strategy generally engineers T cells from a specific patient with a new antigen-specificity, which has achieved considerable success in hematological malignancies, but scarce benefits in solid tumors. Recent studies have demonstrated that tumor immune microenvironment (TIME) cast a profound impact on the immunotherapeutic response. The immunosuppressive landscape of TIME is a critical obstacle to the effector activity of CAR-T cells. Nevertheless, every cloud has a silver lining. The immunosuppressive components also shed new inspiration on reshaping a friendly TIME by targeting them with engineered CARs. Herein, we summarize recent advances in disincentives of TIME and discuss approaches and technologies to enhance CAR-T cell efficacy via addressing current hindrances. Simultaneously, we firmly believe that by parsing the immunosuppressive components, and optimizing CAR-T cell therapy for each patient, the CAR-T cell immunosuppressive components, and malignancies will be substantially enhanced, and novel therapeutic targets will be revealed.

Key words: Tumor immune microenvironment, immunotherapy, chimeric antigen receptor T cell, immunosuppression network, solid tumors

1. Introduction

The past decade has witnessed a revolutionary evolution in the application of immunology to oncology treatment. Immunotherapy has made striking advances in malignancies, far outperforming conventional chemotherapy and radiotherapy. Frustratingly, the immunosuppressive properties of the tumor microenvironment (TME) pose the challenge of limited clinical efficacy and severe side effects to tumor immunotherapy [1]. Priming the tumor immune microenvironment (TIME) is a critical next step in scaling up the success of current immunotherapies [2]. Emerging evidence suggests that the effectiveness of immunotherapy can be maximized by addressing the tumor immunosuppressive microenvironment [3, 4]. Hence, a more detailed dissection and description of immunosuppression in TIME and a deeper understanding of the tumor immunosuppressive profile are necessary for the development and optimization of novel and effective cancer immunotherapy.



Figure 1. CAR-T cell therapy and tumor immunosuppressive microenvironment. A After isolating T cells from the peripheral blood of the patient, engineering the CARs genes into T cells to generate CAR-T cells. Then CAR-T cells are extensively expanded *in vitro* and administered to the patient. **B** TME is the central mediator of tumorigenesis and tumor-promoting function. The solid tumor microenvironment including the extracellular matrix, various immune cells, abnormal tumor vasculature, immunosuppressive molecules, and tumor metabolites prevents CAR-T cells from exerting high cytotoxicity. The tumor-associated stroma such as fibroblasts and mesenchymal cells formed physical barriers against the entry of T cells. The migration of T cells towards tumor lesions was increasingly challenged by dysregulation of adhesion molecules, mismatching of tumor-derived chemokines, and immune cell-expressed chemokine receptors. In addition, the metabolically abnormal TME impeded immune cells and bulk masses of immunosuppressive factors orchestrate a severely immunosuppressive tumor milieu to suppress the efficacy of CAR-T cells.

Chimeric antigen receptor (CAR)-T cell therapy immunotherapy through genetically is novel engineered T cells to express CAR targeting molecules. CAR-T cell therapy aims at introducing CARs genes into patient-derived T cells of peripheral blood in a fairly short period, where the biological properties of T cells are redirected and reprogrammed [5]. T cells are rapidly expanded to derive memory and effector lymphocytes with high affinity in vitro. These T cells are then infused back into the patient to proliferate robustly and elicit potent anti-tumor activity (Figure 1A). These synthetic receptors recognize their corresponding specific antigens using a single-chain variable fragment (scFv) from the variable region in a major histocompatibility complex-independent manner. The majority of scFv possesses binding properties similar to antibodies. CAR-T cell therapy has shown dramatic clinical responses and high rates of complete remission in hematologic malignancies [6]. However, therapeutic effects in solid tumors are not durable, partly owing to physical barriers, cancer heterogeneity, and TIME, which lead to recurrent CAR antigen loss and rapid CAR-T cell exhaustion [7]. Recent strategies constantly focus on harnessing CAR-T cell therapy via engineering CARs, T cells, and interactions with other elements of TIME. Among these, remodeling TIME is one of the most attractive strategies for

promoting the endogenous immune response to achieve a permanent CAR-T cell engagement [8].

The immune characteristics of the tumor microenvironment (TME) have been categorized as one of the ten tumor characteristics [9], which play a decisive role in predicting the clinical outcomes of patients [10]. Recognizing the essence of TIME has paramount implications for battling cancer cells. Here, we review evidence for each of the following perspectives. Firstly, how tumors orchestrate an immunosuppressive microenvironment promoting immune tolerance and evasion. Secondly, how to enhance the effectiveness of CAR-T cell therapy. Ultimately, we further elucidate the utilization of refreshing technologies to reverse the immunosuppressive microenvironment.

2. The Intricate Network Sustaining Immunosuppression in TIME

TME contributes to unfavorable immunotherapy efficacy by preventing CAR-T cells from exerting high cytotoxicity against tumor cells **(Figure 1B)**. As proof: 1) The tumor-associated stroma such as fibroblasts, mesenchymal cells, and various extracellular matrices formed stumbling block against the entry of T cells [11]; 2) The migration of T cells towards tumor lesions was increasingly overshadowed by the 'bad guys', such as dysregulation of adhesion molecules, aberrant tumor-related vasculature, and mismatching of chemokines and their receptors [11]; 3) Cancer cells expressed ligands of suppressive immune checkpoints such as immune-dampening PD-1 ligand (PD-L1)/L2 [12] and recruited more immunosuppressive cells [13, 14] to interfere with the effector T cells (Teffs) cytotoxic function. Additionally, the metabolically abnormal TME characterized by restricted nutrient availability, acidosis, and local hypoxia, impeded immune cell activity [15, 16]. Nevertheless, these straitened circumstances have in turn spurred the design of CAR-T cells to better unleash the potential of immunotherapy. Generally, TME is the central mediator of tumorigenesis and tumor-promoting function, with immunological features possessing indispensable status. The complex immunosuppressive network in TME, that is tumor immunosuppression microenvironment, consists of miscellaneous immunosuppressive cell subsets, secretions, and signals that inhibit the recruitment, proliferation, differentiation, and execution of effector functions of immune cells. Here, we focus on the immunosuppressive landscape of TIME (Figure 1C).

2.1. Representative immunosuppressive cells in TIME

Tumor-associated immune cells possess crucial functions in tumorigenesis, which antagonize and/or promote tumors. Homogeneous immune cells could reshape themselves depending on different tumor ecosystems (Figure 2).

2.1.1 Myeloid-derived suppressor cells (MDSCs)

MDSCs are pathologically activated monocytes and neutrophils with vigorous immunosuppressive behavior and are implicated in the negative modulation of immune responses and poor clinical outcomes [17]. An ocean of extracellular factors could induce MDSCs differentiation or expansion, encompassing granulocyte-macrophage colony-stimulating factor (GM-CSF), macrophage colony-stimulating factor (M-CSF), interferon-gamma (IFN-v), interleukin (IL)-6, prostaglandin E2 (PGE2), IL-13, and vascular endothelial growth factor (VEGF). In contrast, IL-4 and all-trans-retinoic acid can inhibit this procedure [18, 19]. MDSCs could be classified into granulocytic **MDSCs MDSCs** (G-MDSCs) and monocytic (M-MDSCs) according to the phenotypic and morphological characteristics or cell surface markers of MDSCs. M-MDSCs hindered CD8+ T cells via an inducible nitric oxide (iNOS)-mediated pathway [20-22], whereas G-MDSCs suppressed T-cell function through arginase and/or reactive oxygen species (ROS)-dependent mechanisms [23, 24].

Apart from promoting tumor development by forming pre-metastatic niches and angiogenesis [25, 26], MDSCs could also contribute to immunosuppression through the following mechanisms: 1) Inducting other immunosuppressive cells: MDSCs could not only secret IL-10 and transforming growth factor- β (TGF- β) to directly hamper Teffs but also induce the *de novo* generation of regulatory T cells (Tregs) mediated by IL-10 and IFN-y in vivo [21]. CCR5 ligands CCL5, CCL4, and CCL3, produced by tumor-infiltrating M-MDSCs could recruit massive amounts of CCR5+ Tregs [27]. They could also shift macrophages from M1 to an M2-like state with immunosuppressive features. It was demonstrated that MDSCs-produced IL-10 decreased macrophage IL-6 and TNF-a [28]. 2) Blocking lymphocyte homing: splenic or blood-borne MDSCs were shown to execute far-reaching immune suppression through downregulating L-selectin lymph node homing receptors on naïve T and B cells. Furthermore, loss of



Figure 2. Miscellaneous immune cells inside tumors. The tumor-associated immune cells may possess tumor-antagonizing or tumor-promoting capacities. The homogeneous immune cells are able to change their state due to different tumor ecosystem or different stages of tumorigenesis. For instance, macrophages can shift from a hostile to a friendly status after immunotherapy. Elevating the amount of tumor-antagonizing immune cells opens a broad window for tumor treatment.

L-selectin expression could disrupt T cell trafficking. T cells preconditioned by MDSCs have diminished responses to subsequent antigen exposure [29, 30]. 3) Engendering reactive oxygen and nitrogen species: secreting ROS such as hydrogen peroxide, hydroxyl radicals, and superoxide anions was a well-known strategy to eradicate tumor-infiltrating lymphocytes. Enhanced ROS levels could increase the quantity and quality of tumor-infiltrating MDSCs by NF erythroid 2-related factor 2 and VEGF receptors, which might create a positive feedback loop [31, 32]. MDSCs could also generate high levels of reactive nitrogen species (RNS) via activating the iNOS pathway [33]. RNS could induce chemokine CCL2 nitration, hinder T-cell infiltration and engage T-cell apoptosis, leading to the trapping of specific T-cells in the tumor-associated stroma [34, 35]. 4) Competing with immune cells for nutrient metabolites: high expression of arginase I and cationic amino acid transporter 2B by mature MDSCs could rapidly incorporate L-Arginine (L-Arg) and deplete extracellular L-Arg in vitro [36]. L-Arg depletion blocked antigen-specific proliferation of OT-1 and OT-2 cells and the re-expression of CD3zeta in stimulated T cells [37]. 5) Another mechanism by which MDSCs inhibit immune cells included the expression of ectoenzymes regulating adenosine metabolism from ATP [38] and negative immune checkpoint molecules [39].

2.1.2 Tumor-associated macrophages (TAMs)

Proverbially, macrophages are essential components of innate immunity and leukocyte infiltration present in solid tumors, which regulate cancer-related inflammation and constitute vital regulators of tumor initiation and progression. Circulating monocytes could be recruited into the tumor stroma by multiple chemokines and cytokines such as CCL2, GM-CSF, and VEGF family members [40, 41]. TME then prompted the differentiation of monocytes into TAMs [42]. The immunological effects of TAMs are pro- and anti-tumor functions based on the state of macrophage activation. Collectively, M1 macrophages could respond to danger signals that produced type I pro-inflammatory cytokines such as IL-1, IL-12, and TNF-a. Conversely, M2 macrophages expressed scavenger receptors and type II cytokines, such as IL-4, IL-10, and IL-13, promoting antiinflammatory responses [43-45]. M2 macrophages possessed pro-tumorigenic functions including promotion of tumor cell growth [46], drug resistance [47] and metastasis [48], neo-angiogenesis [9, 49], and immune suppression [50]. The M1/M2 balance varied with cancer types. Both M1 and M2 macrophages have high levels of plasticity, with the ability to be converted into each other upon TIME or therapeutic

interventions [51, 52]. For example, hepatocellular carcinoma (HCC)-derived exosomes could reshape macrophages and result in M2-polarized TAMs via inducing pro-inflammatory factors and activating NF-κB signaling [53].

TAMs could regulate the immunological activity of Т cells to foster tumor progression. Pro-inflammatory cytokines produced by TAMs could trigger the accumulation and expansion of CD4+ Th17 cells to foster angiogenesis and overexpressing CTLA-4, programmed death-1 (PD-1), and glucocorticoid-induced tumor necrosis factor receptor (GITR), thereby promoting tumor development [54]. TAMs could also attract Tregs into tumor tissues by producing various chemokines, such as CCL17, CCL18, and CCL22 [55, 56]. Additionally, they could directly handicap the proliferation of CD8+ T cells through the metabolism of L-arginine and the production of iNOS and ROS [34, 57].

TGF- β in TME upregulated Tim-3 expression on TAMs, which promoted tumorigenesis and tolerance via NF- κ B signaling and downstream IL-6 production [58]. It has been also validated that TAMs could produce IL-6 and signal via STAT3 to facilitate the expansion of carcinoma stem cells sustaining carcinogenesis [59]. Likewise, TAMs were demonstrated as a nexus with the prognosis of numerous tumors, such as HCC [60] and breast cancer [61].

2.1.3 Tregs and regulatory B cells (Bregs)

Commonly referred to Tregs as are which CD4+CD25+Foxp3+ cells, Т could chemoattract TME through chemokine gradients such as CCR8-CCL1, CCR4-CCL17/22. Tregs negatively moderate immune responses to maintain autoimmune tolerance and homeostasis. However, excessive suppression of immune responses in the TME promoted tumor progression [62]. Abundant infiltration of Tregs positively correlated with depressed survival in various tumor types, such as pancreatic and melanoma [63]. Intratumoral Tregs were highly immunosuppressive and consumed IL-2 through the high expression of CD25 (IL-2 receptor subunit- α), thereby limiting the activation and proliferation of Teffs dependent on IL-2. Tregs also inhibited and/or killed Teffs by releasing suppressive molecules and producing cytotoxic substances [64, 65]. Upregulating immune checkpoint molecules including the lymphocyte activation gene-3 (LAG-3), CTLA-4, PD-1, and inducible co-stimulator (ICOS), was an alternative strategy for Tregs to suppress Teffs [64].

B cells in TME play a double-edged sword effect either by promoting tumor immunity or enhancing tumorigenesis [66]. It has been shown that any B cell has the potential to differentiate into Bregs. Bregs prohibited the expansion of T cells and other immune-system pro-inflammatory lymphocytes to exert suppression. Despite partial consensus on the immunosuppressive effector functions of Bregs, the field has not yet reached a unified view on their phenotype [67]. Immunohistochemical analyzes showed that the frequency of CD19+IL-10+ Bregs in tongue squamous cell carcinoma was significantly higher than adjacent normal tissue. Increased Bregs could convert CD4+CD25- T cells into CD4+Foxp3+ Tregs in cytological experiments and are associated with cancer progression and worse survival [68]. Correspondingly, B cells enriched in the ascites from ovarian cancer patients were inversely correlated with the frequencies of IFN-g+CD8+ T cells, but positively correlated with Tregs [69]. Another study showed that bone marrow-derived Bregs could abrogate NK cell cell-mediated antibody-dependent cytotoxicity against multiple myeloma cells [70].

2.1.4 Others

Tumor-associated neutrophils (TANs) are another type of immune cell that infiltrates various tumors. TANs could be recruited to TME by IL-8 via CXCR1/CXCR2 receptors and abolish the ability of CD8+ T cells through TNF-a production-mediated NO [71, 72]. Similar to TAMs, TANs are partitioned into anti-tumoral (N1) and pro-tumoral (N2) phenotypes. Type I IFNs could alter neutrophils into N1 phenotypes, which transform the pro-tumor properties of low neutrophil extracellular traps and TNF-a expression into anti-tumor milieus. The constantly changing TIME drove the polarization of TANs. Altered TANs were connected with different prognoses in cancers and modulation of TANs phenotypes may represent a potent therapeutic option [73, 74]. More significantly, TANs infiltration or neutrophil-lymphocyte ratio strongly correlates with cancer development, which may have utility as a predictive biomarker to monitor cancer patients receiving immunotherapy (e.g., melanoma, metastatic renal cell cancer) [75, 76].

Mast cells (MCs) are well known to participate in allergy and inflammation. The ability of MCs in TME has been hypothesized to be either pro- or anti-tumor [77]. The number of infiltrating MCs correlated positively with poor cancer prognosis [78, 79]. Recently, Leveque et al. have demonstrated that tumor-associated mast cells (TAMCs), а heterogeneous population, harbor а distinct phenotype compared to MCs present in the non-lesional homologue of lung cancer [80]. It is further noted that the TAMCs subset expressing alpha E integrin, namely CD103+ TAMCs, appeared more

actively to interact with CD4+ T cells and located closer to tumor cells than their CD103- counterparts. Recruitment of TAMCs was mediated by chemotactic agents released by TME, such as VEGF, CXC chemokine ligand (CXCL)12, PGE2, and platelet-derived growth factor (PDGF) [80, 81]. Activated TAMCs could express CCL5 and IL-33 to further recruit MCs into tumor sites and activate themselves in an autocrine manner. TAMCs could drive angiogenesis, immunosuppression, as well as tumor invasion and metastasis through secreting substantial proteolytic enzymes and growth factors [79, 81]. Nevertheless, some studies have shown that a high frequency of TAMCs correlated with better progression-free survival and overall survival [79, 80]. Together, the immunosuppression of TAMCs in TME remains relatively vague and controversial.

Dendritic cells (DCs) acting as specialized antigen-presentation cells have long been recognized as a critical factor in T-cell-mediated anti-tumor reactions. DCs uptake and cross-present tumor antigens to naïve T cells accompanied by priming and activating CTLs with the ability to eradicate tumor cells [82]. Intriguingly, although tumor-associated dendritic cells (TADCs) could prevent prolongedly steady tumor expansion at early stages, the enduring activity of Т cells was abrogated by microenvironmental immunosuppressive TADCs at late stages, becoming less responsive [83, 84]. The tumor-antagonizing role of DCs within TME faced inauspicious roadblocks[85]. The chemotactic gradient of TME caused a paucity of DCs recruitment and a constellation of immunosuppressive factors accelerated tolerance and DCs dysfunction [86, 87]. Numerous oncogenic signaling axes could restrict the ability of DCs. Ruiz et al. reported that HCC impaired DCs recruitment due to the absence of a tumor-derived chemokine CCL5 via tumor-intrinsic Wnt/ β -catenin signaling [88]. STAT3-mediated signaling also handicapped the differentiation and maturation of DCs by producing IL-10, TGF- β , and indoleamine 2,3-dioxygenase (IDO) [89]. could Furthermore, **TADCs** express immune-inhibitory checkpoints, such as PD-L1 and TIM-3, to impede the activation of CTLs [88, 90]. TADCs could also spark tumor plasticity, growth, and metastasis promoting genomic instability, by neovascularization, and immunometabolism [91, 92].

2.2 The intricate immune inhibitory factors network in TIME

Immune cells, endothelial, stromal, and tumor cells within the TME secreted bulk masses of immunosuppressive factors including chemokines, cytokines, and inhibitory molecules to assist and/or restrain each other, which orchestrates a severely immunosuppressive tumor milieu. These factors bind to corresponding receptors via paracrine, autocrine or endocrine means to modulate tumor growth/invasion/metastasis, and immune responses, thereby reshaping TME and mediating intercellular crosstalk. Some of their specific features have been mentioned above. Herein, we dissect the representative participants in the current data (**Table 1**).

Table 1. Representative immune inhibitory factors mediated pro-/anti-tumor function

Molecules and signaling pathways	Category	Description	References
IL-7, IL-15, IL-21	Pro-inflammatory cytokine	The common γ chain cytokine family	[93, 94]
IL-18	Pro-inflammatory cytokine	 Promote the generation of the stem cell-like memory T cell phenotype Boost the tumor-killing activity of NK and CTL cells A potent inducer of IFN-γ Consider to T and NK cell activities and The Leell palarisation 	[95, 96]
		·Contribute to 1 and INK cell activation and 1n-1 cell polarization	
IL-12	Pro-inflammatory cytokine (anti-tumor immunity modulator)	Activate NK cells and T lymphocytes and induce Th-1 type responses Increase IFN-γ secretion and cytotoxicity	[97]
IL-4, IL-13	Anti-inflammatory/inhibitory cytokine	•Suppress type 1 immunity and cytotoxic T cell development •Induce expansion of monocytes or macrophages within the tumor stroma and antagonize the tumor-suppressing activity of type 1-activated macrophages	
IL-6/STAT3	Pro-inflammatory/carcinogenesis signaling	· Induce MDSCs differentiation or expansion, then result in immunosuppression ·Favor the expansion of carcinoma stem cells sustaining carcinogenesis · Up-regulate IDO production; down-regulate IFN-γ; induce T cells apoptosis and dysfunction	[18, 19, 59]
PD-1/PD-L1	Immune checkpoint molecules	· Expressed on activated T lymphocytes and deliver immunosuppressive signals, drive T cells into a state of exhaustion, tolerance, or dysfunction	
LAG3	Immune checkpoint molecule	· An exhaustion marker and inhibitory receptor · Impair CD4+ and CD8+ TILs functions	[64, 98]
CTLA-4	Immune checkpoint molecule	•Exert an inhibitory signal to T cells, and then make T cells with an inactive state •Enhance Tregs activity and IDO and IL-10 productions in DCs	[64, 99]
Tim3/Galectin-9 signaling pathway CXCL9, CXCL10, CXCL10, CXCL11/CXCR3 axis	Immune checkpoint signaling Th1-type chemokines	·Inhibit the activation and function of CTLs and promote immune cells apoptosis ·Produced by tumors and immune cells, augment Teff and NK cell trafficking into tumors	[90, 98] [100, 101]
		 Promote CTLs and NK cells differentiation, and regulate differentiation of naive T cells to T helper 1 (Th1) cells Possess an opposite role in either pro- or anti-tumor responses 	
CXCL12/CXCR4	C-X-C subfamily chemokine and its receptor	•CXCL12 contributes to the migration of plasmacytoid DCs into TME, tumor proliferation, and metastasis •CXCR4 can only combine with CXCL12 and prevents tumor metastasis and development	[102]
CCL2, CCL3, CCL5	C-C subfamily chemokines	·Recruit macrophages, neutrophils, and Tregs into TME, promote the polarization of M2-macrophages ·Assist the accumulation of immunosuppressive MDSCs and tumor metastasis	[27, 40, 41]
CXCR1	C-X-C chemokine receptor family	·Increase neutrophil recruitment ·Bolster proliferation of tumor-initiating cells, and neoplastic mass formation	[72]
CXCL1/CXCR2 signaling	Immunosuppressive signaling axis	•Stimulate tumor proliferation and self-renewal, correlate with pro-angiogenic and cancer-promoting genes of the tumor, involve in tumor metastasis •Amplify its production and remarkably induce both tumor-promoting and	[103]
IDO	Immunosuppressive modulator	immunosuppressive factors · Suppress T cell cytotoxicity and IL-2, IFN-γ, and TNF-α production ·Enhance Treg-mediated immunosuppression, thus creating a tolerogenic milieu in tumor sites	[50, 104]
VEGF, PDGF	Growth factors	 Promote angiogenesis and immune evasion Mediate recruitment of immune inhibitory cells and other pro-inflammatory signals in TME 	[32, 81]
Hypoxia /HIF1a	Tumor metabolic environmental factors/ "oxygen sensor"	• Increase the MDSCs infiltration and suppressive function • Control the PD-L1 expression on cancer cells and PD-1 expression on T cells, accordingly dampening T cell survival and effector functions	[16, 105]
Hydrogen peroxide, hydroxyl radicals, superoxide anions, and nitric oxide	Reactive species (ROS, RNS)	• Lead to immune cell apoptosis and deficient effector differentiation, • Inhibit the proliferation and function of CD8+ T cells • Contribute to MDSC recruitment into the TME, result in differentiation and	[16, 32-34]
Extracellular adenosine/ Adenosine	immunosuppressive molecule/	expansion of MJSCS . Impair the activation, proliferation, survival and cytokine production of T cells and other immune cells.	[106-109]
orginaling	ығышта	Activation of adenosine receptors A2A and A2B on tumor-infiltrating immune cells suppress the anti-tumor activities of these cells Favor tumor progression and escape from anti-tumor immunity	

2.3 Tumor antigen heterogeneity

In addition to the exogenous factors mentioned above (e.g., immunosuppressive cells and factors of the TIME), the endogenous factors (e.g., tumor antigen heterogeneity) could further exacerbate the immunosuppressive landscape of TIME. Identifying tumor-specific antigens as targets is a challenge for CAR-T cell therapy to win the war against solid tumors. The optimal target surface antigen should possess excellent coverage, high expression on solid tumors, and not affect normal cells to avoid on-target and off-tumor cross-reactions [110]. However, antigenic heterogeneity is a specific feature of solid tumors. Tumor cells could conceal themselves by constantly removing or modifying their representative antigens, which are antigen loss and antigen-low escape, confusing the targeted attacks of CAR-T cells [111, 112]. It has been demonstrated that antigen loss occurred via two distinct mechanisms: antigen escape or lineage switch [113]. Antigen escape, or isoform switch, occurs when cancer cells present different patterns of targeted antigens without being detected by CAR-T cells. This mechanism is probably connected to gene mutations at the antigen locus due to immune pressure by CAR-T cells [114]. Lineage switch is linked to substantial changes in chromatin accessibility and rewiring of transcriptional programs in tumor cells, including alternative splicing [115]. Antigen-low escape indicated that CAR-T cells targeting specific antigens were unable to effectively eliminate specific antigens-low cells [116]. There is an overlap between antigen escape and low antigen density escape.

3. Up-to-date Strategies for CAR-T Cell Therapy to Address the Hostile Immune Microenvironment

With the increasing understanding of TIME, more strategies concentrate on 're-editing' TIME to better arouse tumor-antagonizing immunity, which offers a novel tactic for CAR-T cell therapy.

3.1 Targeting immune suppressive cells

Several approaches of targeting immunosuppressive cells to advance the tumor-killing ability of CAR-T cells have been applied in basic studies (Figure 3A). It has been reported that MDSCs inhibited the cytotoxicity of different generations of disialoganglioside (GD2)-CAR-T cell. The frequency of circulating MDSCs was inversely correlated with the levels of GD2-CAR-T cell in phase I/II clinical trial, further underlining the importance of targeting immunosuppressive cells [117]. Frustratingly, due to the high similarity of MDSCs to normal myeloid cells, it is extremely challenging to specifically target and eliminate MDSCs from TME without detrimental off-target signaling. Nevertheless, flow cytometry analysis confirmed that TRAIL receptor 2 (TR2) was highly expressed on the surface of MDSCs. By co-expressing a costimulatory TR2.41BB receptor to target both tumor cells and MDSCs, CAR-T cells exhibited enhanced persistence and proliferation [118].

The combination of CAR-T therapy with immunomodulatory agents was proposed for obtaining satisfactory therapeutic results [119, 120]. Sun and colleagues performed co-administration of Olaparib, a poly (ADP-ribose) polymerase inhibitor killing cancer cells by participating in DNA defect repair pathways, in combination with CAR-T cells [121]. The co-administration enhanced anti-tumor responses by impeding MDSCs migration through the SDF1a/CXCR4 axis in immunocompetent mouse models of breast cancer. The combination of CAR-T cell therapy with the folate-targeted Toll-like receptor 7 agonists repolarized MDSCs and TAMs from a hostile to the amicable state. Moreover, it also concurrently enhanced the accumulation and activation of both CAR-T cells and endogenous T cells [122].

Engineering CARs to target TAMs is also a pervasive experimental direction. TAMs that expressed appreciable levels of folate receptor beta (FR β) possessed an immunosuppressive M2-like phenotype, which indicated that targeting FR β could limit the outgrowth of solid tumors [123, 124]. Subsequently, Rodriguez-Garcia and colleagues demonstrated that CAR-T cell-mediated selective elimination of FRβ+ TAMs could augment pro-inflammatory monocyte enrichment and endogenous tumor-specific CD8+ T cell influx, delayed cancer progression, and prolonged survival in syngeneic cancer mouse models [124]. By employing a similar strategy, other researchers designed third-generation CAR-natural killer (NK) and -T cells. They specifically targeted human CSF 1 receptor (CSF1R) on TAMs and had favorable cytotoxicity to remove the inhibitory effect of M2 TAMs [125]. However, Wu and colleagues reported that the effects of GPC3-CAR T cells could be further improved, which can at least partially be ascribed to IL12 secretion of TAMs [119]. This implied that TAMs are highly plastic and heterogeneous. And to some extent, early TIME might exhibit anti-tumor effects, while late TIME was more biased toward tumor promotion [83, 84].

Likewise, depletion of other immunosuppressive cells is an attractive way. Using an anti-CD25 antibody with augmented binding to activating Fc gamma receptor (FcyR, an inhibitory receptor that blocks intra-tumoral Tregs depletion), CD25+ Tregs were depleted. This reinforced cancer-infiltrating CD8+ T cells and improved the eradication of established tumors [126]. The third-(CD28-4-1BBz) generation CAR-T cell that the PYAP Lck binding-motif of CD28 domain incorporated two amino acid substitutions to eliminate IL-2-associated Tregs has been invented. CAR-T cell remarkably retarded cancer growth without a need for lymphodepletion [127]. Alternatively, pretreatment of immune cells to lessen the inhibitory effect may serve as bridging salvage chemotherapy for further CAR-T cell therapy [128]. For instance, daratumumab had successfully treated a case-patient with relapsed/ refractory multiple myeloma-transformed plasma-cell leukemia. The immune cell subset analysis of patients revealed dramatical down-regulation of CD38+ NK cells, Tregs, and Bregs [129]. In mice tumor models, CAR-T cells armed with neutrophil-activating protein (NAP, a pluripotent pro-inflammatory protein) could slow tumor occurrence and development, and bolster survival rates of innate immune cells, regardless of host haplotype, target antigen, and tumor type [130]. Taken together, reprogramming cancer-associated immunosuppressive cells to a more favorable niche could provide new dawn for CAR-T cell therapy.

3.2 Targeting the cytokine and/or chemokine milieu

3.2.1 Cytokines

Induced local dissemination of stimulatory factors could counteract the immunosuppressive dilemma to reinforce CAR-T cell potency (Figure 3B). Currently, recombinant cytokine drugs such as IL-2 and IFN- α , have been approved for anti-tumor treatment. This facilitates the generation of "armored" CAR-T cells engineered to secret pro-inflammatory cytokines. To illustrate, CAR-redirected T-cells tailored to release inducible IL-12 could eliminate the antigen-loss cancer cells, recruit macrophages and reinforce the function, and sustain pro-inflammatory



Figure 3. Gene modification strategies for CAR-T cells. A Targeting immune suppressive cells. CAR-T cells exhibit enhanced persistence and proliferation by targeting MDSCs or depleting Tregs. Likewise, repolarizing MDSCs and TAMs from an immunosuppressive to pro-inflammatory phenotype could boost the anti-tumor functions of CAR-T cells. B Targeting cytokines milieu. CAR-redirected T-cell engineered to release inducible IL-12 or IL-18 could eliminate the antigen-loss cancer cells, recruit immune cells and reinforce their functions, and sustain pro-inflammatory responses. An alternative strategy to counteract immunosuppressive TME is to generate an inverted cytokine receptor in which the IL-4 receptor exodomain was fused to the IL-7 receptor endodomain, which removes the side effects of the immunosuppressive cytokine IL-4. C Targeting chemokines milieu. CXCR1 or CXCR2-modified CARs remarkably favored T-cell migration and persistence in TME, leading to the induction of complete tumor regression and urable immunologic memory. IL-7 and CCR2b co-expressing CAR-T cells also boosted the self-survival and migration, increased IFN-Y, Gzms-B, and IL-2 expression, and inhibited tumor growth. D Targeting immune checkpoints. Blocking immune checkpoints sponsored CAR-T cell survival and function of CAR-T cells. Dominant-negative receptors (DNRs) could interfere with the PD-1/PD-L1 pathway, thereby restoring the cytotoxic and cytokine secretion functions of CAR-T cells.

responses [131]. Combining CAR-T cells with recombinant human IL-12 also fueled anti-cancer activity [132]. IL-18-expressing CAR-T cells have also been shown to possess excellent multiplication power and induce deeper B cell aplasia in mice with B16F10 melanoma [133]. Interestingly, Kunert and colleagues verified that IL-12-secreting T-cell receptor-modified T (TCR-T) cells caused severe edema-like toxicity, augmented blood levels of IFN_Y and TNF α , and decreased numbers of peripheral TCR-T cells, while IL-18-secreting TCR-T cells reduced cancer burden and prolonged survival without side effects [134].

More recently, a preclinical study emphasized that CAR-T cells via autocrine IL-23 signaling had the superior anti-tumor capacity and attenuated side effects, with decreased PD-1 expression and increased granzyme B in comparison to those expressing IL-18 [135]. Gene-engineering expressing IL-15 [136], IL-7 and CCL19 [137], and IL-4/21 [138] has also been explored to boost anti-tumor activities through promoting proliferation, survival, activation, and stem cell memory subset of CAR-T cells and endogenous T cells. The above CAR-T cells equipped with cytokines could reverse immunosuppressive TME not only by extending their expansion and lifespan, but also by activating immune effector cells.

Another approach to addressing the efficacy conundrum is to modify CAR-T cells by rewiring inhibitory inputs to stimulatory outputs or being refractory to inhibitory cytokines present in TME. Engineering of TGF- β CAR-T cells demonstrated that CARs could express a dominant-negative TGF-BRII (dnTGF- β RII), thereby blocking TGF- β signaling in T cells. DnTGF-BRII increased CAR-T cell proliferation and long-term in vivo persistence, bolstered cytokine secretion, fought against exhaustion, and induced tumor eradication [139, 140]. In contrast, CAR-T cells engineered to respond to TGF- β allowed themselves to convert immunosuppressive cytokine into triggers of anti-tumor activity [139, 141]. TGF-β-responsive CAR-T cells could proliferate and produce T helper type 1 (Th1)-associated cytokines in the presence of soluble TGF- β , protect nearby cells from the immunosuppressive effects TGF-β, of and significantly improve the anti-tumor efficacy of neighboring CTLs [139, 141]. Notably, TGF-B was associated with the lack of immune response exhibiting low levels of T cell penetration into the tumor center [142]. Thus, targeting this axis could be the most susceptible solution to destroy solid tumors.

'Switch receptor' could also achieve by rewiring negative signals. Mohammed and colleagues [143] generated an inverted cytokine receptor in which the IL-4 receptor exodomain was fused to the IL-7 receptor endodomain to remove the side effects of the immunosuppressive cytokine IL-4. This result manifested that engineered CAR-T cells could transmit inhibitory signals into therapeutic stimulants through an IL-7-induced downstream pathway and possessed strengthened proliferation and superior anti-tumor ability [143, 144]. Pleasingly, combined single-cell transcriptome analysis and CRISPR-Cas9 knockin nominated a novel TGF- β R2:4-1BB switch receptor to improve CAR-T cell fitness and solid tumor clearance [145].

3.2.2 Chemokines

Given the pivotal role of chemokines and their receptors in recruiting immune cells and trafficking CAR-T cells into tumors, numerous studies have endeavored to integrate chemokines and CAR-T cells to battle tumors in recent years (Figure 3C) [146]. Most investigations focused on CXCR2-modified CAR-T cells. Preclinical models demonstrated that IL-8 receptor, CXCR1 or CXCR2, -modified CARs remarkably favored T cell migration and persistence in TME, thereby inducing complete cancer regression and immunologic memory in aggressive tumors such as ovarian, glioblastoma, and pancreatic cancer [147]. CXCR2-modified CAR-T cells could also accelerate trafficking in vivo and tumor-specific accumulation [148]. Whilding and colleagues came to a similar conclusion that CXCR2-expressing CAR-T cells efficiently migrated towards tumor-conditioned media containing IL-8 and had a more favorable toxicity profile of eliciting anti-cancer responses against $\alpha v \beta 6$ -expressing ovarian or pancreatic tumor xenografts [149].

Targeting other chemokines is under active experimentation, some of which have achieved satisfactory therapeutic effects in solid tumors. Some researchers constructed a lentivirus-based CAR gene transfer system targeting CCR4, a chemokine receptor over-expressed in T-cell malignancies and Tregs profiles [150]. CCR4-expressing directed CAR-T cells displayed antigen-dependent potent cytotoxicity against T-cell malignancies in mouse xenograft models. Wang and colleagues designed co-expressing mesothelin (Msln) and chemokine receptors CCR2b or CCR4 CAR-T cells [151]. The Msln-CCR2b-CAR and/or Msln-CCR4-CAR T cells possessed fortified migration and infiltration, specifically exerted cytotoxicity, and expressed high levels of pro-inflammatory cytokines. Compared with traditional CAR-T cells, IL-7 and CCR2b co-expressing CAR-T cells boosted self-survival and migration, enhanced IFN-y, Gzms-B, and IL-2 expression, and obstructed tumor growth [152]. These strategies all pave the way for the clinical application of CAR-T cells.

Undoubtedly, CAR-T cells engineered to acquire

responsiveness to cytokines, chemokines, and their receptors exhibit stronger tumor-antagonizing and trafficking functions than the classical secondgeneration. CAR-T cells against other soluble inhibitors such as PGE2 [153] and VEGF receptor-2 (VEGFR-2) [154] could also conduce to solid tumor regression, shedding light on the potential novel solid tumors treatment regimens. Even so, individuals treated with the aforementioned CAR-T cells might develop off-target effects and various systemic toxicities such as cytokine release syndrome [155], and neurotoxicity [156], even leading to death [157]. It has been hypothesized that gaining insight into cytokine and chemokine expression profiles across different tumor types and different individuals and exploring how to traffic CAR-T cells into TME with personalized medicine could lessen the severity of toxicities. No matter what, the efficacy in patients is required to be further tested in clinical trials.

3.3 Targeting immune checkpoints

Immune checkpoint receptors and ligands, such as PD-1, PD-L1, and CTLA-4, could preclude CAR-T cell cytotoxicity and induce anergy in TME [158]. Checkpoint blockades are a successful therapeutic intervention to date for solid tumors, which have been used concurrently with CAR-T cell therapy in ongoing clinical trials and obtained potent synergistic activities (Figure 3D) [159]. Combination therapy targeting PD-1 has been demonstrated to promote the survival of CAR-T cell and kill PD-L1+ cancer cells via activation-induced cell death [160]. A multitude of studies had also shown that PD-1/PD-L1 pathway interference, including anti-PD-1 antibodies, cell-intrinsic PD-1 shRNA blockades, or PD-1 dominant-negative receptors (DNRs) restored the cytotoxic and cytokine secretion functions of CAR-T cell through tailored to secrete immune-checkpoint blockades or combination therapy [159, 161, 162]. Moreover, the combination therapy of GD2-CAR-T cells with checkpoint blockades was well tolerable and effective in patients with relapsed or refractory neuroblastoma [163]. Likewise, clinically significant antitumor responses following PD-1 blockade combined with CAR-T cells have been reported in a patient with refractory diffuse large B-cell lymphoma and progressive lymphoma [164].

Gene-editing technologies that knock out immune checkpoints could also be applied to abrogate the expression of T cell negative regulators. PD-1 knockout through TALEN technology augmented the persistence and tumor clearance capability of intratumoral T cells and established durable anti-tumor memory [165]. In addition, CRISPR-Cas9-mediated PD-1 disruption enhanced CAR-T cell cytokine production and cytotoxicity towards PD-L1+ cancer cells without attenuating the proliferation [166]. Using the CRISPR/Cas9 system to disrupt universal CAR-T cells with genes lacking TCR and PD-1, Ren and colleagues demonstrated potent anti-tumor effector function *in vitro* and animal models [167]. Knockout of CD3-signaling regulator diacylglycerol kinase for resistance to PGE2 could also augment CAR-T cell abilities [168]. Nevertheless, the safety and feasibility of genome-editing technology to reverse checkpoint-induced inhibitory signaling in clinical patients are still in non-stop exploration.

Similar to PD-1, LAG-3 and Т cell immunoglobulin and mucin-domain containing-3 (Tim-3) known as T cell exhaustion markers functioned as coinhibitory receptors to throttle T cell proliferation and cytokine production [98, 169]. Shapiro et al. found that soluble LAG3 promoted tumor cell activation and anti-apoptotic effects [170]. Afterward, blocking LAG-3 ramped up T cell activation, which rendered LAG-3 evolve into a popular target. Equivalently, targeting Tim-3 is also a prospective strategy [169]. Indeed, checkpoint blockades have revolutionized the field of immuno-oncology, which could be operative against malignancies that fail CAR-T cell therapy and refuel CAR [164]. However, one hypothesis suggests that CAR-T cell therapy for most tumor cells with unexpressed targeted antigen is unlikely to be successful unless combination strategies that enhance bystander effects are applied, and neither anti-PD-1/CTLA-4 antibodies stir bystander effects [171]. Combination therapy could also increase on-target off-cancer toxicity. Inspiringly, Rafiq et al. modified CAR-T cells to secrete PD-1-blocking scFv [172]. ScFv-secreting CAR-T cells were equally effective or superior to combination therapy with CAR-T cells and a checkpoint inhibitor. Notably, this strategy could prevent the toxicities connected with systemic checkpoint inhibition.

3.4 Targeting tumor antigen heterogeneity

In the previous section on targeting immunosuppressive cells and/or factors, we covered part of targeting tumor antigen heterogeneity. Admittedly, as CAR-T cells targeting solid tumors become increasingly effective due to tumor antigen heterogeneity, clinical outcomes will be constrained [112]. Single-target CAR-T cells typically lead to positive selection of antigen-mismatched tumor cells, thereby dampening durable efficacy. Conversely, bispecific CAR-T cells can launch a dual attack on evading cancer cells to avoid ineffective treatment due to heterogeneous target antigen expression and overgrowth of cancers lost a single targeted antigen

[173]. CAR monomers of bispecific CAR-T cells comprised two patterns. One was two distinct scFvs 'hand-in-hand' on one cell, whereas the other was two independent CAR monomers with distinct scFvs on a single cell. Recent studies showed that bispecific CAR-T cells could enhance tumor-suppression capacity through dual antigen recognition and internal activation to effectively eradicate tumor cells [174, 175].

Another way to achieve multiple targets is to create bispecific T cell engagers (BiTEs) linking CD3 scFvs and tumor-associated antigens. BiTEs are bispecific antibodies that redirect T cells to target antigen-expressing tumors. Compared with singletarget CAR-T cells, BiTE CAR-T cells demonstrated prominent activation, cytokine production, and cytotoxicity in response to target-positive tumors [176, 177]. Choi et al. developed a bicistronic construct to drive CAR expression specific for EGFRvIII and EGFR. Unlike EGFR-specific CAR-T cells, BiTE-CAR redirected CAR-T cells, efficiently recruited untransduced bystander T cells against heterogeneous tumors, and was not toxic to human skin grafts in vivo [177]. In conclusion, BiTEs secreted by T cells exhibit robust anti-tumor function, substantial sensitivity, and specificity, establishing a bridge between CAR-T cells and solid tumors [178].

4. Future Directions for CAR-T Cell to Optimize TIME

Despite countless researchers working on the clinical translation of CAR-T cells to overcome the unfriendly TIME, there are numerous hard nuts to crack without any clue. Table 2 lists the ongoing clinical trials of CAR-T cell therapy in solid tumors and no results for all studies in clinical applications. Solid tumors excel at selectively fascinating or avoiding leukocyte subsets and inducing dysfunctional or immunosuppressive phenotypes on resident leukocytes to promote immunosuppression and/or tumor progression [8]. The road for CAR-T cells to conquer the chilly TIME of solid tumors is extremely tortuous. Several reports indicated that aggressive cancer growth was primarily driven by mobilization of immunosuppressive leukocytes in TME rather than loss of tumor immunogenicity following a relatively long incubation period [83, 84]. Due to the spatial and temporal variability of TIME, both immune composition and tumor-specific targets undergo dynamic variation in genotype, phenotype and transcriptome, further illustrating the undisputed primacy of targeting TIME [179].

Table 2. Representative CAR-T cell clinical trials for solid tumors.

NCT Number Phase Status Tumor types		Tumor types	Interventions	Study result
NCT03851146 I	Completed	Advanced Cancer	LeY CAR-T cells	-
NCT03706326 I II	Unknown status	Advanced Esophageal Cancer	Anti-MUC1 CAR-T cells	-
NCT03874897 I	Recruiting	Advanced Solid Tumor	CAR-CLDN18.2 T-Cells	-
NCT02862028 I II	Unknown status	Advanced Solid Tumor	HerinCAR-PD1 cells	-
NCT05287165 I	Recruiting	Digestive System Neoplasms Pancreatic Cancer Colorectal Cancer	IM96 CAR-T cells	-
NCT05275062 I	Recruiting	Gastric Cancer Esophagogastric Cancer Pancreatic Cancer	IM92 CAR-T cells	-
NCT03356795 I II	Unknown status	Cervical Cancer	Cervical cancer-specific CAR-T cells	-
NCT05089266 I	Not yet recruiting	Colorectal Cancer	aPD1-MSLN-CAR-T cells	-
NCT05415475 I	Recruiting	Colorectal Cancer Esophageal Cancer Stomach Cancer Pancreatic Cancer	CEA CAR-T cells	-
NCT04503980 I	Recruiting	Colorectal Cancer Ovarian Cancer	aPD1-MSLN-CAR T cells	-
NCT05341492 I	Recruiting	Lung Cancer Breast Cancer	EGFR/B7H3 CAR-T cells	-
NCT04581473 I II	Recruiting	Gastric Adenocarcinoma Pancreatic Cancer Gastroesophageal Adenocarcinoma	CT041 autologous CAR-T cells	-
NCT05131763 I	Recruiting	Hepatocellular Carcinoma Glioblastoma Medulloblastoma Colon Cancer	NKG2D-based CAR-T cells	-
NCT02932956 I	Active, not recruiting	Liver Cancer	Glypican 3-specific CAR-T cells	-
NCT04489862 I	Recruiting	Non-small-cell Lung Cancer Mesothelioma	aPD1-MSLN-CAR-T cells	-
NCT04864821 I	Not yet recruiting	Osteosarcoma Neuroblastoma Gastric Cancer Lung Cancer	Targeting CD276 CAR-T cells	-
NCT04981691 I	Recruiting	Refractory Malignant Solid Neoplasm	anti-MESO CAR-T cells	-
NCT03356782 I II	Recruiting	Sarcoma Osteoid Sarcoma Ewing Sarcoma	Sarcoma-specific CAR-T cells	-
NCT02107963 I	Completed	Sarcoma Osteosarcoma Neuroblastoma Melanoma	Anti-GD2-CAR engineered T cells	-
NCT03545815 I	Recruiting	Solid Tumor	anti-mesothelin CAR-T cells	-
NCT05437315 I II	Recruiting	Solid Tumor	bi-4SCAR GD2/PSMA T cells	-
NCT05382377 I	Recruiting	Solid Tumor	NKG2D CAR-T cells	-
NCT05373147 I	Recruiting	Solid Tumor	aPD1-MSLN-CAR-T cells	-
NCT04976218 I	Recruiting	Solid Tumor EGFR Overexpression	TGFβR-KO CAR-EGFR T Cells	-
NCT04467853 I	Recruiting	Solid Tumors	LCAR-C18S cells	-

-: No Results Available

Recently, the rapid advancement of nanomedicine has provided inimitable insights into the security and durability of CAR-T cell therapy. Nanoparticles that achieve targeted delivery and prolong the retention time of carried drugs could better remodel the tumor immunosuppressive environment than traditional drugs [180]. Luo and colleagues developed IL-12 nanostimulantengineered CAR-T cell (INS-CAR-T) biohybrids that not only evoked robust anti-tumor efficacy and biosafety via immune feedback but allowed for controllable drug effects [181]. INS-CAR T biohybrids enjoyed various characteristics, including elevated expression of anti-tumor factors, efficient recruitment, deep penetration, and selective proliferation. Several researchers put forward an alternative strategy that CAR-T cells engineered to produce extracellular vesicles containing RN7SL1 by delivering the pattern recognition receptor agonists could effectively stimulate anti-tumor immunity [182]. These CAR-T deliverv systems cells with or functional nanochaperones could be a powerful tool with the possibility to shift the immunologic landscape and

outlook for solid tumors (Figure 4A).

Radioactive material is extensively utilized and may have distinct immunomodulatory effects locally and systemically. In addition to boosting local expression of multiple cytokines and promoting vascular normalization, radiotherapy could activate endogenous target antigen-specific immunity to yield complementary benefits, thereby maximizing the effect of CAR-T cells [183-185]. Radionuclide-based molecular imaging afforded the visualization and therapeutic monitoring of CAR-T cells through cellular radiolabeling approach or gene imaging strategies in determining whether CAR-T cells homed and infiltrated into the tumor bed, as well as their survival and persistence in TIME [186]. The endogenous cell imaging could reflect the immune status and functional information of T cells and even delineate the developmental trajectory of immune cells. Dynamic monitoring will help researchers understand cellular behavior in vivo, allowing better infusion timing and dosage optimization to avoid potentially lethal systemic toxicity (Figure 4B) [187]. As for assessing the prognosis and suitability of



Figure 4. Cutting-edge technologies to augment the security and durability of CAR-T cells. There are increasingly revolutionary technologies and researches applied in CAR-T cell therapy, which provide a bright guide to reshaping TIME. A Nanoparticles could achieve targeted delivery and prolong the retention time of carried drugs. Nanostimulant-engineered CAR-T cells not only could evoke robust anti-tumor efficacy and biosafety via immunofeedback but allow for controllable drug effects on CAR-T cells. I CAR-T cells penetrate the tumor location and elicit the first killing. II Secreting pro-inflammatory factors to trigger immune cells and endogenous T cells at the right moment could initiate secondary killing and form a positive anti-tumor cycle from CAR-T cells. III The cancer cells with CAR antigen loss could be recognized and killed by CAR-T cells delivering peptide antigen to cancer cells. B Radioactive material is extensively utilized and may have distinct immunomodulatory effects both locally and systemically. The combination of radiotherapy and CAR-T cell therapy could maximize the effect of immunotherapy. The separated T cells are directly labeled and genetically modified. The radiolabeled CAR-T cells are infused into tumor-bearing mice and monitored by PET/SPECT imaging. If the tumor worsens in tumor-bearing mice, then redesign and relabel the CAR-T cells with radioactive material.

CAR-T cell therapy, types of TIME could be considered as a novel biomarker to stratify the overall survival risk of untreated tumor patients and tertiary lymphoid structure (TLS) could be used as a sign of effective immunotherapies [179, 188]. Upregulation of TLS in most tumors could foster CAR-T cell immunotherapy [188].

Until recently, tumor-on-chip offered a renewed direction for preclinical CAR-T cell research. Compared with conventional in vivo animal models and in vitro planar cell models, emerging tumor-on-chip platforms integrating microfluidics, tissue engineering, and 3D cell culture have successfully mimicked the key structural and functional properties of TME in vivo [189-191]. Several studies have attempted to generate functional immune cells from human pluripotent stem cells in combination with neo-platforms, such as the generation of T-cell progenitors from hematopoietic organoids [192] and the invention of hematopoietic organs-on-chips [193]. The studies pushed the limit of pluripotent stem cells to produce immune cells useful for CAR-T therapy. Thus, tumor/organ-on-chip platforms hold great promise as more accurate and realistic models for investigating the immunosuppressive mechanisms of TIME as well as the drug toxicity and efficacy of CAR-T cell therapy before being applied to individuals. Additionally, using a comprehensive single-cell gene expression and TCR sequencing dataset for pre- and post-infusion CAR-T cells, Wilson et al. [194] found a unique signature of CAR-T cell effector precursors present in pre-infusion cell products. These effector exhibited precursor CAR-T cells functional superiority and decreased expression of the exhaustion-associated transcription factor, consistent with post-infusion cellular patterns observed in patients. Engineering alternative immune cells (such as NK and macrophages) to express CAR targeting molecules is also an optimal strategy. CAR-NK cells or macrophages were shown to induce a pro-inflammatory TIME and boost anti-tumor T cell activity [195, 196]. Collectively, these nascent technologies or preclinical studies offer great potential for therapeutic applications.

5. Conclusion

CAR-T immunotherapy is revolutionizing the paradigm of cancer therapy. However, unique obstacles posed by solid tumors remain a Gordian knot for CAR-T immunotherapy. The intricate interactions among immune cells, tumor cells, immune molecules, and cytokines form TIME that throttles immune responses and encourages tumor development. By precisely mapping the complex regulatory network of TIME and drawing the optimization strategies, we could provide patients with more effective and tailored CAR-T immunotherapy. To address the hostile TIME and mitigate or even obviate the risk of serious adverse cutting-edge technologies, reactions, such as nanoparticle delivering systems, radionuclide-based imaging, and novel tumor-on-chip molecular platforms are gradually applied for CAR-T immunotherapy. Indeed, the current understanding of TIME is only the tip of an iceberg. Nevertheless, we believe that by progressively regulating the harsh TIME to reshape a friendly microenvironment, CAR-T immunotherapy will produce a more astonishing breakthrough.

Abbreviations

BiTE: bispecific T cell engager; Breg: regulatory B cell; CAR: chimeric antigen receptor; CTL: cytotoxic lymphocyte; CTLA-4: cytotoxic T lymphocyteassociated antigen 4; CXCL: CXC chemokine ligand; DC: dendritic cell; DNR: dominant-negative receptor; FcγR: Fc gamma receptor; FRβ: folate receptor beta; GD2: disialoganglioside; GITR: glucocorticoidinduced tumor necrosis factor receptor; GM-CSF: granulocyte-macrophage colony-stimulating factor; G-MDSC: granulocytic MDSC; HCC: hepatocellular carcinoma; ICOS: inducible co-stimulator; IFN-y: interferon-gamma; IL: interleukin; iNOS: inducible nitric oxide; LAG-3: lymphocyte activation gene-3; L-Arg: L-Arginine; MC: Mast cell; M-CSF: macrophage colony-stimulating factor; MDSC: myeloid-derived suppressor cell: M-MDSC: monocytic MDSC; Msln: mesothelin; NAP: neutrophil-activating protein; NK cell: natural killer cell; Nrf2: NF erythroid 2-related factor 2; PD-1: programmed death-1; PDGF: platelet-derived growth factor; PD-L1: PD-1 ligand; PGE2: prostaglandin E2; RNS: reactive nitrogen species; ROS: reactive oxygen species; TADC: tumor-associated dendritic cell; TAM: tumor-associated macrophage; TAMC: tumorassociated mast cell; TAN: tumor associated neutrophil; TCR: T-cell receptor; Teff: effector T cell; TGF- β : transforming growth factor- β ; Th1: T helper type 1; Tim-3: T cell immunoglobulin and mucin-domain containing-3; TIME: tumor immune microenvironment; TLS: tertiary lymphoid structure; TME: tumor microenvironment; TR2: TRAIL receptor 2; Treg: regulatory T cell; VEGF: vascular endothelial growth factor.

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Author contributions

ZQL, XWH, and QD provided direction and

guidance throughout the preparation of this manuscript. ZKZ, ZQL, and QD wrote and edited the manuscript. QD reviewed and made significant revisions to the manuscript. HX, JXL, HYL, QD, and ZQL collected and prepared the related papers. All authors read and approved the final manuscript.

Competing Interests

The authors have declared that no competing interest exists.

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